

**INSTITUTE
FOR CANCER
RESEARCH**

**ANNUAL
REPORT
2019**

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The photographic theme of this year's
Annual Report is *Humans at ICR*.
- The most important part of ICR is
by far its human resources and our
collective competence.

FRONT PAGE:
Optics inside a flow cytometer. Flow
cytometry is used in our research in
cancer immunology, immune-oncology
and immune monitoring as well as in
cell cycle and cell signalling analyses.
Advanced technology and competence
provided by the Department of Core
Facilities is a pillar for cutting-edge
research at the Institute for Cancer
Research. Advanced infrastructure
is financed by OUH, UoO, RCN,
Radiumhospitalets Legater, and Norsk
Hydros Fond.

PAPER: 150/300 Profimatt
CIRCULATION: 800

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Taking our legacy into the future

- Who we are and what we do

WHO WE ARE

1954

LEGACY IN RESEARCH, INNOVATION AND EDUCATION

The Institute for Cancer Research (ICR) was founded in 1954 representing 65 years of continuous advances in cancer research. Through the years, our work has significantly contributed to better understand cell cycle and cell division, cancer cell biology, cancer genetics, cancer immunology, radiation biology, molecular oncology, tumor biology, and metastasis. In addition, we have contributed to novel methods and implementation of a number of new technologies to help cancer diagnostics and monitoring.

RESEARCH

There is also a proud history of translational research and innovation, for example on photomedicine leading to companies such as Photocure and PCI Biotech and with numerous other past and present translational and innovation projects at the ICR in various stages of development.

INNOVATION

Education and dissemination are an integrated part of ICR activities and many masters- and PhD-students and postdocs have received their training at the ICR and have pursued successful careers in research, clinical medicine, administration or in the biotech and pharma sector.

EDUCATION

IN NUMBERS

350

RESEARCH STAFF

Approximately 350 research staff in 25 research groups, 6 core facilities and 1 admin unit (organization see pages 8-9)

PAST 5 YEARS (2015-2019)

ICR researchers have raised more than 1.0 billion NOK extramural funding (of total 1.5 billion accounted)

1.00

BILLION NOK

1650

WORKING YEARS

Put in 1650 working years (FTEs)

1100

PUBLICATIONS

Leading to production of more than 1100 (1112) publications (mean IF 6.1, median IF 4.5)

150

EDUCATED PhDs and MScs

Educated 66 PhDs and 84 MScs that have graduated

(for 2019 Key Figures see pages 10-11).

WHAT WE DO

VISION: EXCELLENCE IN FIGHTING CANCER

WE AIM TO:

- Stay at the cutting edge of cancer research
- Educate the leaders of tomorrow in cancer research, cancer diagnostics and treatment
- Take a prominent part in developing of new OUH and national strategies for advanced molecular cancer diagnostics, cancer precision medicine and experimental cell therapy (see also page 15)

VALUES: QUALITY, INTEGRITY, TEAMWORK, VISION

VISION

- Objectives
- Ambition
- Global influence
- Cutting-edge
- Passion

QUALITY

- Respect
- Excellence in research
- Trust
- Competence

INTEGRITY

- Ethics
- Responsibility
- Loyalty
- Demands

TEAMWORK

- Synergy
- Diversity
- Generosity
- Partnerships
- Solidarity
- Openness

EXCELLENCE IN FIGHTING CANCER

GOALS 2019-2020

- Strengthen translational research
- Strengthen contact, coordination and collaboration with clinicians and diagnostic staff in OUH CCC and beyond
- Build further excellence in research
- Establish new SAB
- Increase internationalisation and technology development

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Introduction by the Director

I am proud to present you with our Annual Report for 2019. The Institute for Cancer Research (ICR) is an institution with approximately 350 employees plus students organised in 25 research groups complemented with cutting-edge core facilities. The ICR is a premier institution in basic and translational cancer research on a national and international arena and has a strong prior track record in translation and innovation. More than one third of the ICR staff is international and come from 37 different countries.

In 2019 we worked in the ICR Leadership group, among the Group Leaders and in all ICR departments and groups to set out and integrate our Vision and Values (Quality-Integrity-Teamwork-Vision, see the previous page). We also set out Goals for 2019-2020, both as presented under the heading “The ICR into the future!” in the 2018 Annual Report and on the preceding pages in this report. Among our goals for 2019 and 2020 are to (1) strengthen translational research, (2) to improve contact, coordination and collaboration with clinicians and diagnostic staff in the Oslo University Hospital Comprehensive Cancer Centre (OUH-CCC) and beyond, (3) to build further excellence in research where we would like to see originality, depth, quality, and international value.

With respect to goals 1 and 2, I think there are numerous outstanding examples of collaborative translational and clinical projects throughout this report. Furthermore, the ICR is a central component in the new and developing strategies for precision medicine and for cell therapy in the Division for Cancer Medicine and the OUH-CCC. To build further excellence (goal 3), we would like to see a further increase in the quality of the scientific output. Against this backdrop, it is interesting to see that the median impact factor of the papers produced is up from 4.1 in 2017 to 5.0 in 2018 and 4.5 in 2019. Moreover, the fact that the output in terms of the number of papers is increasing, with 241 papers in 2019 and more than 40 in the two first months of 2020 (versus 220 and 186 papers in 2017 and 2018, respectively). This is excellent, particularly as almost half of the production also has first and/or senior authors from ICR. ICR innovation activities also appear to be progressing well with 10 new DOIs, 7 new patent applications filed and 5 patents granted from ICR inventors. In addition, company collaborations and interaction appear to be increasing.

More than 2/3rds of the total ICR funding in 2019 (234.4 of 333.5 MNOK) came from extramural grants which are higher than ever before. ICR scientists have also been able to obtain substantial new funding and secured major new grants from the Norwegian Cancer Society, the Research Council of Norway, and the South-Eastern Norway Regional Health Authority as well as private and international sources.

We celebrate our victories at the ICR, and in 2019 we have marked the fact that two of the newly awarded OUH Strategic areas went to the ICR-led projects TEAM-ACT (led by Ragnhild A. Lothe) and STRATCELL (led by Karl-Johan Malmberg). We have celebrated a number of grants from the RCN including several young talent grants and including the start of Digital Life PINPOINT project, and multiple grants from the Cancer Society, the Regional Health Authority for South-Eastern Norway and other sources. ICR is also well represented in the new Cancer Society Expert Groups for Lung Cancer (headed by Åslaug Helland) and Pancreatic Cancer (headed by Caroline Verbeke). We have also had the official opening of the privately funded InvaCell project that formalized a collaboration between the ICR and the Curie Institute (with Harald Stenmark and Philippe Chavrier as PIs, donor Trond Paulsen). At Christmas, we celebrated the award of a highly prestigious ERC Consolidator Grant to Johanna Olweus. We have marked the fact that the University of Oslo (UiO)/OUH was granted a prestigious 6-mEUR Horizon2020 grant, RESCUER, for precision medicine research on breast cancer (headed by Vessela Kristensen and with strong ICR participation) and that Åslaug Helland received a large KLINBEFORSK grant. The ICR has also received very significant grants for new instrumentation from the Radium Hospital Foundation and from Norsk Hydro's Fund for Cancer Research in 2019. We also celebrated internally the award of the OUH Early Career Award to Anita Sveen, the EU Innovation Radar Prize to Kristian Berg and Theodossis Theodossiou, the award of the Ragnar Mørk's Prize for Outstanding Research to Karl-Johan Malmberg and the award of the new ICR Prizes Researcher-of-the-year and Employee-of-the-year for 2019 to Theodossis Theodossiou and Peter Wiedswang, respectively. Furthermore, OUH awarded prizes in their bi-annual assessment of best papers to ICR researchers Marthe Løvf and Rolf Skotheim (May 2019), Muhammad Ali, Zsolia Foldvari, Eirini Giannakopoulou and Johanna Olweus (Nov 2019) and Marte Sneeggen and Kay Oliver Schink (Nov 2019).

In the Department of Cancer Genetics, we welcomed Tero Aittokallio as a new Group Leader for the Computational Systems Medicine in Cancer Group from September 2019 (recruited from Finland) and in Department of Cancer Immunology, Jon Amund Kyte has been appointed Group Leader for the Immunotherapy Against Solid Cancers Group from 2020.

ICR's strong standing in Norwegian research is also illustrated by the fact that members of the institute lead a Centre of Excellence (CoE) Centre for Cancer Cell Reprogramming (CanCell, Director Harald Stenmark) and the three K.G. Jebsen centres for Cancer Immunotherapy (Director Johanna Olweus), Colorectal Cancer Research (Director Ragnhild A. Lothe), both

“Research and innovation with patient benefit in mind”

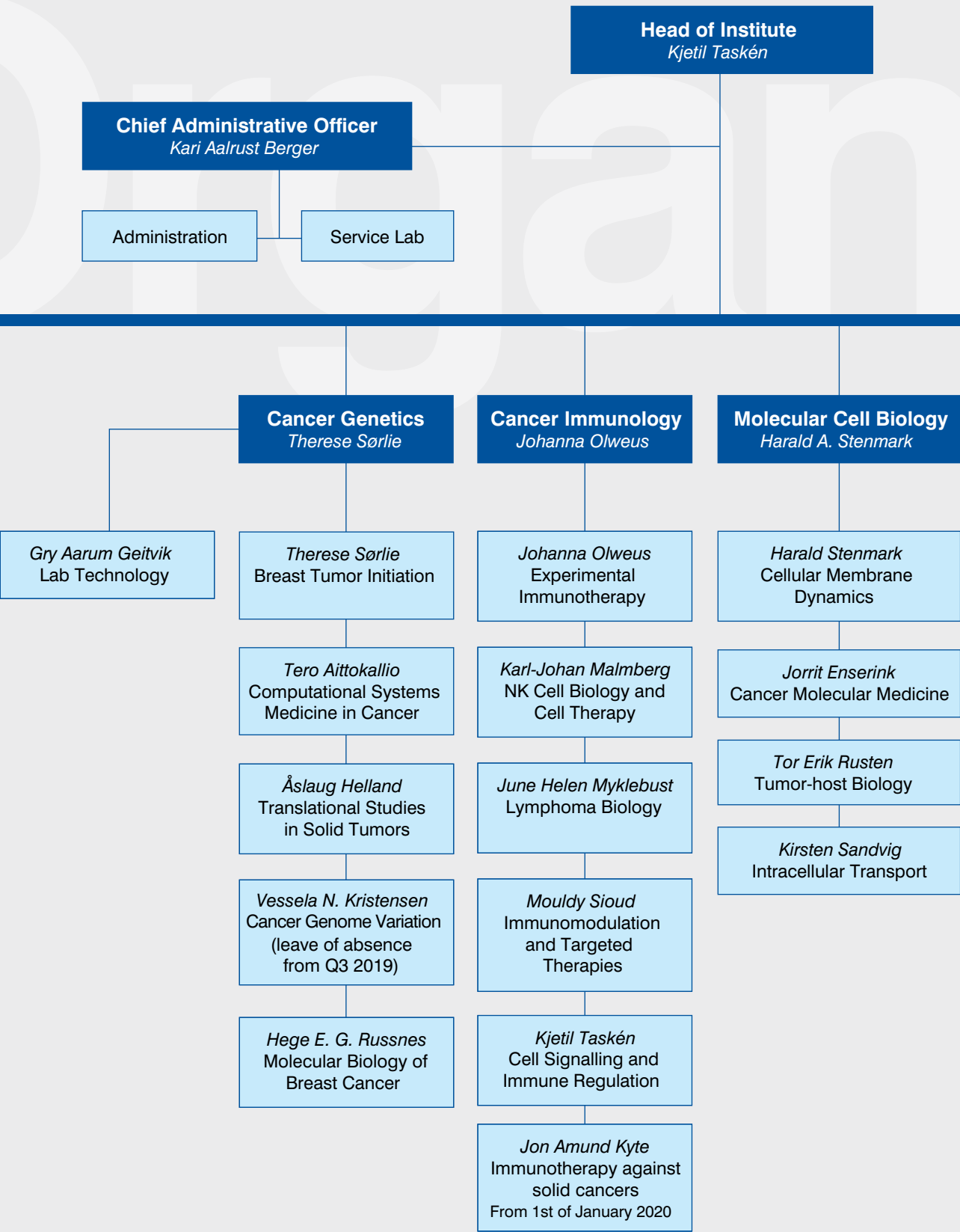
in the extension phase, and for B Cell Malignancies (Deputy Director June Myklebust), all with strong participation from ICR groups.

The recent advents of the Covid-19 pandemic have had major impacts world-wide and on the Norwegian society at large as well as on OUH and UiO research activity at this time. At the ICR, as part of the hospital and with good support from the leadership at all levels, we decided to keep the institute open throughout the pandemic. - This in recognition of the importance of the science we do and to support the hospital operations. With appropriate risk assessments, precautions and personal protection measures such as social distancing, hygiene measures, reduced density at work and somewhat reduced activity, this has so far gone well with strong efforts from all our employees in these difficult circumstances.

The ICR sets out to maintain the excellent science and outstanding production and to further build excellence by organising more collaborative efforts at all levels to deal with grand challenges in cancer medicine and to position the ICR in national and international alliances and consortia. We aim to be a significant partner for the clinical activities in the Division of Cancer Medicine and the OUS OEI-accredited Comprehensive Cancer Centre (CCC). We continue feeding results into a translational research path and to have **patient benefit** in mind in all aspects of research and innovation.

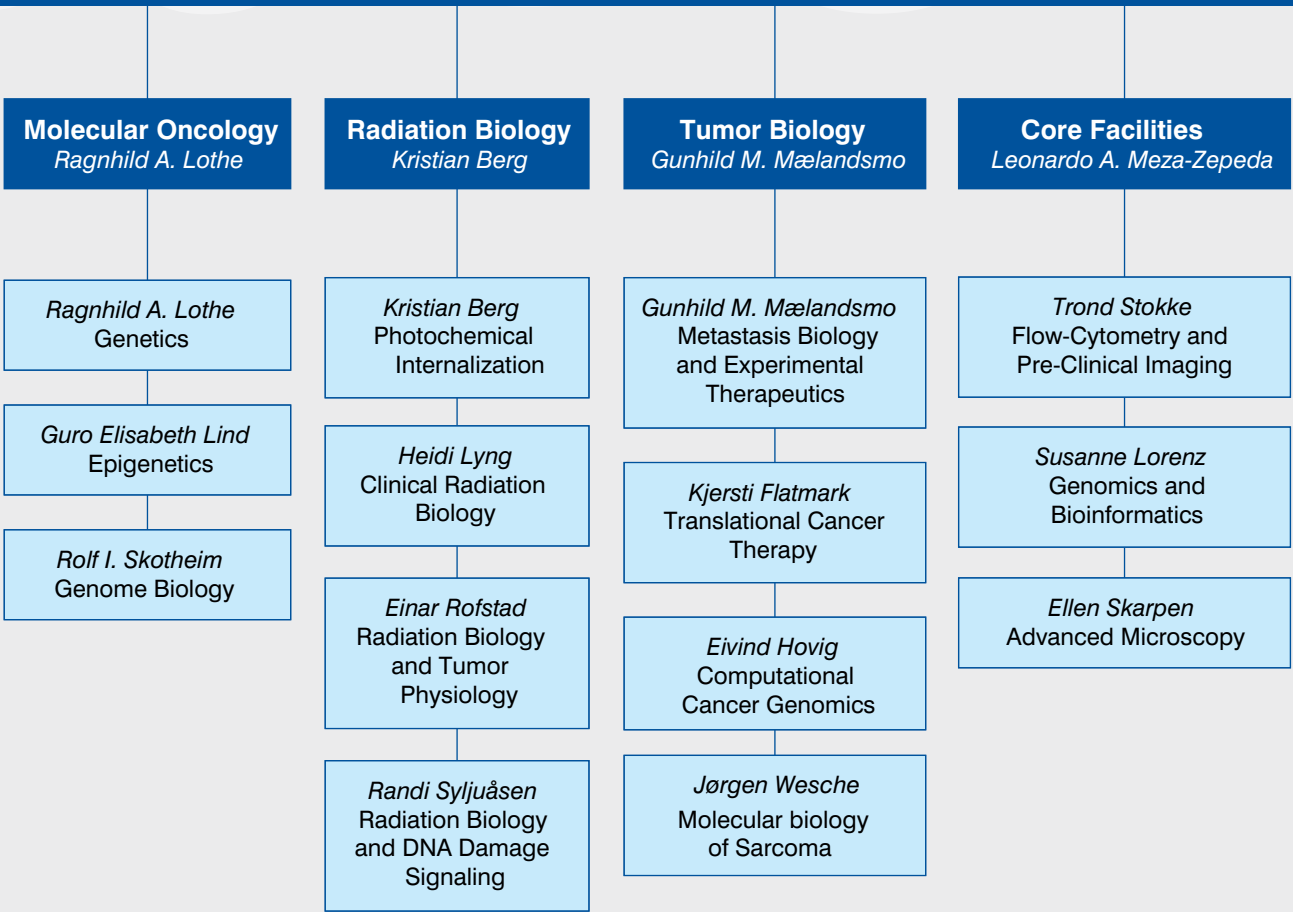
April, 2020
Kjetil Taskén, Head of the ICR

Organisation



The Institute for Cancer Research

Institute for Cancer Research is organized in 6 research departments with 25 research groups, and one Department of (6) Core Facilities.

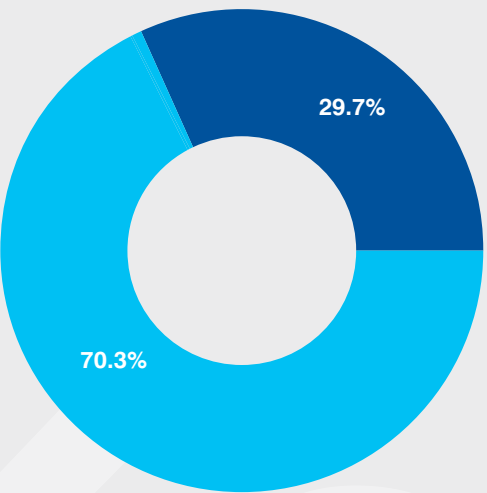


Key figures 2019

Funding

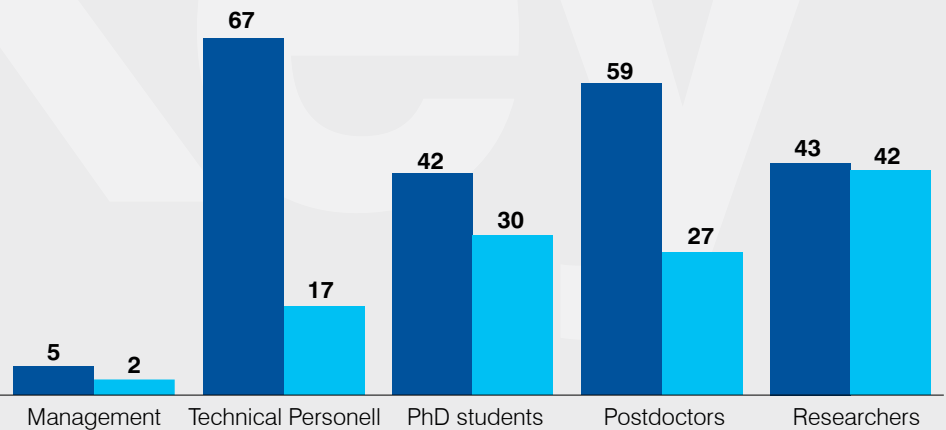
Percent

Actual Institute expenditure for 2019 by internal and external funding sources (total 333,5 MNOK = approx. 32,8 M€)



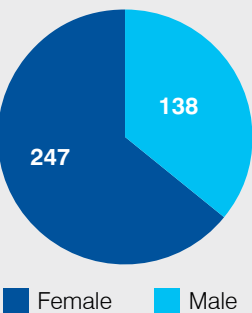
Internal funding
External funding

Employees



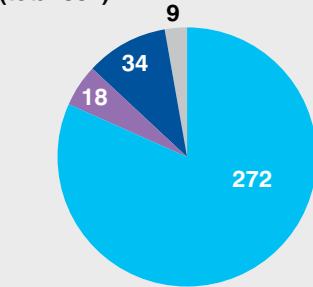
(FTEs)
Female
Male

Employees by Gender (total 385)



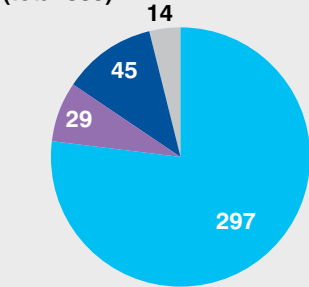
Female
Male

FTEs by Employer (total 334)



ICR
OUH*
UiO
Other
*other than ICR

Employed by (total 385)

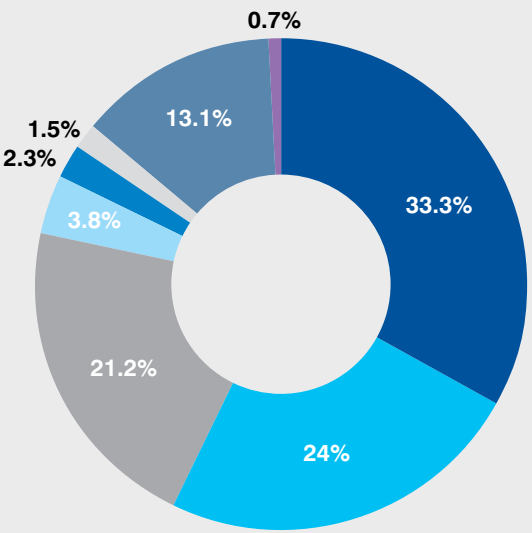


ICR
OUH*
UiO
Other
*other than ICR

External funding by source

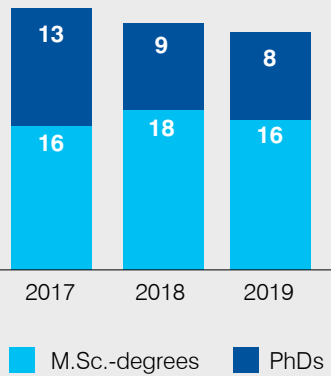
Percent

Sources of external competitive funding for 2019, based on actual expenditure (total 234,4 MNOK = approx. 23 M€)



South-Eastern Norway Regional Health Authority
The Research Council of Norway
The Norwegian Cancer Society
University of Oslo
EU
Other international sources
Other private sources
Other public sources

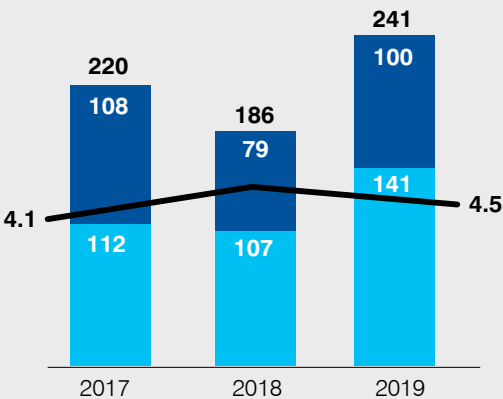
Completed PhDs and M.Sc.-degrees



M.Sc.-degrees
PhDs

Articles published

First or last authorship
Co author
Impact factor median



IMPACT FACTOR

Mean
Median

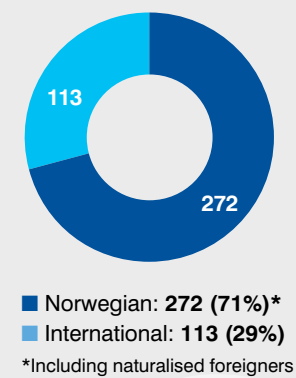
5.8
4.1

6.5
5

6.1
4.5

International Staff Distribution

37 nations are represented
113 people in total are from outside Norway



01

Countries represented by one person

- Bosnia
- Colombia
- Croatia
- Czech Republic
- Equador
- Estonia
- Finland
- Ireland
- Iceland
- Latvia
- Macedonia
- Mexico
- Netherlands
- Pakistan
- Russia
- Singapore
- Slovakia
- Switzerland

02

- People
- Denmark
 - Japan
 - Portugal
 - USA

03

- People
- Austria
 - Greece
 - Iran
 - Serbia

04

- People
- France
 - Hungary

05

- People
- Great Britain
 - Lithuania
 - Poland

06

- People
- Italy

07

- People
- China
 - Spain

08

- People
- Sweden

11

- People
- India

13

- People
- Germany

Oslo University Hospital Comprehensive Cancer Centre

Oslo University Hospital was designated a Comprehensive Cancer Centre by the OECC in 2017, after a process where activities both in clinical practice and in research were evaluated. The cancer research performed by the Institute for Cancer Research (ICR) is a corner stone in our OECC-accredited Comprehensive Cancer Centre (CCC).

The Institute is situated in close proximity to clinical cancer departments and diagnostic laboratories at the Radium Hospital, a cancer-oriented part of Oslo University Hospital where also the Cancer Registry of Norway is located. The collaboration between the ICR and clinical research groups in Oslo University Hospital is an important factor to increase our activities in clinical research. More patients into clinical trials is an expressed aim for our Comprehensive Cancer Centre, and several investigator initiated clinical trials have been developed in collaboration between researchers at ICR and clinical research groups at all locations of Oslo University Hospital. Another of ICR's neighbours is

Oslo Cancer Cluster, - which consist of several biotech and pharmaceutical companies. The proximities between all these actors, provides an excellent environment for synergies and collaboration.

Several Centres of Excellence also including basic research are located at the Institute for Cancer Research, enriching the Comprehensive Cancer Centre with competence and expertise. The extensive international collaboration involving researchers at ICR is also an important asset for the CCC.

Further development of the tight bonds between ICR, clinical researchers and the Cancer Registry, will be important in the coming years and with the new hospital building and proton center that will open for patients in 2023/24. In the integrated organisation of cancer-related activities, the ICR will be a key participant in the further development of Oslo University Hospital as a leading cancer centre in Europe.

Sigbjørn Smeland

Head of Division of Cancer Medicine,
Chair, OUH CCC Board

Åslaug Helland

Research Director, Division of Cancer Medicine
Head OUH, CCC Research Committee

Precision Cancer Medicine at OUH Comprehensive Cancer Centre

The implementation of genomic medicine and individualised treatment has been lagging behind in Norway and other countries in Northern Europe with publicly funded health care systems. Against this backdrop, the Head of the Division for Cancer Medicine and Chair of the OUH Comprehensive Cancer Centre appointed a PCM working group¹ that started its activities in January of 2019. The working group decided to make a concrete plan for PCM implementation and delivered the results of its work to the Division and the CCC Board in June 2019:

VISION AND OBJECTIVES FOR IMPLEMENTATION OF PRECISION CANCER MEDICINE

- *Patients who are referred to OUH with cancer and where advanced molecular cancer diagnostics will be instrumental for selection of treatment should be offered such diagnostics at the right time during the course of the disease.*
- *Cancer patients should have opportunity to receive individualised treatment where this is shown to impact on clinical outcome or where it is probable that it would give patient benefit.*
- *OUH will learn and build competence in advanced molecular cancer diagnostics and individualised treatment and contribute to development and production and dissemination of new knowledge in this area.*

ACTION POINTS NECESSARY FOR IMPLEMENTATION OF PCM, THREE MAIN AREAS:

- Establishment of a platform that can provide advanced (experimental) molecular cancer diagnostics. Develop such service also in the standard process to examine and diagnose patients.
- Increase the volume of clinical trials and number of patients in trials in the precision medicine area (industry studies, researcher-initiated trials).
- Offer individualised cancer treatment in the ordinary standard of care in OUH according to national and international guidelines.

¹ The Cancer Precision Medicine Working Group (Jan-Sept 2019) has consisted of Kjetil Taskén (ICR, chair) Tormod Guren (Onc), Ragnhild Lothe (ICR), Per Magnus Mæhle (CCC), Hege Russnes (ICR/Pat), Gunnar Sæter (CCC), and with Live Fagereng (ICR) as secretary.

² From Oct 2019 the CCC PCM Working Group consists in addition to the above members of: Monica Cheng Munthe-Kaas (Ped), Nils Tore Vetthe (ClinPharm), Ben Davidson (Pat), Espen Enerly (CRN), Åslaug Helland (Onc/CCC), Yngvar Fløysand (Hem), Kristina Lindemann (GynOnc), Turid Vederhus (Rad), Torunn Berge (HSE, observer)



PROGRESS:

- In the spring of 2019, the Division for Laboratory Medicine responded to the needs of the cancer area with respect to action points I and II (above) and decided to establish a Section for Experimental Diagnostics and Research Support in the Department of Pathology to provide a genomic medicine platform to ensure delivery of necessary molecular cancer diagnostics (Head Hege Russnes). The CCC PCM working group also on behalf of the CCC in June organized a national meeting on implementation of advanced molecular cancer diagnostics.
- In the fall of 2020, the PCM working group was reinforced and expanded² and now reports directly to the CCC Governing Board. The work has next focussed on attracting PCM clinical trials (action point II). Specifically, the CCC PCM working group has looked at the possibility of establishing a national PCM trial modelled on the DRUP study in the Netherlands (IMPRESS-Norway trial, national coordinator and PI, Åslaug Helland) matched by a national infrastructure for precision diagnostics, InPreD. The work has involved raising national support, discussion with key stakeholders, planning of a public-private partnership with industry participation and national consensus as well as organising a second national meeting in January 2020.

Departments

- 18 Department of Cancer Genetics
- 26 Department of Cancer Immunology
- 34 Department of Molecular Cell Biology
- 40 Department of Molecular Oncology

- 46 Department of Radiation Biology
- 52 Department of Tumor Biology
- 58 Department of Core Facilities



Department of Cancer Genetics

Headed by Therese Sørli

Our aim is to improve risk estimation, achieve earlier diagnosis and improve prediction of treatment responses and other clinical outcomes for patients with early and advanced stages of both solid tumors and hematologic malignancies. Our research is translational in nature and include functional studies, molecular classification, data integration and pan-cancer analyses. We work towards facilitating the implementation of discoveries into clinical use. A common theme across groups is to achieve a deeper molecular understanding of inter- and intra-tumor heterogeneity and tumor evolution using patient cohorts and mouse models.

We are an interdisciplinary team of 50 researchers, including medical doctors, molecular biologists, computational biologists and highly specialized engineers organized in four research groups and one lab-technology unit. Two of the group leaders hold part-time clinical positions and three have affiliated positions at UiO. The lab technology unit reinforces the department's expertise in state of the art technology and improves exchange of knowledge across research groups and cancer types. This is a key asset leading to increased quality of the department's laboratory work and project management.

For translational studies, we have established a pipeline for high-quality biobanking and secure data handling of patient cohorts with long-term follow-up that enables omics analysis of tumors down to single cell levels. Our clinical database consists of > 3000 subjects from consecutive studies and clinical trials with high quality follow-up information. We are involved in the following studies:

- NeoAva - Neoadjuvant chemotherapy in breast cancer with/without bevacizumab.
- IBCT - Improved Breast Cancer Therapy in the neoadju-

- vant and metastatic setting
- EMIT -Establishment of Molecular profiling for Individual Treatment decisions in Early BC; three-phase study including randomized and observational clinical trials.
- OPTIMA-optimal personalized treatment of early breast cancer using multi-parameter analysis
- ComIT - evaluation of the benefit of radiation in combination with immune therapy for lung cancer
- TREM - Lung cancer patients with EGFR mutations and primary TKI-resistance
- ThoRaT - Lung cancer patients receiving radiotherapy
- NorPACT-1 and 2 - Neo-adjuvant chemotherapy for pancreatic cancer
- ICON - A randomized phase IIb study evaluating immunogenic chemotherapy combined with ipilimumab and nivolumab in patients with luminal B breast cancer
- ALICE - atezolizumab combined with immunogenic chemotherapy in patients with metastatic triple-negative breast cancer
- Oslo2- observational study with comprehensive biobanking

We are part of extensive collaborations at the institutional, national and international levels, and partnering in several national and international networks and consortia; for instance, the National Breast Cancer Research Network, the Norwegian Cancer Society expert groups on lung and pancreatic cancer, the Regional Research Network on Extracellular Vesicles, Personalized Cancer Treatment and Metaflammation, International Cancer Genome Consortium (ICGC), EuroPDX, the Breast Cancer Association Consortium (BCAC); EU funded projects (EpiMark, Cancer-ID, Gender Net Plus, HERCULES and RESCUER). The total number of publications in 2019 was 68.

Breast Tumor Initiation

Group Leader: Therese Sørli



“Understanding cell fate decisions in tumor progression”

ABOUT

Our group studies molecular aspects of breast tumor initiation and progression with a special emphasis on identifying cell(s) of origin of molecular subtypes and the transition from in situ to invasive breast cancer. The group counts 10 members, including the group leader and professor (TS), three postdocs, three PhD students, one master student, and two engineers as well as two affiliated scientists (in 20% and 10% positions) and one affiliated breast surgeon. Two members are MDs and one is DVM. We have a broad expertise in laboratory technologies that includes high-throughput genomic technologies, in vivo lineage-tracing, 2D and 3D in vitro culture techniques, in situ hybridization, confocal microscopy, and FACS analysis. We use patient cohorts and mouse models (transgenic and patient derived xenograft - PDX) in our studies. We also have expertise in bioinformatics and statistical modeling.

AIMS

Our aim is to identify the cellular origins of breast tumors and the mechanisms underlying tumor initiation and further development into the heterogeneous molecular subtypes. Through an increased understanding of how early lesions progress to more advanced stages, we aim to contribute to improved strategies for early intervention and more precise treatment.

PROJECTS

- Characterize breast cancer subtype-specific progression pathways
- Explore the role of LGR5-expressing cells in the mammary tumorigenesis
- Investigate the role of Hedgehog/GLI1 signaling in breast cancer progression
- Investigate the role of FOXA1 and FGFR1 in endocrine resistant breast cancer
- Genomic and functional analysis of therapeutic targets

RECENT ACHIEVEMENTS

8 publications by group members in 2019. One Master of Science completed.

Computational Systems Medicine in Cancer

Group leader: Tero Aittokallio



“A systems view of disease networks to pinpoint critical targets and their interaction partners”

ABOUT

The group was launched in September 2019 when Tero Aittokallio moved from Finland (FIMM, University of Helsinki and University of Turku). Our group has expertise in integrating multi-omics profiling and clinical information from cancer patients using mathematical and statistical approaches, such as machine learning and network modeling. We carry out multidisciplinary projects in collaboration with other researchers from the institute and as part of international projects, where we develop, test and implement novel practices of how to use artificial intelligence (AI) and machine learning (ML) models in translational and clinical studies. We believe that combining functional, molecular and genomic profiling information is critical for next-generation precision medicine applications, where integrative modeling and clever use of patient-level big data will pinpoint effective and selective targets for personalized therapies.

AIMS

The modeling aim is to develop novel supervised learning approaches to identify multi-omic features predictive of clinical outcomes for individual patients by means of efficient AI/ML models that maximize the accuracy of outcome predictions using minimal panels of biomarkers. The medical aim is to optimize treatment outcomes for individual patients using the maximally predictive models and minimal biomarker signatures that enable real-time and cost-effective diagnostics and prognosis.

PROJECTS

- Multi-omics prediction of clinical outcomes for precision oncology applications
- AI-guided treatment optimization by means of cost-effective biomarker panels
- Decision support systems for real-time patient monitoring and adaptive trials

RECENT ACHIEVEMENTS

In 2019, the group published 11 articles in peer-review journals and 2 book chapters. We are partnering in several international translational projects, including ERA-PerMed project JAK/STAT TARGET, EU-H2020 project HERCULES, and the UK Breast Cancer Now Catalyst Programme for predictive markers for TNBC drug responses. In 2019, the group received 3-year open-project grant from South-Eastern Norway Regional Health Authority for AI-guided treatment optimization by means of multi-omics biomarker panels.

Translational Studies in Solid Tumors

Group leader: Åslaug Helland



”With starting point in clinical studies and clinical questions, we do molecular analyses aiming at improving outcome for cancer patients”

ABOUT

Our group focuses on translational studies on solid tumours, with a special interest in pancreatic, lung and colorectal cancers. By increasing the understanding of the underlying biology of tumour development, we will improve precision medicine in cancer care. Several of our projects include material from patients included in clinical studies, and we have detailed clinical data from all patients. Predictive biomarkers and mechanisms of resistance is in focus.

We are organized into three project groups, headed by Elin H. Kure (Professor USN), Odd Terje Brustugun and Åslaug Helland (Professor UIO), with a total of 17 members. Seven of these are MDs. We are three researchers, three postdocs, seven PhD-students, one study nurse and three engineers.

AIMS

The ultimate goal is to increase our ability to offer personalised cancer treatment, and thereby improve prognosis.

Sub aims are:

- Identification of biomarkers in blood samples, for diagnostics, follow-up and prognostication
- Identification of tumour biomarkers for prediction of therapy response and for prognostication

PROJECTS

- Molecular characterization (-omics) of pancreatic- and lung cancers
- Proteogenomic analysis of pancreatic tumors and characterization of circulating biomarkers in free plasma and exosomes
- Identification of circulating plasma biomarkers in colorectal cancers (the Nordic VII clinical trial)
- Protein (TMA) analyses in lung cancers
- Serum predictive markers for therapy response
- Elucidate mechanisms for therapy resistance
- Expand biomarker identification material to stool (microbiome) and urine
- Investigate combination of radiotherapy and immunotherapy
- Gender differences in side effects on immunotherapy

RECENT ACHIEVEMENTS

In 2019, the group published 18 papers in peer-reviewed journals. We have had several talks at national and international meetings, and are PIs on >20 translational and clinical studies. An ERA-network we are part of received funding (Gender-net). We are partners in several South-Eastern Norway Regional Health Authority networks (NORMAN, ReMics, NIRO), and received approximately 36.2 mill NOK in research funding (Elin Kure 1.2 mill, Odd Terje Brustugun 2 mill, Åslaug Helland 33 mill).

Cancer Genome Variation

Group leader: Vessela Kristensen



“Germ-line and somatic genetic and epigenetic variants in cancer and pharmacogenomics”

ABOUT

The Kristensen group with 2 senior scientists, 3 postdocs, 2 PhD students, 1 MSc and 2 research technologists moved Q3-2019 to Institute for Medical Genetics (IMG) at OUH. The group will continue to work towards intensive and fruitful collaboration between ICR and IMG and with the Institute for Clinical Medicine, University of Oslo. Group members work closely together and in collaboration with breast clinicians, pathologists and oncologists. A remaining project group at ICR consists of 1 senior scientist and 3 PhD students. This project group is headed by Thomas Fleischer (Epigenomics of Breast Cancer) and works to characterize epigenetic alterations in breast cancer, and assess the implications for breast cancer pathogenesis and utilization for precision medicine (<http://ous-research.no/fleischer/>).

AIMS

The Cancer Genome Variation group is working to understand how genetic variation affects occurrence of somatic alterations, gene expression patterns and genome wide copy number alterations in human tumours (<http://ous-research.no/kristensen/>)

PROJECTS

Together with our collaborative network we received several major grants in 2019: from EU Horizon 2020 (RESCUER: 63 mill NOK), EU EraCoSysMed (PI, 3 mill NOK), South-Eastern Norway Regional Health Authority (Open Project, 9 mill NOK; PhD student, 3 mill NOK). Projects include:

- **Genome variation:** In the breast cancer association consortium we identified 160 new breast cancer risk loci (published in *Nature Genetics*) and contributed to the fine-mapping of these loci (credible causal variants CCV).
- **Epigenomics of Breast Cancer:** Identification of epigenetically regulated cancer-driving pathways in breast cancer, and functional validation using CRISPR epigenetic engineering.
- **MicroRNAs in breast cancer:** Alterations in miRNA expression caused by neo-adjuvant treatment with chemotherapy and bevacizumab is associated with proliferation and response to therapy (published in *Molecular Oncology*). Identification of miRNAs as potential master regulators of the methylome in breast cancer.
- **Non-canonical transcriptomes:** Characterization of alternate exon usage and long non-coding RNA expression in breast cancer and connection to epigenetic profiles and patient outcomes.
- **Immune signaling:** Identification of patient groups based on immune signaling identifies a bad prognosis immune infiltration type independently of PAM50 classification and other known clinicopathological features (published in *Nature Communications*)

RECENT ACHIEVEMENTS

Publication activity. 22 publications in 2019, popular at <https://radiumlegat.no/Prosjekter/Vessela-Kristensen>,

Molecular Biology of Breast Cancer

Group Leader: Hege G. Russnes



“Exploring inter- and intra-tumor heterogeneity to improve molecular classification of breast cancer”

ABOUT

The group was founded January 2018 and have a total of 2 scientists, 1 postdoc, 1 research engineer, 1 nurse/MSc, 1 MD-PhD student and 1 MD student. In addition, 1 prof. emerita (A-L Børresen-Dale, UiO), 1 researcher (group leader A. Matheliér, NCMM), 1 oncologist (L. Ottestad) and 1 professor in bioinformatics (O. C. Lingjærde, UiO) are associated with the group (part-time).

Focus of the group is to reach a more fundamental understanding of the biological dynamics of breast cancer through high-throughput molecular analyses of DNA, mRNA, miRNA and protein, both at diagnosis and during disease progression.

As co-PI/partners in several clinic trials we perform “state of the art” analyses of patient material, both on bulk tumors, single cells and liquid biopsies, at various stages of the disease. The group is active in the IMI/EU funded project CancerID aiming at standardizing liquid biopsies for cancer diagnostics. Hege G. Russnes is also senior consultant at Dept. of Pathology, OUH where she is Head of “Section for experimental pathology and research support”, a lab developing and performing molecular diagnostics for clinical trials. She is also appointed “Young Associated Investigator” at NCMM (Centre for Molecular Medicine Norway).

AIMS

Our group aims at defining, refining and implementing molecular classification and biomarker analyses to improve stratification of cancer patients into treatment relevant groups. We are addressing the impact of inter- and intra-tumor heterogeneity as well as host response on disease progression, response to therapy and patient outcome.

PROJECTS

- LATE: Characterizing in time and space the metastatic process of hormone receptor positive breast cancer
- TREAC: Towards personalized treatment for patients with aggressive breast cancer
- Somatic Genetics of Breast Cancer, from single genes to whole genome sequencing
- SN - Sentinel Lymph Node in Breast Cancer- revealing the interaction between tumor subtypes and host immune response.
- CANCAN: CANcer specific Copynumber alteration Analysis
- CARMA: Copy Aberration Regional Mapping Analysis
- Liquid biopsies – optimizing methodologies for nucleic acid and single cell detection, capture and analysis

RECENT ACHIEVEMENTS

In 2019 we published 9 articles in peer-reviewed journals (in addition to 19 publications by our affiliated members). The LATE project received major funding from South-Eastern Norway Regional Health Authority (open project support) and from the Wellcome Sanger Institute (coll. MD/PhD Lucy Yates). The TREAC project, a collaboration with Elin Mortensen (University of Northern Norway), received a generous grant from Jeanette and Søren Bothners legat. In addition to being active partners in several clinical trials (ICON, ALICE, OPTIMA, I-BCT) we are coordinating the EMIT trial (A. Langerød). The EU/IMI collaboration CancerID finished December 2019, and we are active partners in the initiation of The European Liquid Biopsy Society (ELBS).





Department of Cancer Immunology

Headed by Johanna Olweus

ABOUT

The Department of Cancer Immunology (DCI) consists of five research groups (Olweus, Taskén, Sioud, Myklebust, Malmberg) and one project group (Kyte). Four DCI members are full professors at the University of Oslo. Groups at the DCI are partners in two K.G. Jebsen Centers (Cancer Immunotherapy and B-cell malignancies) and several EU-funded research programs in cancer immunotherapy. The groups provide complementary expertise in molecular and cellular immunology, including a broad experimental tool-box for antigen discovery and studies of immune cells at the single cell level. The aim is to decipher the molecular regulation of key cellular components of the innate and adaptive immune system, including dendritic cells (DC), B cells, T cells, regulatory T cells (Treg) and NK cells. The key driving force is to develop better tools for cancer diagnostics and new therapeutic strategies. The latter include investigator-initiated clinical trials to alleviate immune suppression and improve the use of check-point inhibition, and the design of gene-edited T- and NK cells for adoptive cell therapy.

PROJECTS

- Lymphocyte biology, by deciphering
 - ontogeny and function of B, T and NK cells
 - tumor heterogeneity (signaling and mutanome)
 - immune cell recognition elements (antigen discovery)
- Biomarkers, by profiling of
 - lymphocyte repertoires
 - the tumor and its microenvironment
 - T-cell receptors and humoral immunity

- Therapeutics, by
 - genetically engineered T and NK cells
 - immune priming with siRNA and antigen-targeting to DC
 - genetically engineered human antibodies and lytic peptides
 - cell therapy across HLA barriers to overcome immune tolerance
 - clinical trials using experimental immunotherapy
 - small molecules

RECENT ACHIEVEMENTS (2019)

- 27 publications; 70% with first/last authors from DCI (mean/median IF 10.1/6.4), including first and/or senior authorships in high-impact journals like Nature Communications, Nature Protocols, Cell Reports, Leukemia and Haematologica
- Awarded ERC Consolidator Grant (2 mill Euro) to Olweus for the project “Outsourcing Cancer Immunity to healthy donors” (only scientist in Life Science category in Norway)
- Launched Strat-Cell, a strategic research program (5 years) in cell therapy at Oslo University Hospital (Malmberg Director, Olweus/Kyte co-Directors)
- Filed four new DOFIs, three patent applications filed and two granted.
- 1 graduated Msc

CLINICAL TRIALS:

- Recruited first 300 patients to the ASAC trial that examines the effect of reversing prostaglandin E2-mediated inhibition of tumor immunity in patients with metastatic colorectal cancer by the end of 2019 (www.asac.no)

Experimental Immunotherapy

Group leader: Johanna Olweus



“Overcoming tolerance by T-cell based cancer immunotherapies”

ABOUT

The group counts 16 members (F/M 70/30); 1 full professor (JO), 6 postdocs, 5 PhD students and 3.5 engineers, and two associated clinicians. Four members have MD background. 13 members are recruited from abroad. The group is partner in K.G. Jebsen Center for Cancer Immunotherapy (2013-); (JCIT). Olweus is Director of JCIT, which was awarded maximal prolongation in 2016 (two years), till 2020

AIMS

To find new immunotherapeutic T-cell based strategies that overcome the major challenge of self-tolerance in cancer, and couple clinical trials with penetrating mechanistic analyses. Two principally distinct approaches are pursued in an interdisciplinary and translational program:
Strategy 1: Use of T cell-based alloreactivity to target self-antigens.
Strategy 2: Identification and targeting of cancer-specific neo-antigens.

PROJECTS

Strategy 1

- Identification of novel TCRs targeting self-antigens and epitope discovery

Strategy 2

- Identification of novel TCRs targeting neoantigens and epitope discovery
- Characterization of anti-tumor reactive T cells in biobanked material from patients responding to immunotherapy

RECENT ACHIEVEMENTS

Olweus was awarded an ERC Consolidator Grant (2 mill Euro over 5 yrs), as the only Norwegian scientist in the Life Science category. The group described a technology for identification of neoantigen specific T cells from healthy donors (Ali/Foldvari/Giannakopoulou et al, Nature Protocols, 2019), and wrote an invited perspectives article for Blood (Olweus and Lund-Johansen, 2019). The group continued their research collaboration with biotech company Kite Pharma (acquired by Gilead) on development of T-cell receptors to target cancer. Olweus was elected member of the Executive Board of CIMT, organizing the largest and most influential annual cancer immunotherapy meeting in Europe. She was invited speaker at numerous international conferences in 2019, including the CIMT Winter School in immunology (Innsbruck), Synthetic Systems Immunology (Ascona), MDS Symposium (Copenhagen), Cancer Research UK Symposium (London).

NK Cell Biology and Cell Therapy

Group leader: Karl-Johan Malmberg



“Towards the next generation NK cell therapy”

ABOUT

The Malmberg Lab in Oslo counts 18 members (F/M: 10/8); 1 full professor (KJM), 2 scientists, 1 project manager, 7 postdocs, 5 PhD students, 3 engineers. Malmberg is a visiting Professor at the Karolinska Institute (KI) and a partner in the K.G. Jebsen Center for Cancer Immunotherapy. Affiliated to the group is the Kyte Project group in Translational Cancer Immunotherapy led by Dr. Jon Amund Kyte, with 10 members: Project group leader (JAK), 3 researchers, two postdocs, 1 MD PhD student, 3 technicians. Dr. Kyte is also a senior consultant in oncology and Head of Dept. Clinical Cancer Research and is appointed Group Leader at DCI from 2020.

AIMS

The Malmberg group seeks to develop new strategies for cell-based immunotherapy based on insights into the functional regulation of natural killer (NK) cells. A combination of single-cell assays, including live cell imaging, high-dimensional immune profiling by mass cytometry, flow cytometry and RNA-seq is used to decipher the cellular and molecular mechanisms involved in calibration of effector functions in human NK cells. The Kyte group aims to develop improved cancer treatment based on two strategies for immunotherapy: 1) Development of cell therapy by use of novel tumor-targeting chimeric antigen receptors (CARs) and concepts for countering tumor tolerance 2) Development of optimized regimes for combining checkpoint inhibitors with chemotherapy or radiotherapy. The Kyte group has from 2017 started three clinical trials, in breast cancer (ALICE, ICON) and head and neck cancer (REPORT).

PROJECTS

- 1) Functional Diversification of human NK cell repertoires
- 2) Cell therapy with iPSC-derived NK cells

RECENT ACHIEVEMENTS

- Gained new insights into the functional regulation of human NK cells (Goodridge et al, Nature Communications 2019)
- Deciphered molecular pathways involved in NK cell homeostasis (Jacobs et al., J Immunology 2019, Pfefferle et al., Cell Reports 2019)
- Completed a one-year research sabbatical at UCSD resulting in new collaborations with world leading iPSC environments (Sætersmoen et al., Seminars in Immunopathology 2019).
- Launched Strat-Cell, a strategies research program (5 years) in cell therapy at Oslo University Hospital.
- Innovation: Initiated a 2-year collaborative agreement with EMD-Serrono to develop functional modulators of NK cells in vivo.

Lymphoma Biology

Group leader: June Helen Myklebust



“Understanding B-cell lymphoma biology to identify new therapeutic targets and treatment strategies”

ABOUT

The group counts 13 members with research background in medicine, biology and biotechnology, and includes 1 professor, 1 associate professor, 1 scientist, 7 postdocs (including visiting postdoc at MGH/Broad Institute, Boston), 2 PhD students and 1 technician. Five members have MD background and four members are recruited from abroad. The group is part of the KG Jebsen Centre for B-cell malignancies.

AIMS:

The group performs translational studies in B-cell lymphoma to define tumor clonal evolution, cancer driver genes, actionable targets and new targets for immunotherapy.

PROJECTS

We use single-cell technologies to characterize tumor cells and intratumor immune cells in patient biopsies. This includes high-dimensional flow cytometry, mass cytometry and mass cytometry imaging (Hyperion), and single-cell RNA sequencing. We also use CRISPR/Cas9 genomic editing combined with immunological assays to characterize potential cancer driver genes, and have established patient-derived xenograft (PDX) mouse models for pre-clinical drug testing. Ongoing projects are:

- Clonal evolution and recurrent mutations associated with therapy relapse
- Functional characterization of recurrent driver mutations
- Novel targets for immune checkpoint blockade
- Cancer sensitivity drug screen and preclinical testing
- Clinical register studies

RECENT ACHIEVEMENTS

The group published 7 original papers, 1 commentary and 1 news&views (Nature) in 2019: Our in-depth analysis of the tumor microenvironment identified TIGIT and PD-1 as relevant targets for co-checkpoint blockade in B-cell lymphoma (Josefsson, Cancer Immunol Res). Our collaborative effort with Cell therapy has led to preclinical development of CD37 CAR T cell therapy (Köksal, Blood Adv and patent filed). We developed the clinical-genetic prognostic tool BTK-FLIPI to identify follicular lymphoma at high risk for adverse disease (Steen, Haematologica), and contributed to convergence of risk prediction models (Silva, Haematologica). Several members had oral presentations at international meetings; the most prestigious being our collaborative project with lymphoma clinicians, the Hovig group and Genomics core facility to map tumor clonal evolution in follicular lymphoma, selected for oral presentation at the American Society of Hematology (Bai, ASH 2019).

Immunomodulation and Targeted Therapies

Group leader: Mouldy Sioud



“Innovative approaches for cancer therapy”

ABOUT

The group has 8 members, including 1 postdoc, 2 research engineers, 1 PhD student, 3 master students, and the group leader (DEA Pharm, PhD) with formal research experience in immunology, molecular and cell biology, microbiology, and medicine. Sioud is a visiting professor at University of Tunis. To date, the group has published 197 peer-reviewed original articles (mean IF = 5.865) and reviews, with 1st and/or last authorship on 85% of the papers.

Traditional cancer treatments like radiation and chemotherapy have major side effects because they not only affect the tumors, but also the healthy parts of the body. Hence, there is a need for improvement. The group is mainly focused on the development of therapeutic antibodies, lytic peptides, and small RNA modulators to target and kill cancer cells or counteract immune suppression. With respect to clinical translation, we developed the first RNA interference-modified dendritic cell cancer vaccine that is now available to patients under compassionate use (Sioud 2019, *Cancers*, IF 6.2).

AIMS

The principal aim is to develop antibodies, antibody derivatives (e.g., antibody-photosensitizer conjugates, chimeric antigen receptors, bispecific antibodies), and lytic peptides for use in cancer immunotherapy.

PROJECTS

- Profiling the cell surface proteome using high-throughput phage display technologies
- Development of therapeutic human antibodies and lytic peptides
- Fingerprinting immune responses in cancer patients
- RNA interference and CRISPR technologies as immune modulators

RECENT ACHIEVEMENTS

- Reported the first lytic peptide targeting tumor macrophages and leukemia cells (Sioud et al. *Cancers* 2019)
- Developed new antibody-photosensitizer conjugates to target and kill cancer cells (Gebretensaie et al. manuscript in preparation)
- Developed new human antibodies to modulate immune cell functions
- Described a new mechanism of action of extracorporeal photopheresis immunotherapy involving IDO+ tolerogenic dendritic cells (Sagar et al. *Cancers* 2020).
- Three publications
- Sioud edited a new book on RNA interference and CRISPR technologies (Springer-Nature: <https://www.springer.com/gp/book/9781071602898>).
- The group obtained funding from South-Eastern Norway Regional Health Authority and Radforsk

Cell Signalling and Immune Regulation

Group leader: Kjetil Taskén



«We want to perturb tumor immune evasion mechanisms to boost anti-tumor immunity»

ABOUT

In 2019 the group counted 18 members (F/M: 10/8), 1 full professor (KT), 1 senior consultant, 2 researchers, 8 postdocs, 1 PhD student, 1 MD/PhD student, 2 technicians and 2 M.Sc. students. The group is part of K.G. Jebsen Centres for Immunotherapy and B Cell Malignancies.

AIMS

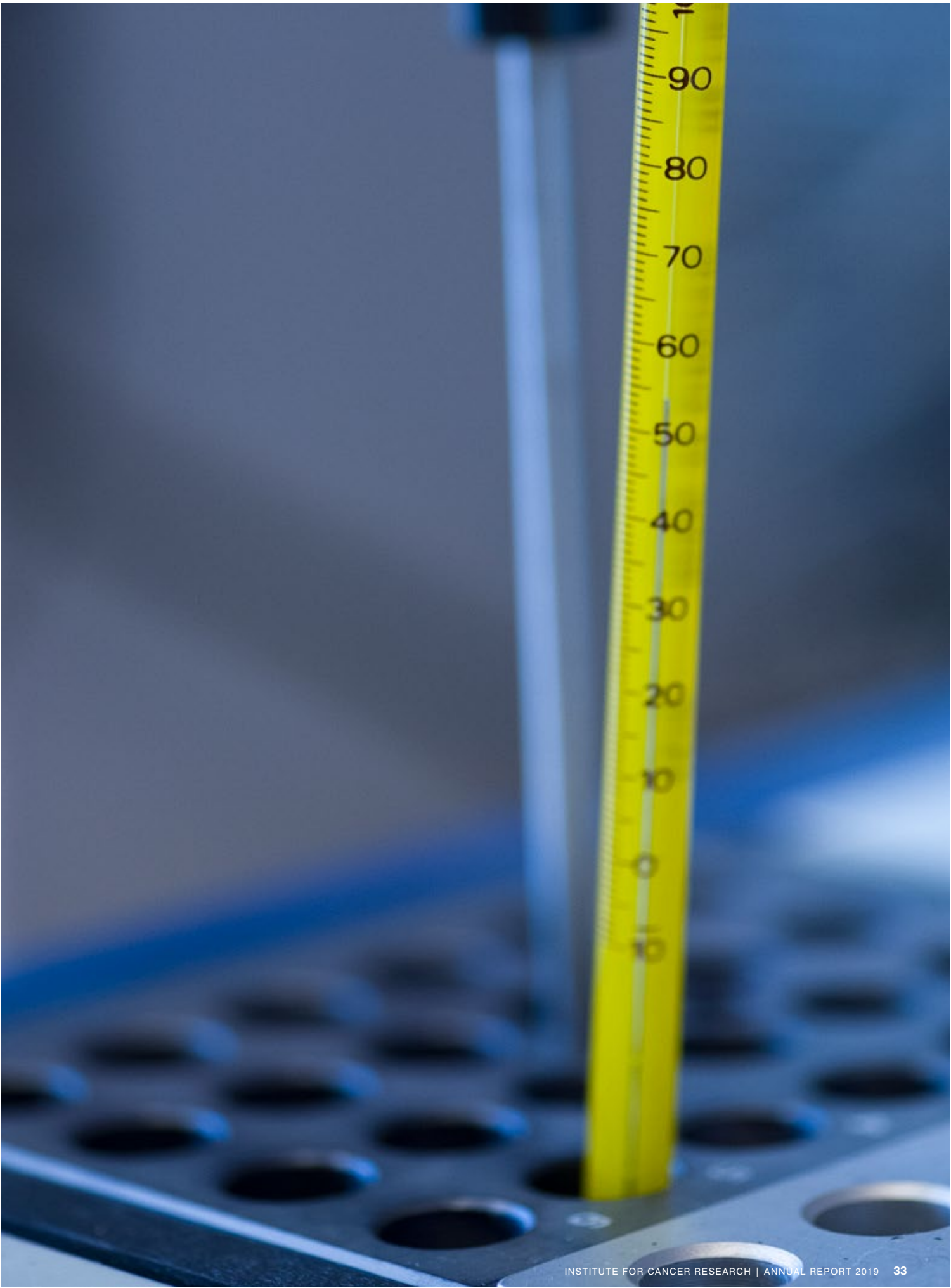
The Taskén group aims to understand intracellular signalling networks, the anchoring and localization of signaling complexes through scaffold proteins, how these signalling networks mediate physiological and pathophysiological processes and can be perturbed using drug-like small molecules. We aim to understand tumor immune evasion mechanisms, and how we can interfere to boost anti-tumor immunity. We proceed with cancer drug sensitivity screening (CDSS) to explore individual drug responsiveness and resistance patterns in patient cancer cells and aiming to develop models to assist individualised clinical decisions in precision medicine in oncology and haematology.

PROJECTS

- T cell function in cancer and immune-related diseases
- Identification of regulatory T cell targets that can be perturbed to reverse tumor immune suppression
- Role of Prostaglandin E2, cAMP and AKAPs in signaling and regulation of T cell function
- Targeting the cAMP signalling pathway for cancer immunotherapy
- CDSS in chronic lymphocytic leukemia and multiple myeloma, understanding drug synergies and predicting effective drug combinations in individual patients (RCN-Digital Life PINPOINT project)
- Acetyl salicylic acid clinical intervention study in metastatic colorectal cancer (ASAC)

RECENT ACHIEVEMENTS

Publication highlights include studies on CLL patients in our precision medicine programme where we predict clinical dosing of combinatorial treatments based on biomarker studies (Leukemia), on regulatory T cells and tumor immune suppression by idelalisib (J. Immunol.) and on signalling complexes orchestrated by CD28 and CD2 (J. Immunol.). Furthermore, in co-authored papers we have contributed to understanding metabolic regulatory programmes for aerobic glycolysis in T cells (Nature) and TLR8 signaling in T cells (Nature Commun., in press). 1 DOFI was submitted, 1 patent was filed and 1 granted. One M.Sc. student graduated.





“Uncovering
the cellular
basis of cancer
development”

Department of Molecular Cell Biology

Headed by Harald Stenmark

The department has a staff of 80 (including 11 MSc students and trainees) of 23 nationalities and hosts 4 research groups (Enserink, Rusten, Sandvig and Stenmark), 9 project groups, and a departmental service unit. Research in the department comprises various aspects of cancer relevant cell biology, including membrane traffic, cell signaling, cell metabolism and cell division. In addition, the department carries out biotechnological research on nanoparticles and translational research on leukemia drug sensitivity, cancer-related fungal infections, and cancer derived exosome biomarkers.

A common goal for researchers in the department is to uncover novel molecular mechanisms that control cellular functions related to cancer. The department's research strategy is to combine molecular cell biology with biochemistry, genetics, drug screening and advanced imaging in order to address relevant scientific questions. Recent key achievements from the department's scientists include studies on autophagy

and tumour growth, growth factor signalling and intracellular transport, molecular mechanisms of cell division, exosome secretion and biomarkers for prostate cancer. In general, the department's groups have been successful in obtaining national and international external funding.

The groups of Harald Stenmark, Jorrit Enserink and Tor Erik Rusten are members of a Norwegian Centre of Excellence, CanCell, headed by Harald Stenmark. Harald Stenmark also heads the Norwegian Advanced Light Microscopy Infrastructure Network, NALMIN, and the Chinese-Norwegian Partnership for Education and Research in Cancer Cell Biology, ChiNoCell. Stenmark's group is member of a Convergence Environment under UiO Life Science, called “Programmable cell-like compartments”. Jorrit Enserink participates in a project under Norwegian Centre for Digital Life, “Pipeline for individually tailoring new treatments in hematological cancers”. Enserink's group also participates in the EU Horizon 2020 project, RESCUER.



Cellular Membrane Dynamics

Group leader: Harald Stenmark



“Understanding how remodelling of cellular membranes contributes to cancer”

ABOUT

The group studies the dynamics of cellular membranes with the aim of understanding their relevance to cancer. Cellular membrane dynamics processes studied by the group include endocytosis, autophagy, and cell division. The group has 34 members from 12 nations and is headed by Harald Stenmark (group leader) and Camilla Raiborg (assistant group leader). Its 5 project groups are led by Andreas Brech, Kaisa Haglund, Camilla Raiborg, Kay O. Schink, and Antoni Wiedlocha.

AIMS

The principal aim of the group is to understand how membrane dynamics contribute to tumour suppression and how their dysfunction may promote cancer development.

PROJECTS

- Phosphoinositides in regulation of membrane dynamics
- Mechanisms of autophagy and lipid droplet biogenesis, and their role in cell metabolism
- Centrosome dynamics in cancer
- Cytokinesis in development and carcinogenesis
- The β -catenin destruction complex in physiology and cancer
- Membrane dynamics in promotion of genome integrity
- Mechanisms of cancer cell invasion

RECENT ACHIEVEMENTS

- New mechanism of tumour suppression – intracellular retention of matrix metalloproteinases (Sneeggen et al., Nature Communications 2019). Awarded publication prize from Oslo University Hospital.
- ESCRT proteins promote autophagy by mediating sealing of newly formed autophagosomes (Zhen et al., Autophagy, 2019).
- Novel mechanism for recruitment of the abscission machinery to the midbody during cytokinesis – (Lie-Jensen et al., Current Biology, 2019). Dedicated a commentary article in Current Biology.
- Comprehensive review on ESCRT proteins in sealing and scission of cellular membranes (Vietri et al., Nature Reviews Molecular Cell Biology, 2019).
- Major grants in 2019 to Kay O. Schink, Harald Stenmark, and Viola Nähse.
- Marte Sneeggen obtained her PhD in November 2019.
- Harald Stenmark was invited speaker at international conferences in Okinawa, Paris, Trondheim, Glasgow, Beijing, Edinburgh, Essen and Strasbourg in 2019.
- Nina Marie Pedersen and Lene Malerød were selected for oral presentations at the international conference on “Cell Signaling and Intracellular Trafficking in Cancer Biology” in Turin, and Yan Zhen and Andreas Brech received prizes for best posters at the 9th International Symposium on Autophagy in Taipei.
- New collaboration on cancer cell invasion, InvaCell, with the group of Philippe Chavrier at the Curie Institute, Paris. InvaCell is generously sponsored by Trond Paulsen.
- Members of the group published 13 original papers and 3 reviews in 2019.

Cancer Molecular Medicine

Group leader Jorrit Enserink



“Identifying weak points in the molecular networks that drive cancer”

ABOUT

The group currently consists of one group leader, two project group leaders, seven post-docs, one clinician in a 20% post-doc position, two PhD students, four MSc students and one Erasmus student. A large fraction of the group consists of scientists from abroad, including the Netherlands, Austria, Spain, Colombia and the UK. The group is mainly focused on basic cancer biology and personalized medicine, using a wide range of models such as budding yeast and fruit flies, human and mouse cell lines, and primary human cancer samples.

AIMS

Research is aimed at understanding cancer cell biology, with the ultimate goal of developing precise cancer treatments. A major focus is on hematopoietic cancers, including –but not limited to– Acute Myeloid Leukemia (AML). A second research theme is to better understand cellular responses to sudden environmental changes.

PROJECTS

- High-throughput drug combination screens to identify drug synergies and to reveal correlations between driver mutations and drug sensitivity profiles

- Development of novel immune therapy for AML
- Modeling of leukemia in fruit flies to better understand the genetic networks that drive AML
- Genome-wide CRISPR-Cas9 screens in leukemic cells to identify novel mechanisms of resistance to tyrosine kinase inhibitors
- Cellular responses to nutrient stress; particularly the identification of the upstream pathways that control the dynamics of autophagy

ACHIEVEMENTS

- Five MSc degrees were completed and the group published three articles.
- Funding obtained: A research grant from The Norwegian Cancer Society to Jorrit Enserink and a Young Research Talent grant from The Research Council of Norway to Dr. Helene Knævelsrud.
- The group is also part of the RESCUER project headed by Dr. Vessela Kristensen, which was awarded by the EU Horizon 2020 scheme in 2019.

Tumor-host Biology

Group leader Tor Erik Rusten



“Tumor-host interactions during cancer progression”

ABOUT

The research group counts 12 members representing 7 nationalities in 2019 (Australia, India, Iran, Hungary, France, Spain and Norway): 1 group leader, 1 scientist, 5 postdocs and 1 PhD student, 1 technician and 1 master students and 2 Erasmus exchange students. Cancer can be regarded as organ development derailed. Our overarching interest is to understand the mechanisms by which tumor and host cells mutually engage each other in reciprocal interactions that promote cancer progression. These interactions occur locally in the tumor microenvironment, but also systemically causing organ dysfunction such as in cancer cachexia - the catastrophic wasting of muscle and adipose tissue. We believe that studying these processes can uncover new ways to intercept carcinogenesis. To mechanistically probe how tumor cells and non- tumor cells and organs communicate to foster tumor growth and cause cancer cachexia we develop novel genetic tools in *Drosophila*. These tools will allow us to selectively and independently manipulate tumor and either tumor microenvironment or somatic organs in vivo. We investigate the molecular genetic mechanisms and cell biology of genetically defined experimental tumor-host interactions using a mix of human cell culture and the fruit fly *Drosophila melanogaster*, as an animal model system. In this work we employ a wide array of techniques and collaborate with experts in cell biology, genetics, imaging, electron microscopy, tumor biology, metabolism, bioinformatics and clinical cancer cachexia in order to survey, measure and mechanistically understand these complex aspects of cancer biology.

AIMS

The principal aim of the group is to understand tumor-host interactions that facilitate cancer progression in order to uncover novel ways to intercept cancer.

PROJECTS

- Oncogene-induced epithelial disintegration and invasion.
- Tumor-microenvironment interactions and growth support.
- Mechanisms of cancer cachexia.
- Roles of autophagy in metabolic reprogramming, nutrient mobilization and breakdown of muscle and adipose tissue during cancer cachexia.

RECENT ACHIEVEMENTS

- EMBO long-term postdoctoral fellowship was awarded to Dr. Swarupa Panda to work on “Mechanisms of non-autonomous support of tumor cell growth”
- Tor Erik Rusten was invited at 5 international research conferences (including a Gordon Research Conference) in Heidelberg, Lisbon, Utrecht, Barcelona and Santa Fe in 2019, one of these as keynote speaker
- In 2019 one MSc was completed and the group published one review, one preview and contributed to two original research articles

Intracellular Transport

Group leader: Kirsten Sandvig



“All the way from basic research to translation”

ABOUT

Sandvig's group, counting 15 members, is interested in the mechanisms of endocytosis, intracellular transport and secretion. Uptake of receptors and ligands and correct intracellular sorting of endocytosed molecules are crucial for maintenance of a normal differentiated phenotype. In some studies we are using protein toxins such as ricin and Shiga toxin, which are well established as markers for studies of membrane traffic, and which can be used as agents in cancer diagnosis and therapy. Our expertise is also applied to investigate uptake of nanoparticles for drug delivery, and since 2013 we have had a large grant from the Norwegian Research Council to build national competence in nanomedicine. This project, “Biodegradable nanoparticles in cancer diagnosis and therapy”, headed by Sandvig, involved collaboration with ten other Norwegian groups working in academia, research institutes, hospitals and industry, and was running until March 2019. The Sandvig group has also been involved in an INNO INDIGO granted project, which started April 2016 and ended September 15th, 2019. INNO INDIGO is an innovation-driven initiative for the development and integration of Indian and European research. Our nanoparticle research has, based on results from these projects, obtained further support from the Norwegian Cancer Society for 2020-2023. We also characterize exosomes from prostate cancer cells and patients with the goal of detecting lipid, RNA and protein biomarkers. Our research spans all the way from basic to translational medicine, including innovation. The work published by the Sandvig group is well cited, and Sandvig's H-index is now 73 (~340 publications). The group has extensive national and international collaborations.

AIMS

The projects aim at increasing our knowledge about intracellular transport, nanoparticles for drug delivery, and exosomal biomarkers in order to provide a rational basis for diagnosis, treatment and prevention of cancer.

PROJECTS

- Characterization of intracellular transport
- Biodegradable nanoparticles in cancer diagnosis and therapy
- Exosomes and prostate cancer

RECENT ACHIEVEMENTS

- Mechanistic studies of different types of endocytosis and intracellular transport. Our competence concerning cellular membranes and lipid species made us publish a perspective article (Skotland et al., Nature Communications, 2019) about the role of specific lipid species and what should be done in the future to increase our understanding of cellular membranes.
- We have performed further studies of exosome biogenesis and release, as well as of biomarkers for prostate cancer. Investigations of cytotoxic effects of different types of nanoparticles with and without drugs were performed both in vitro and in vivo.
- In 2019 the group published on the different topics; 10 articles in different journals.
- Kirsten Sandvig obtained a research grant of 8 MNOK from the Norwegian Cancer Society on a project that will investigate the efficacy of drug-containing nanoparticles in breast and colorectal cancers.



”Biological discoveries for improved precision cancer medicine”



Department of Molecular Oncology

Headed by Ragnhild A. Lothe

Our main research programs are devoted to colorectal cancer and prostate cancer, but with a longstanding project portfolio also on other solid tumor types. Our expertise in biomedical research spans across several disciplines, and we have a broad range of advanced technologies and analytical tools established in the lab. The department hosts three research groups and three project leaders, in total ~10 % of the total staff at the Institute (full time equivalent). Six of the senior scientists are affiliated with the University of Oslo as professors (three), associate professor, or lecturers (two). We are devoted to teaching and supervision, and six academic degrees (4 MSc and 2 PhD) with main supervisors from our Department were successfully completed last year.

Our research groups are current partners of the K. G. Jebsen Colorectal Cancer Research Centre, an OUH Strategic Research Area, and several national and international scientific networks. The latter includes national multicenter studies on colorectal cancer and bladder cancer, European cooperation studies on colorectal cancer (COST action), a European multicenter study on the rare tumor type malignant peripheral nerve sheath tumor, the European network for study on cholangiocarcinoma, and the Global Testicular Cancer Consortium.

A few selected results from last year:
We showed that multifocal primary prostate cancers have an exceptional degree of intra-patient genomic heterogeneity (Løv et al., Eur Urol), and this challenges the usefulness of known molecular classifiers (Carm et al., Sci Rep). Furthermore, we published a comprehensive review on biomarker-guided therapy for colorectal cancer (Sveen et al., Nature Rev Clin Oncol), and demonstrated the clinical impact of tumor microenvironment markers relative to cancer genomic markers in patients with colorectal cancer (Dienstmann et al., Ann Oncol; Berg et al., Oncogene). A national multicentre study and innovation project to monitor bladder cancer patients with cancer-specific epigenetic markers in urine samples documented strong discovery and validation results (unpublished), and externally financed by major grants from the Norwegian Research Council and National Health Regions (KLINBEFORSK)

Our main research goals for the next three to five years are three-fold, (i) to decipher spatio-temporal tumor heterogeneity and its clinical relevance in colorectal cancer and prostate cancer, (ii) to monitor minimal residual disease, early recurrence and clonal evolution by analyses of repeated liquid biopsies and tumor samples, and (iii) to predict treatment responses by combined molecular profiling and ex vivo drug screening of patient-derived organoids in clinical trials and translational studies.

Genetics

Group leader: Ragnhild A. Lothe



“Molecular tumor heterogeneity and personalized pharmacogenomics - the basis for next generation clinical trials”

ABOUT

Our main research program involves translational studies of primary and metastatic colorectal cancers (CRC), using genomics, drug screening, digital pathology and functional analyses. The group has 23 members and includes two project groups in Cell signaling and Computational oncology.

AIM

Our overarching goal is to translate novel biomedical knowledge into improved patient stratification and treatment of CRC.

PROJECTS

- Prognostic and predictive biomarkers
- Modeling tumor heterogeneity and clonal evolution
- Pharmacogenomics of solid tumors using patient-derived organoids (PDO)
- E3 ubiquitin ligases in intercellular communication and carcinogenesis

RECENT ACHIEVEMENTS

In 2019 we published 20 papers, including in Ann Oncol (2), Oncogene (2), PNAS, and Nature Rev Clin Oncol. Kaja Christine Graue Berg defended her PhD at the Faculty of Medicine, U of Oslo. Two early career scientists were rewarded for their excellence, Anita Sveen received “Early Career Award” from OUH, and a major grant from the Norwegian Cancer Society; Peter W. Eide received the “Young researcher award” from Onkologisk Forum (Annual National Oncology meeting).

We published a comprehensive review on biomarker-guided therapy for CRC (Sveen et al., Nature Rev Clin Oncol 2019), showed that splice variants of KRAS has prognostic relevance beyond its mutation status (Eilertsen et al., Int J Ca 2019) and we concluded that the long noncoding RNA MIR31HG is a bona fide prognostic marker in CRC (Eide et al., Int J Ca 2019). We showed that TP53 mutations may modulate the immune microenvironment in a particularly immunogenic subgroup of CRCs, inferring a poor patient survival (Smeby et al., ESMO Open 2019). In multicentre studies we confirm that tumor microenvironment markers are key determinants for risk of relapse in early-stage CRC (Dienstmann et al., Ann Oncol, 2019; Glaire et al., Brit J Ca 2019).

Much effort is put into the pharmacogenomics project: multiple ex vivo cultures from CRC liver metastases have been successfully established as PDOs, and screened for sensitivity to 40 anticancer agents. Intra-patient inter-metastatic pharmacological heterogeneity is not pronounced and variation in drug sensitivities is reflected at the transcriptomic level, suggesting great potential to develop gene expression-based predictive signatures to guide experimental therapies (paper in revision). A protocol for a clinical trial implementing such personalized pharmacological models is currently in development.

Epigenetics

Group leader: Guro E. Lind



“Epigenomics – reversible changes in cancer and a source for clinical biomarkers”

ABOUT

In the group of Epigenetics we are studying DNA methylation alterations by integrating genome wide methylation profiling (sequencing and arrays), with detailed candidate gene characterization. The research is performed across cancer types, with a main focus on gastrointestinal and urological cancer. In 2019 the group counted eleven members: 3 postdocs/scientists, 2 PhD students, 3 engineers, one study nurse, one MSc student and the group leader, and includes a project group.

AIMS

- 1) To identify and develop epigenetic biomarkers with clinical impact, including markers for early detection and monitoring of cancer.
- 2) To explore the inter- and intra-tumor diversity of epigenetic aberrations in cancer, and effect on patient outcome.

PROJECTS

- Epigenetic subclassification and prognostic markers for colorectal cancer
- Methylome-based early detection and monitoring of bladder and gastrointestinal cancer
- Epigenetic heterogeneity in gastrointestinal cancers

RECENT ACHIEVEMENTS

During 2019, the group has continued the development of a urine-based test for monitoring of bladder cancer patients. A set of biomarkers, identified in-house using methylome sequencing, demonstrated high accuracy in a test series and was recently validated using a blinded prospective international patient series (manuscript). A cohort of 50 post-surgery individuals is being followed at Aker for two years, in order to compare the urine test to the current gold standard method (cystoscopy). Full follow up time will be reached in 2020. Preliminary data indicate that the urine test enables earlier detection of bladder cancer recurrence.

The data have laid the foundations of a national multi-center study aiming at demonstrating the clinical utility of the urine test. Hospitals from all health regions are participating, and the first patient was recruited in December. All together 500 patients will be followed for a two-year period. The study is funded through “klinisk behandlingsforskning” and received additional funding from the Research council in 2019.

In 2019 Marine Jeanmougin was appointed leader of a project group with expertise in biostatistics. One PhD degree (Heidi Pharo) and one MSc degree (Ida Marie Børresen) were completed from the group in 2019.

Genome Biology

Group leader: Rolf I. Skotheim



“Molecular heterogeneity of prostate cancer - Improved precision diagnostics”

ABOUT

The research group studies how genomes and transcriptomes are altered in cancer cells by using both computational and wet-lab based approaches. Mutation analyses in cancer benefit from knowing which genes and variants that are expressed, and the group has specialized in RNA-level analyses. Further, most projects focus on prostate cancer. An interdisciplinary set of expertise is beneficial for genome-scale cancer research, and the personnel have their education across the disciplines biology, informatics, and medicine. Through 2019, the group consisted of twelve members.

AIM

To improve the diagnosis and management of cancer by utilizing genome technologies.

PROJECTS

- Interfocal heterogeneity of prostate cancer
- RNA variation caused by aberrant splicing and as a source of cancer biomarkers
- Fusion gene identification and characterization

RECENT ACHIEVEMENTS

During 2019, the group continued the development of a large prostate cancer research program. The first large in-depth genomic heterogeneity analysis of primary

prostate cancer was reported (Løvfi et al., Eur. Urol, 2019). Here, whole-exome sequencing of 89 tumor foci from 41 patients revealed that different tumor foci within the same patient only exceptionally share any somatic mutations. A follow-up study pointed out that existing molecular signatures are useless in the clinic if inter-focal heterogeneity is not considered (Carm et al., Sci. Rep., 2019). This is important for how genetics can inform on treatment decisions as information from all tumor foci is necessary to conclude about the cancer. The group continued to develop bioinformatics pipelines for high-throughput sequencing of DNA and RNA. A paper in press describes a tool for sensitive fusion gene detection and with biomedical data from testicular cancer (Zhao et al., NAR Genomics and Bioinformatics). In addition to these three articles, the group contributed to four publications from internal and external collaborations.

In 2019, three MSc degrees were achieved. Karina Borlaug got her degree at Dept. Informatics, whereas Susanne Kidd and Linn Olsen got theirs from Dept. Molecular Biosciences.

Important research funding was obtained from the Norwegian Cancer Society and South-Eastern Norway Regional Health Authority.





“Our goal is to develop new predictive methods and treatment strategies for improved radiation therapy”

Department of Radiation Biology

Headed by Kristian Berg



The Department has more than 60 employees organized in 4 research groups and 6 project groups. The research at the department is focused on the biological responses to ionizing and non-ionizing radiation, including γ -radiation, radiation from radionuclides, ultraviolet radiation, visible light as well as proton therapy. Our department is actively engaged in revealing the mechanisms involved in DNA repair, cell-cycle regulation, genomic instability and immune responses after radiation, the impact of hypoxia on the radiation response and predictive markers for the outcome of radiotherapy of cancer patients. The department is also involved in delivering radionuclides to cancer tissue. Another research area is the use of visible light to activate photosensitive compounds, thereby generating reactive oxygen species, which are used for cancer detection and therapy. In that respect a technology has been developed, photochemical internalization (PCI), which enables site-directed intracellular delivery of anticancer therapeutics.

OUR GOALS ARE

- To understand the molecular and physiological mechanisms behind tumour response to radiation, and to develop predictive methods and treatment strategies;
- To utilize acquired knowledge to establish new radiation-based treatment regimens with improved specificity and efficacy towards cancer cells;
- To develop treatment strategies utilizing non-ionizing radiation combined with photosensitizing agents, either to induce tumour cell kill directly or via internalization of therapeutic or sensitizing agents.

KEY ACHIEVEMENTS THE LAST 3-4 YEARS INCLUDE

- Pancreatic carcinoma xenografts treated with sunitinib show less abnormal microvessels but larger hypoxic regions after treatment than before treatment;
- Cervical cancer patients with tumors showing high hypoxic fraction in combination with high interstitial fluid pressure have particularly poor prognosis with a 5-year survival rate of only 13 %;
- A novel mechanism of activation of the DNA damage kinase;ATR (Ataxia Telangiectasia and Rad3-related) was identified;
- PCI has been found to efficiently enhance antigen presentation during anti-cancer vaccination.;
- A production unit for biomolecular therapeutics has been established. Several new recombinant targeted protein toxins have been developed and are under preclinical evaluation;
- Documented for the first time that protons can activate photosensitizers used in photodynamic therapy (PDT) to mimic responses as in PDT;
- A new method to image hypoxia in prostate cancer based on integration of images reflecting oxygen consumption and supply has been developed;
- We gained important new knowledge about the regulation of translation in response to cellular stress as well as about the function of the stress-response kinase GCN2 in human cells;
- Combinatorial drug partners were identified that overcome resistance to CD37-targeted radioimmunotherapy in B cell lymphoma;
- Described a key mechanism required for formation of Hh-signaling competent cilia that is potentially exploitable as treatment target for Hh-dependent cancers.

Photochemical Internalization

Group leader: Kristian Berg



“Our goal is to develop and optimize the PCI technology for treatment of solid cancers”

ABOUT

Group members: 21, including 5 researchers, 5 postdocs, 3 PhD students, 6 technical positions and 2 MSc students, including the project groups of Asta Juzeniene, Pål Kristian Selbo, Anette Weyergang and Theodossis Theodossiou.

Project *Photochemical Internalization* (group leader Berg, project leaders Selbo, Weyergang and Theodossiou): Endocytic entrapment is a hurdle for therapeutic utilization of macromolecular therapeutics with intracellular targets. Photochemical internalisation (PCI) is a technology for cytosolic release of therapeutic macromolecules subjected to endocytosis. PCI is invented and developed in our research group and is currently evaluated in clinical trials.

Project *Targeted alpha therapy* (project leader Juzeniene): Metastases are the primary cause of death in cancer patients. Targeted alpha-particle therapy is a promising treatment for eradicating *micrometastases*.

AIMS

Project *Photochemical internalization*: The main aim of our group is to develop and optimize the PCI technology for treatment of solid cancers with a curative endpoint.

Project *Radionuclide therapy*: The main goal is to develop a novel technology with potentially broad therapeutic applications for cancer micrometastases by means of dual targeted alpha particle radiation.

PROJECTS

- Design and development of recombinant immunotoxins for activation by PCI
- Light-controlled delivery of cancer immunotherapeutics including PCI of 1) immunotoxins targeting cancer stem cells (CSCs) and 2) CSC-derived vaccines.
- Utilize new vehicles for targeted delivery of small molecular drugs to endocytic vesicles for activation by PCI
- Evaluation of the vasculature as a target for PCI and further utilize these effects to reach a curative endpoint
- Using mitochondria-powered chemiluminescence to non-invasively treat inaccessible tumours
- Utilizing other radiation sources to induce PCI effects
- Targeted alpha radionuclide therapy for bone and visceral metastases of osteosarcoma, prostate and breast cancer

RECENT ACHIEVEMENTS

- Documented for the first time that protons can activate photosensitizers used in photodynamic therapy (PDT) to mimic responses as in PDT; Published in *Nature Comm.*
- The new FET-OPEN project FRINGE (Co-ordinator: Theodossiou) was launched in 2019,
- New grants in 2019: PhD stipend from HSE (Selbo); Project grant from Radforsk (Weyergang)
- Theodossis A. Theodossiou awarded the prize researcher of the year at the Institute for Cancer Research
- The EU Innovative Science 2019 awarded to the Lumiblast project (Theodossiou and Berg)
- No. of papers in 2019: 8
- MSc thesis: 2

Clinical Radiation Biology

Group leader: Heidi Lyng



“Our goal is to discover biomarkers and molecular targets for combination therapies with radiation”

ABOUT

Group members: 10, including one researcher, four postdocs, three PhD students, and two technicians.

The focus is to develop molecular biomarkers that can identify patients at risk of radiotherapy failure and to find targets for therapeutic intervention in a combined radiotherapy regimen. The research projects are run in close collaboration with medical doctors and physicists at the hospital.

The work is based on tumor biopsies and blood samples collected at diagnosis or during therapy. Whole genome methodology is used, including microarray and sequencing techniques, which provides new insight into the disease and enables identification of novel biomarkers and drug targets. We also explore whether functional tumor images (MRI/PET) can aid the translation of the molecular findings into radiotherapy practice.

AIMS

- Understand molecular mechanisms behind radioresistance of cervical and prostate cancer
- Develop biomarkers and find drug targets for personalized radiotherapy of patients with these diseases

PROJECTS

- Genetic alterations and chemo-radioresistance in cervical cancer
- Molecular hypoxia biomarkers in cervical and prostate cancer
- Combined molecular and imaging biomarkers in cervical and prostate cancer

RECENT ACHIEVEMENTS

In 2019, the group published 4 articles. Major findings were:

- Discovery of a miRNA candidate biomarker for hypoxia related treatment resistance in cervical.
- Identification and exploration of an aggressive cervical phenotype with elevated oxidative phosphorylation and proliferation activity.

Radiation Biology and Tumor Physiology

Group leader: Einar K. Rofstad



“Our goal is to identify strategies for personalized radiation therapy of cancer”

ABOUT

Group members: 7, including 2 researchers, 4 postdocs, 2 PhD students, and 1 technician.

The focus of the group is to improve the outcome of radiation therapy of cancer. Poor outcome is a consequence of radiation resistance and elevated metastatic propensity of the primary tumor, and our research is based on the hypothesis that poor outcome is caused primarily by an abnormal physicochemical tumor microenvironment. Xenografts of human cervical carcinoma, pancreatic carcinoma, and malignant melanoma are used as preclinical tumor models. In addition to standard molecular biology methods, important methods include intravital microscopy, MRI, and probe measurements of physicochemical parameters.

AIMS

To reach the primary goal, our research is divided into two arms with the following aims:

- To develop MRI-based methods that can provide information on the physicochemical microenvironment, metastatic propensity, and radiocurability of tumors
- To develop treatment strategies for normalizing the physicochemical microenvironment, decreasing the metastatic propensity, and enhancing the radiocurability of tumors

PROJECTS

- Clinical and preclinical MRI of the physicochemical microenvironment of tumors
- Treatment strategies targeting the physicochemical microenvironment of tumors

RECENT ACHIEVEMENTS

Anette Hauge defended her Doctor Philosophiae (Dr Philos) thesis at the Faculty of Mathematics and Natural Sciences, University of Oslo in 2019. The thesis (The Tumor Microenvironment and Its Assessment with DCE-MRI and DW-MRI) consists of 4 first-author and 3 second-author papers.

Our group developed novel MRI methods for characterizing the physicochemical microenvironment of tumors in 2019.

Important scientific findings in 2019:

- Highly elevated tumor interstitial fluid pressure is a stronger biomarker of the outcome of locally advanced carcinoma of the uterine cervix than high fraction of hypoxic tumor tissue.
- Antiangiogenic treatment of cervical carcinoma with bevacizumab may cause increased tumor hypoxia, owing to treatment-induced vessel pruning.

Radiation Biology and DNA Damage Signaling

Group leader: Randi Syljuåsen



“Our goal is to obtain new knowledge about cellular responses to radiation and utilize it to improve cancer therapy”

ABOUT

Group members
14.3 including 5 researchers, 1 postdoc ,5 PhD students and 3.3 technicians.

Theme

In radiotherapy ionizing radiation is used to cause DNA damage in cancer cells. The DNA damage is rapidly sensed and a network of intracellular DNA damage signaling cascades is activated. These signaling cascades influence whether the cells die or survive, through regulation of DNA repair, cell cycle checkpoint and cell death pathways. Our group works at the border between basic and translational radiation research. We conduct basic projects to identify novel mechanisms of DNA damage signaling, in addition to more applied projects to understand how inhibitors of DNA repair and checkpoints can be used in an optimized manner for cancer treatment. Three project groups, headed by Beata Grallert Trond Stokke and Sebastian Patzke, are members of our group.

AIMS

- Obtain new knowledge about cellular responses to radiation, with focus on cell cycle checkpoints, DNA damage signaling and repair, and explore how such knowledge can be used to improve radiotherapy.

PROJECTS

- Pre-clinical exploration of the effects of checkpoint kinase (CHK1, WEE1, ATR) and PARP inhibitors
- Functional roles of Protein phosphatase 1 (PP1) targeting subunits in DNA damage signaling
- Identification of drugs that inhibit DNA repair after radiation, through flow cytometry-based compound screens
- Centrosomal and ciliary proteins in cell cycle regulation and genome integrity- roles and targets in cancer etiology, diagnostics and treatment
- The role of GCN2 and translational regulation in the cell cycle and cellular stress

RECENT ACHIEVEMENTS:

In 2019, a total of 10 articles were published. Members of the group were senior/first authors on 6 of these, published in Nucleic Acids Research, Cell Reports, Frontiers in Oncology, Cell Cycle, Current Genetics and Bioessays. One new grant was obtained from the Norwegian Research Council.



Department of Tumor Biology

Headed by Gunhild M. Mælandsmo

The department has four research groups and 54 employees with a common vision to improve treatment of metastatic cancer. Our strategy is, through basic and translational research in the areas of cancer biology and computational science, to advance systems understanding and identify novel intervention strategies. We emphasize multidisciplinary competence and collaboration between researchers, clinicians and patients to stimulate the necessary synergy for improved cancer care.

We are performing basic, translational and clinical research, and our scientific goal is to provide knowledge for clinical translation of precision cancer medicine. We will do so by contributing with expertise in genomics and bioinformatics and by utilizing patient samples as model systems for investigation of therapeutic efficacy. We have a large collection of patient-derived cell lines and xenografts from different types of human cancer. The models are utilized for mechanistic studies of disease progression and treatment responses, and for preclinical evaluation of novel drugs and drug combinations. Together with patient samples, collected in consecutive, longitudinal or study-specific biobanks, the patient-derived models serve as resources to identify novel targets and biomarkers.

To foster a strong link between translational and clinical research we have several researchers holding part-time clinical positions. An ambition for the department is to participate in design and conduct of clinical trials, and to provide molecular and bioinformatics competences in

multidisciplinary tumor boards in the area of precision cancer medicine.

Key achievements over the last 3-4 years include:

- Project leader responsibilities in collaborative projects in precision cancer medicine:
NCGC - The Norwegian Cancer Genomics Consortium, NoSarc - Norwegian Sarcoma Consortium, MetAction - Actionable targets in cancer metastasis, MOVEMBER - The Norwegian Prostate Cancer Biomarker Consortium, The EuroPMP Cost Action, Biobank Norway 2 - multicenter biobanking of prostate cancer tissue in Norway
- Project leader responsibilities for clinical trials:
NeoAva: Bevacizumab in combination with neoadjuvant chemotherapy in breast cancer
I-BCT: DNA targeted chemotherapy to breast cancer patients with an aggressive phenotype
ImmunoPeCa: Immunotoxin in Peritoneal Carcinomatosis
- Co-PI responsibilities in the METIMMOX multicenter trial which will investigate the combination of oxalip-
latin and checkpoint inhibition (nivolumab) in micro-
satellite stable metastatic CRC
- All research groups have during the period received major funding (Norwegian Cancer Society, H2o2o, South-Eastern Norway Regional Health Authority)
- The department has published 66 papers, with 28 as first or last author, and educated three PhDs and one M.Sc. in 2019



Metastasis Biology and Experimental Therapeutics

Group leader: Gunhild M. Mælandsmo



“Cellular plasticity - the route to resistance and metastasis”

ABOUT

Employees: The group has 20 members with multidisciplinary background and expertise (cell- and molecular biologists, medical doctors, laboratory- and animal technicians), including two MDs in shared clinical positions. Several of the technicians are trained in specialized technologies and compose resources for all groups in the department.

Research focus: Investigations on mechanisms of resistance and metastasis for improved cancer treatment.

Methodology: Molecular and functional analyses utilizing clinical samples, human cell cultures and patient-derived models (ex vivo, in vitro and in vivo).

AIMS

Metastatic progression is the major clinical challenge and the principal cause of death in cancer patients. To combat metastatic cancer our goal is to understand the underlying biological mechanisms causing therapy resistance, invasion and re-growth. Knowledge on these processes is vital for identification of molecules that can be utilized as prognostic and predictive biomarkers or as targets for therapy. We are working with malignant melanoma, breast cancer and prostate cancer.

PROJECTS

1. Basic research revealing mechanisms causing treatment resistance and metastasis
 - Molecular and cellular determinants regulating cancer cells plasticity, with special emphasis on the role of tumor-stroma interactions

2. Preclinical research investigating novel drugs and drug combinations
 - Mechanistic studies and assessment of treatment efficacy in patient-derived models in vivo and ex vivo
 - Biomarker discovery by molecular and functional techniques
 - Response evaluation of experimental drugs (often in collaboration with commercial partners, eg.: Lytix Biopharma and Arctic Pharma)
3. Clinical trials in precision medicine – clinical and translational efforts towards biomarker discovery
 - NeoAva: bevacizumab in combination with neoadjuvant chemotherapy for patients with locally advanced breast cancer
 - I-BCT: extended DNA-targeted chemotherapy to breast cancer patients with an aggressive phenotype

RECENT ACHIEVEMENTS

- The group was credited with 20 publications in 2019, of which nine with group members as first and/or last author; two PhD degrees completed
- One clinical intervention trial in breast cancer open for inclusion (I-BCT)
- Successful establishment of a user board for melanoma research
- Funding: Two innovation grants from the South-Eastern Norway Regional Health Authority, one researcher grant for biomarker stratification in breast cancer and two partnerships in new H2020-projects
- One patent approved

Translational Cancer Therapy

Group leader: Kjersti Flatmark



“New treatment for metastatic colorectal cancer”

ABOUT

In 2019, the Translational Cancer Therapy group comprised 16 members (including part-time employees and students) with a broad variety of expertise, including basic biologists, translational scientists, and clinician-scientists. Our approach to research is translational, encompassing methods from biochemistry and cell biology, preclinical drug testing in animal models, biomarker studies in clinical cohorts and interventional clinical trials. Extensive collaboration has been established within the Institute, with clinical departments, nationally and internationally.

AIMS

Our long-term aim is to make new, efficacious treatment(s) available to patients with colorectal cancer (CRC). This will be accomplished by bringing the clinic and lab together in translational research projects utilizing 1) preclinical models for efficacy and mechanistic studies 2) clinical cohort studies to understand CRC biology and identify biomarkers and therapeutic targets 3) clinical intervention trials to test new drugs and therapeutic concepts in patients.

PROJECTS

- Peritoneal metastasis – molecular targets and new therapies
 - Personalizing CRC therapy – identification of biomarkers and therapeutic targets in locally advanced and metastatic CRC, involving generation and use of our extensive biobanks, molecular and bioinformatics analyses (subprojects include: genomics, microRNA, mRNA, and immune cell analysis, participation in the BigMed project)
 - Novel drugs and therapeutic concepts in models of peritoneal and liver metastases
- Translational studies within the METIMMOX multicenter trial (Colorectal Cancer Metastasis – Shaping Anti-Tumor Immunity by Oxaliplatin), which will investigate the combination of oxaliplatin and checkpoint inhibition (nivolumab) in microsatellite stable CRC
- Commercial development of MOC31PE and BM7PE immunotoxins for cancer therapy

RECENT ACHIEVEMENTS

- Group members were credited with 25 publications in 2019
- The multicenter METIMMOX trial has included almost half of the planned number of patients, and interesting responses have been registered
- Our COST Action, EuroPMP, European Research Network on rare cancer pseudomyxoma peritonei currently comprises >80 members from 21 European countries.
- Two patent applications filed

Computational Cancer Genomics

Group leader: Eivind Hovig



“Enabling the transition to clinical utility”

ABOUT

The 10-member group has strong interest in the development of computational approaches in cancer genomics, with focus on melanoma systems biology, and prostate cancer functional genomics. Further activity is centered on computational aspects of deep sequencing for cancer, with downstream analysis. The group facilitates moving precision cancer medicine towards the clinic, leveraging participation in the BigMed RCN-financed ICT lighthouse project.

The group collaborates with deCODE, Iceland, on the characterization of a familial cancer biobank, with 16000 genotyped samples, and where 2000 Norwegians have been whole genome sequenced. We pursue both the heritable cancer information from this effort, as well as a characterization of the ethnic Norwegian population genetics as a communal resource for Norway.

AIMS

- to develop and apply novel methodologies for computational studies of cancer-related processes, including statistical genomics, drug prediction algorithms, and mutational processes
- to contribute insight on heritable cancer-predisposing DNA-variation in the Norwegian population
- to characterize geographical stratification aspects of the Norwegian population
- to develop solutions for precision cancer medicine towards the clinic
- to understand signaling and epigenetic processes in melanoma and prostate cancer

PROJECTS

- Development of solutions for integrative cancer sequencing towards diagnostic utility, for patient stratification, and prediction of treatment response
- Participation in national and international efforts for standardization and development of best practice methods, including the 1 Million genomes community effort, partner of the BigMed project, and participating in the Center of innovation Big Insight for the knowledge economy.
- Melanoma signaling perturbations for systems understanding, including MITF, BRAF and CDKN2A aberrations
- Biomarkers in prostate cancer epigenomics
- Identification of familial cancer predisposing variants in Norwegian families with overrepresentation of cancer
- Clinical implications of Lynch syndrome genetics in Europe and Latin America

RECENT ACHIEVEMENTS

- Group members were credited with 15 publications in 2019, with 8 as first or last author
- Cancer society research funding for prostate cancer to Alfonso Urbanucci

Molecular Biology of Sarcomas

Group leader: Jørgen Wesche



“Towards precision medicine to improve treatment of sarcomas”

ABOUT

The 14-member group has a long standing interest in the biology of mesenchymal tumors (sarcomas). The current focus is on precision medicine for sarcomas. To achieve this, the group has broad expertise in basic cell biology, genomics and translational research and, in addition, one MD in a shared clinical position. The group is part of a Centre of Excellence (CanCell).

AIMS

The group aims to improve the treatment of sarcoma patients by investigating how genetic changes (mutations) affect signalling in sarcoma cells, and how tumors evolve and become resistant. We combine extensive genomic characterization of clinical cohorts with preclinical investigation in sarcoma cell lines and xenografts to better understand the biology of sarcomas. Sarcomas are rare cancers with poor treatment options, and we aim to use our biological knowledge to identify new treatments opportunities by repurposing approved drugs for other cancers types.

PROJECTS

- Sarcoma cell biology – Gaining understanding of the development and progression of rhabdomyosarcoma, liposarcoma and osteosarcoma, which will lead to the identification of biomarkers and novel drug targets. A main focus is the study of the role of fibroblast growth fact receptors (FGFRs).
- Norwegian Sarcoma Consortium (NoSarC) – Biobanking and genomic characterization of patient material of 3-4 national cohorts of sarcomas (~500 samples).
- Exploration of “liquid biopsies”, as a non-invasive methods for detection of tumor-derived DNA in blood, to monitor disease progression, treatment response and tumor evolution.
- Dissecting drug resistance in gastrointestinal stromal tumors - Revealing the underlying mechanisms of early drug resistance in light of the complexity of the tumor to identify novel treatment modalities.

RECENT ACHIEVEMENTS

- 10 publications in 2019
- The group obtained a grant from the South-Eastern Norway Regional Health Authority (PhD project).
- Group visit to Vall d'Hebron, Barcelona, resulted in a new important collaboration
- 1 Master degree was completed



“Providing cutting-edge technology and competence to excel research”



Department of Core Facilities

Headed by Leonardo A. Meza-Zepeda

The Department of Core Facilities runs six regional and national technology platforms financed by the South-Eastern Norway Regional Health Authority and the Research Council of Norway, providing advanced competence, infrastructure, and services to regional, national and international users. The Department aims to deliver easy access to cutting-edge advanced technologies and competence, and to improve research quality through an optimal choice of technology, ultimately increasing the scientific competitiveness of our users. The Department of Core Facilities is organized in three units; Advanced Microscopy, Genomics and Bioinformatics, and Flow Cytometry and Pre-Clinical Imaging, with a total of 20 employees. More information at www.ous-research.no/corefacilities.

ADVANCED LIGHT MICROSCOPY

Unit Leader: Ellen Skarpen

Scientifically responsible: Harald Stenmark

Facility staff: 2

The Core Facility for Advanced Light Microscopy (ALM) offers services in various types of ALM, including confocal microscopy, live-cell microscopy and superresolution microscopy. Current instruments include a Zeiss LSM 880 FAST airyscan microscope, a Zeiss LSM 710 confocal microscope, and an OMX Blaze microscope for structured illumination and STORM super-resolution imaging, as well as live-cell microscopy. The core facility

cooperates with nodes at the Rikshospitalet and Ullevål campuses of OUH, as well as the light microscopy core facility at the Institute for Biosciences, University of Oslo. The core facility for ALM is a Eurobioimaging node, providing state-of-the-art technologies in the fields of biological and medical imaging. Services include training, courses, access to microscopes and microscopy performed by competent core facility personnel.

ADVANCED ELECTRON MICROSCOPY

Unit Leader: Ellen Skarpen

Scientifically responsible: Andreas Brech

Facility staff: 1

The Core Facility for Advanced Electron Microscopy (AEM) includes nodes at the Radium Hospital and Rikshospitalet, and provides service, training, and access to microscopes for ultrastructural studies. Available techniques at the facility include conventional plastic embedding, immune EM, correlative light/electron microscopy (CLEM), high-pressure freezing, electron tomography, cryo-EM and STEM. The facility staff is actively developing new methods in order to offer state-of-the-art microscopy solutions for researchers. We cooperate with the imaging platform at the Institute for Biosciences, University of Oslo and are part of the Norwegian Advanced Light Microscopy node within EuroBioImaging.

BIOINFORMATICS

Unit Leader: Susanne Lorenz
Scientifically responsible: Eivind Hovig
Facility staff: 5

The Bioinformatics Core Facility (BCF) provides service, support and advice in the field of bioinformatics. By combining biology with computer science, statistics and mathematical modelling, we offer support for analysis and interpretation of biological data for basic and translational research. The BCF has a specific focus on developing and providing solutions for precision medicine and cancer-related projects. Through dedicated support from the South-Eastern Norway Regional Health Authority, we are actively assisting in developing the infrastructure for precision diagnostics at the hospital. Our service also includes support with data handling and storage by providing approved solutions for sensitive data. The operation of the BCF is aligned with the national and local Elixir bioinformatics infrastructure project, and also interacts with USIT, University of Oslo.

HIGH-THROUGHPUT SEQUENCING (GENOMICS)

Unit Leader: Susanne Lorenz
Scientifically responsible: Leonardo A. Meza-Zepeda
Facility staff: 5

The Genomics Core Facility (GCF) provides state-of-the-art high-throughput genomic services to the scientific community. The GCF offers advanced technologies and competence to study genome structure, dynamics and function using high-throughput sequencing, nanoString, and microarray technologies. Our services include solutions to study the transcriptome, genome, and epigenome from multi-gene analysis to a genome-wide level. Highly experienced service personnel provide advanced support to clinical, translational and basic research projects. We have extended our services for single-cell transcriptome analysis to the single-cell analysis of T & B cell receptors, scATAC-Seq, and feature barcoding to study protein expression. In addition, we have established a collaboration with the Department of Pathology at OUH to develop the infrastructure for advanced precision diagnostics towards clinical studies with dedicated support from the South-Eastern Norway Regional Health Authority. The GCF is a member of the Norwegian Consortium for Sequencing and Personalised Medicine (NorSeq) running the National platform for sequencing technology. The core facility collaborates with the sequencing node at the Ullevål campus for sequencing services.

FLOW CYTOMETRY

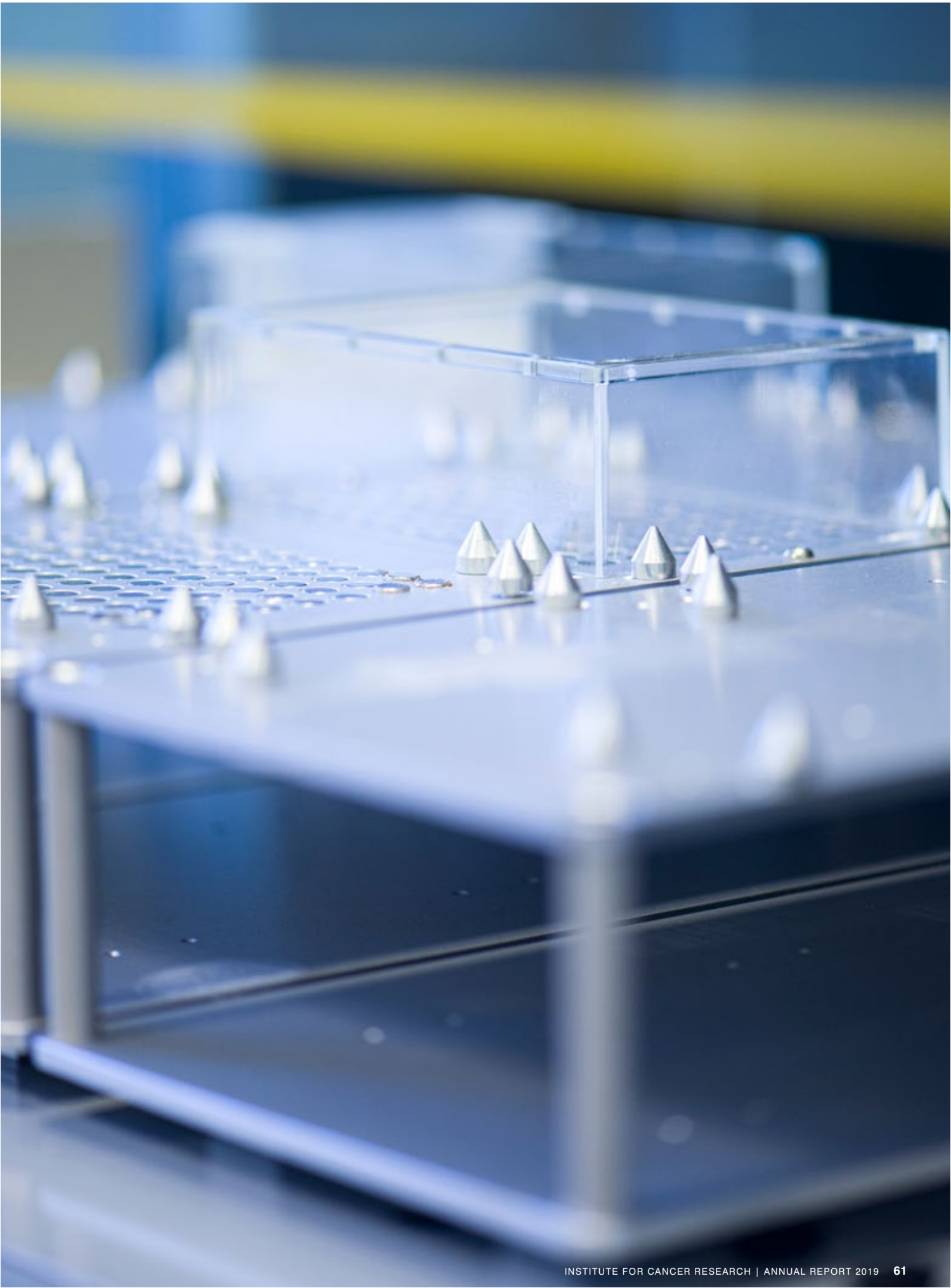
Unit Leader: Trond Stokke
Scientifically responsible: Trond Stokke
Facility staff: 4

The Flow Cytometry Core Facility (FCCF) provides analysis and sorting services for users within the South-Eastern Health Region of Norway. We have recently installed a state-of-the-art analyser (BD Symphony) with 5 lasers that may measure up to 28 fluorescence parameters simultaneously. In total, the core facility provides services using 3 analysers and two sorting instruments. Flow cytometry analysis is performed by the users themselves. Sorting experiments are either performed by core facility personnel (in the BD Aria), or by the users in the Sony SH100 sorter. The FCCF has possibilities for high throughput screening, processing and staining of cells in 96- or 384-well plates, followed by an automated analysis. We also have a “mass-spec flow cytometer” (Helios). This instrument can measure up to 60 parameters simultaneously at single-cell resolution. We have recently installed an add-on to the Helios, Hyperion, which allows for imaging of sections labelled with up to 60 heavy metal-tagged antibodies. The FCCF collaborates with facility nodes at the Rikshospitalet and Ullevål campuses of OUH.

PRECLINICAL IMAGING FACILITY

Unit Leader: Trond Stokke
Scientifically responsible: Tord Hompland
Facility staff: 2

The Preclinical Imaging Facility provides access to state-of-the-art non-invasive imaging equipment for mice and rats. The equipment is situated within the animal facility and consists of a 7T Bruker MRI, IVIS spectrum and Zeiss Stereo Microscope for optical imaging, and a Multirad 225 small animal irradiator capable of doing x-ray imaging. The facility also provides all the necessary equipment for in vivo research on small animals or tissue/organ samples from larger animals or humans. A comprehensive library of standard off-the-shelf imaging protocols are available, and custom-protocols can be developed upon user request. We are at present developing a protocol for synchronization of images obtained by MRI, IVIS and X-ray imaging. The service offered by the core facility includes design, development and running of the imaging experiment, as well as post-processing of the data in addition to instrument-specific training. The core facility collaborates with the Preclinical MRI node at the Ullevål campus.



A blue-tinted photograph of a laboratory setting featuring a microscope and various glassware.

Research centres

search

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Centre of Excellence

The Centres of Excellence (CoE) scheme is a national programme under the auspices of the Research Council of Norway. The granted CoE (CanCell) is funded for 5 + 5 years (if recommended following a mid-term evaluation). The total basic funding is ~167 million NOK.

A blue-tinted photograph of a woman with long dark hair looking through a microscope.

centres

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K. G. Jebsen Centres

The K.G.Jebsen Foundation supports Centres for Medical Research of high international quality as part of its program for translational medicine. The centres are selected in collaboration with the Norwegian Medical Faculties and University Hospitals for a period of 4 years with the possibility of a 2-year extension. The selected Centres receive 16-18 million NOK in basic funding over the first four years from the Foundation and support from the host institutions, University of Oslo (KG Jebsen Centre for Cancer Immunotherapy, KG Jebsen Centre for B Cell Malignancies) or Oslo University Hospital (KG Jebsen Colorectal Cancer Research Centre).



“Reprogramming
of cancer”

Centre for Cancer Cell Reprogramming (CanCell)

Headed by Harald Stenmark



ABOUT

Centre for Cancer Cell Reprogramming (CanCell) is a Norwegian Centre of Excellence (CoE) which opened 01.01.2018 and has a planned CoE funding period of 10 years. CanCell is led by Harald Stenmark at Institute for Cancer Research, whereas Anne Simonsen at Institute of Basic Medical Sciences is co-director. The centre consists of 6 research groups and has 7 associate members (Eivind Hovig, Åslaug Helland, Yngvar Fløisand, Philippe Collas, Arnoldo Frigessi, Emmet McCormack, and Terje Johansen) and 4 international visiting professors (Kristian Helin, Ivan Dikic, Eileen White, and Eyal Gottlieb). By the end of 2019, CanCell had 100 members of 34 (!) nationalities.

AIMS

CanCell's vision is to identify novel vulnerabilities of cancer cells that can be targeted for cancer cell reprogramming. The centre's founding hypothesis is that pathway intersections between chromatin regulation, membrane dynamics, cell signaling and metabolism during cancer progression represent potential “Achilles’ heels” of cancer cells. These will be identified through close cooperations between specialists within these four cellular processes, and will be targeted by genetic and pharmacological regimens to achieve reprogramming of cancer cells into harmless (or dying) cells.

PROJECTS:

- Membrane dynamics in cancer
- Autophagy in immunity and cancer
- Molecular medicine of leukemia
- Tumour-host interactions
- Molecular biology of sarcomas
- Mechanisms of epigenetic regulation in cancer

RECENT ACHIEVEMENTS

- Demonstration of distinct functions of ATG16L1 isoforms in membrane binding and LC3B lipidation in autophagy-related processes (Lystad et al., *Nature Cell Biology*, 2019).
- Identification of NIPSNAP proteins as “eat-me” signals for autophagic degradation of damaged mitochondria (Abudu et al., *Developmental Cell*, 2019). This was a collaboration between Anne Simonsen's group and the group of CanCell associate member Terje Johansen.
- Demonstration that ESCRT proteins seal the autophagosome during starvation-induced autophagy and autophagy of damaged mitochondria (Zhen et al., *Autophagy*, 2019). This was a collaboration between the CanCell groups of Harald Stenmark and Anne Simonsen.
- Identification of a new mechanism of tumour suppression – intracellular retention of matrix metalloproteinases (Sneeggen et al., *Nature Communications*, 2019).

- Demonstration that Centralspindlin recruits ALIX to the midbody during cytokinetic abscission in fruit flies via a mechanism analogous to virus budding (Lie-jensen et al., *Current Biology*, 2019).
- Demonstration that desumoylation of RNA Polymerase III lies at the core of the Sumo stress response (Nguéa et al., *Journal of Biological Chemistry*, 2019).
- Demonstration that cancer mutations in fibroblast growth factor receptor 2 prevent a negative feedback loop mediated by the ERK1/2 pathway (Szybowska et al., *Cells*, 2019).
- CanCell scientists published 41 papers in 2019, many of these in leading journals. Two PhD students, Marte Sneeggen and Ignacio Cuervo, successfully defended their theses. Eight MSc students were graduated.
- In 2019, major grants were obtained by Jorrit Enserink, Leonardo Meza-Zepeda, Anne Simonsen, Harald Stenmark, Kay O. Schink, Swarupa Panda, Helene Knævelsrud and Viola Nähse. Two of these were international grants – EMBO long-term fellowship to Swarupa Panda and Horizon-2020 Initial Training Network grant to Anne Simonsen.

GROUP LEADERS/STEERING COMMITTEE

CanCell was established by the following 6 group leaders, who also serve as CanCell's steering committee:

Harald Stenmark, Institute for Cancer Research, OUS, and Institute of Clinical Medicine, UiO

Anne Simonsen, Institute of Basic Medical Sciences, UiO, and Institute for Cancer Research, OUS

Jorrit Enserink, Institute for Cancer Research, OUS, and Department of Biosciences, UiO

Ragnhild Eskeland, Institute of Basic Medical Sciences, UiO

Tor Erik Rusten, Institute of Clinical Medicine, UiO, and Institute for Cancer Research, OUS

Jørgen Wesche, Institute for Cancer Research, OUS, and Institute of Basic Medical Sciences, UiO

“Our goal is to develop new therapeutic strategies that overcome immune tolerance to target cancer”

K.G. Jebsen Center for Cancer Immunotherapy

Headed by Johanna Olweus



Kristian Gerhard Jebsen Foundation

ABOUT

In the K.G. Jebsen Center for Cancer Immunotherapy (JCIT) hosted by the University of Oslo, five Norwegian and one Dutch research group have come together to find new immunotherapeutic strategies that overcome the major challenge of self-tolerance in cancer. JCIT was granted prolongation following the first 4-year period, throughout 2019. The partnering groups of JCIT span complementary competencies ranging from basic proteomics, cell signaling and T-cell receptor engineering to expertise in experimental clinical immunotherapy trials. This places the center in a unique position to pursue novel therapeutic opportunities, and the strong focus on translating therapeutic opportunities is a fundamental characteristic of JCIT. Results from basic research are pursued through the necessary translational steps to testing in patients, and in-depth mechanistic studies of patient material obtained in experimental clinical trials are performed with the aim of improved designs of immunotherapeutic strategies.

AIMS

Three principally distinct approaches are pursued in an interdisciplinary and translational program: 1. Use of T-cell and NK-cell based alloreactivity to overcome self-tolerance. 2. Identification and targeting of cancer-specific neo-antigens. 3. Enhancement of effector T-cell function by counteracting inhibitory mechanisms.

PROJECTS

- Epitope discovery to identify targets for immunotherapy
- Identify immune cells expressing immune receptors that recognize and kill cells that express the identified targets
- Molecular cloning, genetic transfer and profiling of immune receptors
- Enhance cytotoxic lymphocyte function by overcoming inhibitory mechanisms
- In vivo evaluation of immune modulating therapies

RECENT ACHIEVEMENTS

- Deciphered molecular mechanisms of NK cell homeostasis, education and differentiation (*Goodridge et al, Nature Communications, 2019, Jacobs et al., J. Immunology 2019 and Pfefferle et al., Cell Reports 2019*).
- Described a technology for identification of neoantigen specific T cells from healthy donors (*Ali et al, Nature Protocols, 2019*)
- Olweus was awarded an ERC Consolidator Grant (2 mill Euro over 5 yrs) for an application based on results generated within the center – “Outsourcing cancer immunity”.
- Reported low and variable tumor reactivity of the intratumoral TCR repertoire in human cancers (*Scheper et al, Nat. Med. 2019*)
- Included 10 patients in Lymvac-2, an experimental immunotherapy trial combining intratumoral immunotherapy with anti-PD1 for treatment of patients with follicular lymphoma, in collaboration with Merck (Kolstad PI).
- Started GMP virus production for first clinical trial

- in Norway using an in-house generated immune receptor (JCIT TCR) for cancer immunotherapy.
- Recruited first 300 patients to the ASAC trial that examines the effect of reversing prostaglandin E2-mediated inhibition of tumor immunity in patients with metastatic colorectal cancer by the end of 2019 (www.asac.no).
- Malmberg completed a research sabbatical at UCSD resulting in new collaborative efforts with leading iPSC-NK/T cell environments, including Kaufman Lab at UCSD, and Sadelain Lab at MSKCC (*Sætersmoen et al, Seminars in Immunopathology 2019, Hong et al., Journal of Clinical Investigation in press*).
- Demonstrated that idelalisib preferentially inhibits human regulatory T cells, enhancing anti-tumor immunity (*Chelappa et al, J Immunol., 2019*) and characterized the CD28 and CD2 interactomes in T cells (*Skånland and Taskén, J. Immunol. 2019*)
- Characterized how metabolic regulation through the FoxK1/2 master switch affects T cell function (*Sukonina et al, Nature, 2019*) and contributed to understanding TLR8 signaling in T cells (*Meås et al., Nature Commun., in press 2020*).
- Obtained funding for proteomics infrastructure from The Norwegian Research Council (total 50M NOK, Lund-Johansen partner in NAPI consortium) and funding from the Norwegian Cancer Society for identification of HLA_bound peptides from tumor-associated antigens.
- Obtained funding from UiO for production of antibodies to peptides in the ImmunoLingo convergence consortium (total 11.5 mill NOK, Lund-Johansen partner).

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<http://www.med.uio.no/klinmed/english/research/centres/kgj-cancer-immunotherapy/>

GROUP LEADERS/ STEERING COMMITTEE

Johanna Olweus (MD, PhD, Director), Dept of Cancer Immunology, Institute for Cancer Research, Oslo University Hospital (OUH)-Radiumhospitalet and University of Oslo (UiO)

Karl-Johan Malmberg (MD, PhD, deputy Director), Dept of Cancer Immunology, Institute for Cancer Research, OUH-Radiumhospitalet and UiO

Arne Kolstad (MD, PhD), Dept of Oncology, OUH-Radiumhospitalet

Kjetil Taskén (MD, PhD), Dept of Cancer Immunology, Institute for Cancer Research, OUH-Radiumhospitalet and UiO

Fridtjof Lund-Johansen (MD, PhD), Dept of Immunology, OUH-Rikshospitalet

Ton Schumacher (PhD), The Netherlands Cancer Institute, Amsterdam

“From basic research and preclinical studies to precision medicine for B-cell malignancies”

K.G. Jebsen Centre for B-cell malignancies

Headed by Ludvig A. Munthe and June H. Myklebust



Kristian Gerhard Jebsen Foundation

ABOUT

The K.G. Jebsen Centre for B-cell malignancies was established in June 2018, and bridges 4 translational research groups with 3 clinical groups; placing us in a unique position to translate pre-clinical results into clinical trials in lymphoma, B-cell leukemia and multiple myeloma (MM). The centre utilizes “cutting edge” technologies for deep profiling of patient samples, cancer drug sensitivity screens for precision medicine, and development of novel approaches for immunotherapy. Collectively, the Centre represents a multidisciplinary integration of life science research with preclinical development of personalized medicine, drug discovery and cell-based immunotherapy, as well as clinical trials and establishment of best practice on how to treat B-cell malignancies.

AIMS

The centre aims to identify, develop and test new therapeutic options for patients with B-cell malignancies.

PROJECTS

- Identify molecular biomarkers to guide precision medicine and to identify high risk patients
- Deciphering signal integration and interactions with the tumor microenvironment to reveal actionable targets
- Develop novel therapeutics: identify antigens for vaccination, T-cell epitope discovery, and CAR T cell design
- Preclinical testing: immunotherapy and personalized medicine
- Translating results into clinical initiatives – from bench to bedside and back

RECENT ACHIEVEMENTS/CLINICAL TRANSLATION

- The clinical portfolio includes 58 clinical trials at different stages, majority are researcher initiated.
- Enrolled most MM patients in clinical trials
- Ixazomib improved progression-free survival (PFS) after autologous stem cell transplantation in MM (Dimopolous, *Lancet* 2019).
- The addition of isatuximab to pomalidomide-dexamethasone or pomalidomide to bortezomib-dexamethasone improved PFS in refractory/relapsed MM (Attal, *Lancet* 2019; Richardson, *Lancet Oncol* 2019).
- Consensus recommendations on imaging in monoclonal plasma cell disorders (Hillengass, *Lancet Oncol.* 2019).
- Phase II trial testing CD19 CAR T-cell therapy in Diffuse Large B-Cell Lymphoma demonstrated 52% overall response rate (Schuster, *NEJM* 2019).
- Chemotherapy can be reduced in aggressive Non-Hodgkin's Lymphoma with favorable prognosis (Poeschel, *Lancet* 2019)
- Established B-cell directed therapy as treatment of-choice in chronic cold agglutinin disease (Berentsen, *J Blood Med* 2019)
- Relapse risk and loss of lifetime after modern combined modality treatment of young patients with Hodgkin Lymphoma (Bicler, *J Clin Oncol* 2019).

- Investigated inheritance of susceptibility to malignant blood disorders (Jonsson, *Sci Rep.* 2019).
- Stromal cell PKC-beta inhibition enhances chemosensitivity in B-cell malignancies and overcomes drug resistance (Park, *Sci Transl Med*, in press 2019).
- Developed in vitro phospho-flow cytometry assay for biomarker discovery and dose prediction in CLL (Skånland, *Leukemia*, in press 2019).
- Distinct subtypes of diffuse large B-cell lymphoma can be defined by hypermutated genes (Alkods, *Leukemia* 2019).
- Developed the prognostic predictor BTK-FLIPI to identify follicular lymphoma patients at high risk for adverse disease (Steen, *Haematologica* 2019), and contributed to convergence of risk prediction models (Silva, *Haematologica* 2019).
- Identified TIGIT and PD-1 as relevant targets for co-checkpoint blockade in B-cell lymphoma (Josefsson, *Cancer Immunol Res* 2019).
- Identified tumor-reactive T cells in B-ALL patients (Bürgler, *Oncogene* 2019).
- Bone marrow Th1 T cells induce activation and proliferation of leukemic cells from B-ALL patients (Traxel, *Oncogene* 2019).
- Preclinical development of CD37 CAR T-cell therapy (Köksal, *Blood Adv* 2019; patent filed).
- The PI3Kδ inhibitor idelalisib enhances anti-tumor immunity preferentially through inhibition of regulatory T cells (Chellappa, *J Immunol*, 2019), and CSK binds CD28 upon activation and mutes downstream signaling (Skånland, *J Immunol* 2019).
- B-cell receptor ligation induces display of V-region immunoglobulin peptides on MHC class II molecules to T cells (Hutsky, *PNAS* 2019)
- FOXK1 and FOXK2 regulate aerobic glycolysis (Sukonia, *Nature* 2019).


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<https://www.med.uio.no/klinmed/english/research/centres/kgj-b-cell-malignancies/>

GROUP LEADERS/STEERING COMMITTEE

Ludvig A. Munthe (MD, PhD, Centre Director)^{1,2}
 June H. Myklebust (PhD, Assistant Director)^{2,3}
 Geir E. Tjønnfjord (MD, PhD)^{2,4}
 Harald Holte (MD, PhD)⁵
 Hilde Schjerven (PhD)^{1,6}
 Erlend B. Smeland (MD, PhD)^{2,3}
 Kjetil Taskén (MD, PhD)^{2,3}

- ¹ Dept. of Immunology, Div. for Laboratory Medicine, Oslo University Hospital (OUH)
- ² Institute for Clinical Medicine, University of Oslo
- ³ Dept of Cancer Immunology, Institute for Cancer Research, OUH
- ⁴ Dept. of Haematology, 5Dept. of Oncology, Div. for Cancer Medicine, OUH
- ⁵ Dept. of Laboratory Medicine, University of California, San Francisco.



“High quality
translational
research to the
benefit of
colorectal cancer
patients”

K.G. Jebsen Colorectal Cancer Research Centre

Headed by Ragnhild A. Lothe



ABOUT

Colorectal cancer (CRC) is a major global health burden, and the focus of our Centre is to meet the current challenges in disease management by improved patient monitoring and stratified treatment. The Centre is hosted by the Clinic for Cancer Medicine, Oslo University Hospital (OUH). The Centre PIs are also partners in the recently appointed OUH Strategic Research Area TEAM-ACT (2019-24).
Home page: www.colorectalcancer.no

GROUP LEADERS/STEERING COMMITTEE

- **Professor Ragnhild A. Lothe** (MSc, PhD, Centre leader), Dept. of Molecular Oncology, Institute for Cancer Research, OUH and Institute for Clinical Medicine, University of Oslo (UiO)
- **Professor Arild Nesbakken** (MD, PhD, deputy Centre leader), Dept. of Gastrointestinal Surgery, OUH and Institute for Clinical Medicine, UiO
- **Professor Rolf I. Skotheim** (MSc, PhD), Dept. of Molecular Oncology, Institute for Cancer Research, OUH, and Dept. of Informatics, UiO.
- **Senior Consultant Marianne G. Guren** (MD, PhD), Dept. of Oncology, OUH
- **Professor Guro E. Lind** (MSc, PhD), Dept. Molecular Oncology, Institute for Cancer Research, OUH and Dept. of BioSciences, UiO

Our Centre has an active Patient Advisory Board, established in 2016.

AIM

Translate biomedical knowledge in the context of tumor heterogeneity into improved stratified medicine of CRC.

PROJECTS

- Clinical and molecular biomarkers for improved risk stratification of patients
- Model tumor heterogeneity and clonal evolution to monitor early relapse and treatment failure
- Pharmacogenomic profiling of organoid models derived from the patients' own tumor cells to guide therapy selection, identify biomarkers for response prediction, and develop synergistic drug combinations

RECENT ACHIEVEMENTS AND CLINICAL TRANSLATION

In 2019, we were granted a 6-year strategic research area from OUH (TEAM-ACT) to continue the translational and clinical research program on primary and metastatic CRC established in the K.G. Jebsen Centre. TEAM-ACT is led by professor R. A. Lothe and associate professor Anita Sveen. Sveen also received a major grant from the Norwegian Cancer Society to further develop the platform for pre-clinical pharmacogenomics of patient-derived models to

enable testing and optimization of immunotherapy. Peter Wold Eide received the Young Researcher Award from the national annual meeting in oncology (“Onkologisk Forum”) for his studies in CRC genomics, and Sveen received the Early Career Award from OUH. Heidi Pharo and Kaja Christine Graue Berg defended their PhDs at the UiO.

2019 was a successful year for the Centre with respect to scientific production. Together with collaborator Scott Kopetz at The MD Anderson Cancer Centre, USA, a practice changing clinical trial of targeted combination therapy in metastatic CRC was published in NEJM (Kopetz et al., 2019), in addition to a comprehensive review on biomarker-guided therapy for CRC (Sveen et al., Nature Rev Clin Oncol 2019). Totally 31 peer-reviewed papers were published within a broad range of topics. Epidemiological research demonstrated a rising incidence of CRC among young people (Araghi M et al., Lancet Gastroenterol Hepatol 2019), and compared survival after surgery for CRC in Norway to other high-income countries (Benitez Majano S et al., Lancet Oncol, 2019). Molecular studies were published of circulating biomarkers for management of CRC (Hamfjord J et al., Ann Oncol 2019), the strong influence of tumor microenvironment features on patient prognosis (Dienstmann et al., Ann Oncol, 2019; Glaire et al., Brit J Ca 2019), tumor heterogeneity and multi-level genomics (Berg et al., Oncogene 2019; Brunsell et al., Eur J Surg Oncol; Smeby et al., ESMO Open 2019), as well as advanced technologies (Eilertsen et al., Cancer Letters 2019; Lopes et al., Lab Invest 2019).

Unpublished results of drug screening and molecular profiling of organoids derived from 30 patients with multiple CRC liver metastases, gene expression-based classification in a tumor heterogeneity context, and identification of new high-level amplifications in CRC were recently presented at the Keystone Symposium on “Cancer Evolution and Combinatorial Cancer Therapies” in Banff, Canada.

Key opinions:

Centre members published several editorials and reviews related to a variety of challenges in CRC: “The complexity of biomarkers” (Sveen et al., Nat Rev Clin Oncol); “The global challenge of CRC” (Guren MG, Lancet Gastroenterol Hepatol); “Gene expression-based modelling of targeted therapies” (Sveen et al., Ann Oncol 2019); “Combination therapies with HSP90 inhibitors” (Kryeziu et al., BBA Rev Cancer); “Circulation biomarkers for early detection and management” (Marcuello et al., Mol Aspects of Medicine). Also modelling costs and survival of several treatment strategies of CRC was published (Joranger et al., Eur J Health Econ).

International Collaboration

- USA

CANADA

PORTUGAL

SPAIN

FRANCE

UNITED KINGDOM

GERMANY

ITALY

DENMARK

NORWAY

SWEDEN

FINLAND

POLAND

AUSTRIA
- GREECE

AUSTRALIA

ICELAND

IRELAND

THE NETHERLANDS

BELGIUM

SWITZERLAND

CZECH REPUBLIC

HUNGARY

CROATIA

INDIA

SINGAPORE

ISRAEL

RUSSIA

TUNISIA

- AUSTRALIA**
 - Garvan Institute, Sydney
 - Kinghorn Cancer Centre, Sydney
 - Monash University, Melbourne
- AUSTRIA**
 - Medical University of Vienna, Vienna

- BELGIUM**
 - Catholic university of Brussels, Brussels
 - Ghent University, Ghent
 - Katholieke University Leuven, Leuven
 - Universiteit Hasselt, Genk

- CANADA**
 - McGill University, Montreal
 - Princess Margaret Hospital, Toronto
 - University of Ottawa, Ottawa

- CROATIA**
 - University of Zagreb, Zagreb

- CZECH REPUBLIC**
 - Charles University, Prague
 - Institute of Experimental Biology, Masaryk University, Brno
 - Institute of Molecular Genetics, Academy of Sciences of the Czech Republic, Prague
 - National Institute of Public Health, Prague

- DENMARK**
 - Aalborg University Hospital, Aalborg
 - Aarhus University Hospital, Aarhus
 - Copenhagen University Hospital, Copenhagen
 - University of Copenhagen, Copenhagen
 - University of Southern Denmark, Odense

- FINLAND**
 - Biomedicum Helsinki, University of Helsinki, Helsinki
 - Finnish Institute of Molecular Medicine, Nordic EMBL partner, Helsinki
 - Tampere University of Technology, Tampere
 - Zora Oy, Espoo

- FRANCE**
 - Centre National de Génotypage, Paris
 - EurOPDX - European Consortium on Patient-derived Xenografts, Paris
 - Institut Gustave Roussy, Paris
 - Institut National de la Sante et de la Recherche Medicale, Paris
 - Institute Cûrie, Paris
 - Institute of Systems and Synthetic Biology Genopole, UEVE, CNRS, Évry
 - International Agency for Research on Cancer (IARC), Lyon
 - Université Lyon, Villeurbanne
 - Université Paris-Sud, Orsay

- GERMANY**
 - EMBL, Heidelberg
 - Institut für Biochemie, University of Stuttgart, Stuttgart
 - Jacobs University, Bremen
 - University of Bayreuth, Bayreuth
 - University of Bochum, Bochum
 - University of Cologne, Cologne
 - University of Freiburg, Freiburg
 - University of Heidelberg, Heidelberg
 - University of Mainz, Mainz
 - University of Marburg, Marburg

- GREECE**
 - National and Kapodistrian University of Athens, Athens
 - National Centre for Scientific Research "Demokritos", Athens
 - University of Ioannina, Ioannina

- HUNGARY**
 - University of Szeged, Szeged
- ICELAND**
 - University of Iceland, Biomedical Center, Reykjavik

- INDIA**
 - Indian institute of Technology, Hyderabad
 - Savitribai Phule Pune University, Pune

- IRELAND**
 - National Institute for Bioprocessing Research and Training (NIBRT), Dublin

- ISRAEL**
 - Technion - Israel Institute of Technology, Haifa
 - Weizmann Institute, Rehovot

- ITALY**
 - IFOM, Milan
 - International School for Advanced Studies, Trieste
 - Istituto Nazionale di Tumori, Milano
 - The Rizzoli Institute, Bologna
 - University of Bologna, Bologna
 - University of Padova, Padova
 - University of Salento, Lecce

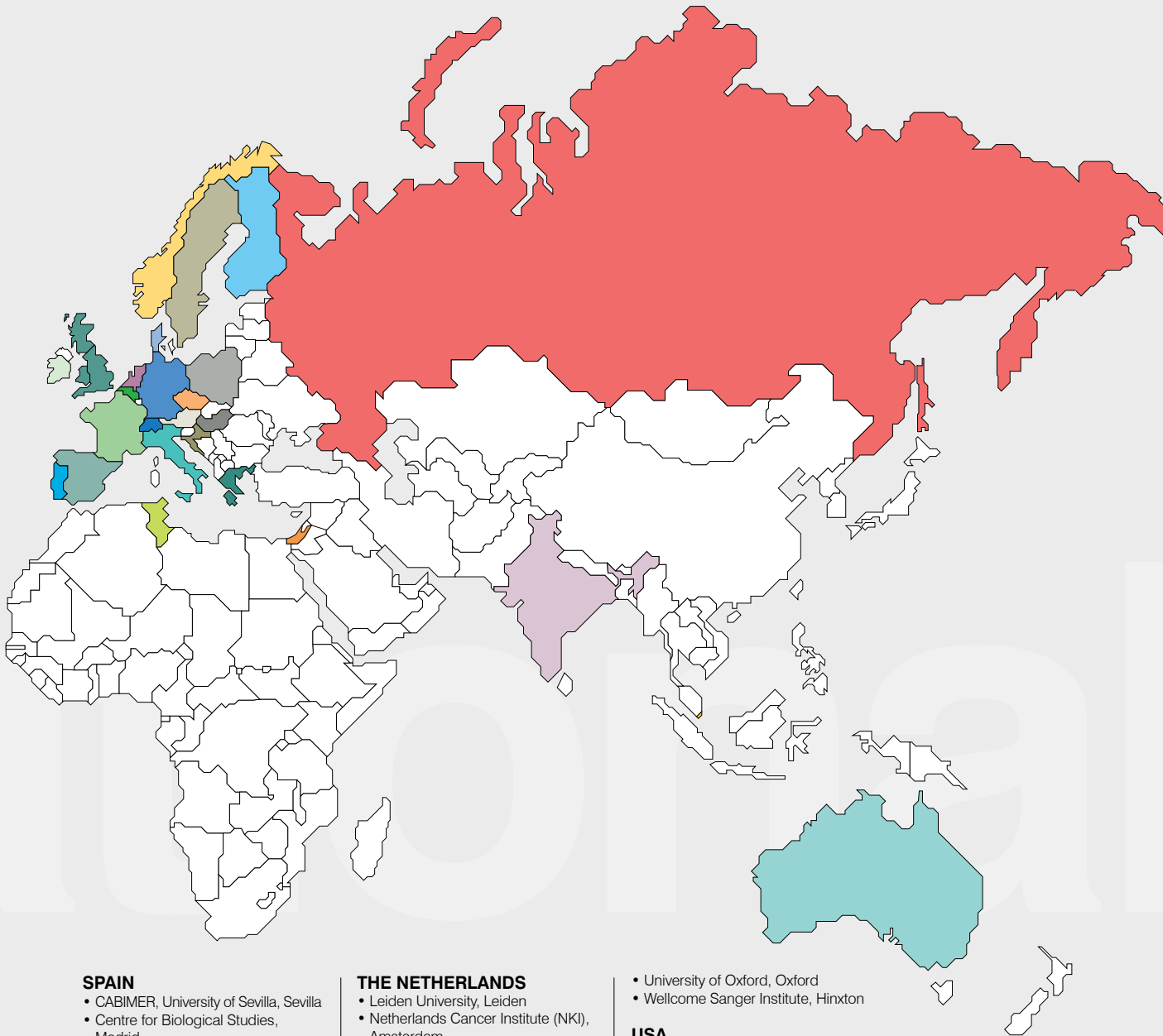
- NORWAY**
 - Cancer Registry of Norway, Oslo
 - Haukeland University Hospital, Bergen
 - Norwegian University of Life Sciences, Ås
 - Norwegian University of Science and Technology, Trondheim
 - Stavanger University Hospital, Stavanger
 - Trondheim University Hospital- St. Olavs Hospital, Trondheim
 - University hospital of North Norway, Tromsø
 - University of Bergen, Bergen
 - University of Oslo, Oslo

- POLAND**
 - Faculty of Biotechnology, University of Wroclaw, Wroclaw
 - Jagiellonian University, Kraków
 - University of Gdansk, Gdansk

- PORTUGAL**
 - Institute of Molecular Pathology and Immunology, University of Porto
 - Portuguese Oncology Institute, Porto

- RUSSIA**
 - Institute of Cytology and Genetics, Novosibirsk

- SINGAPORE**
 - Cancer Science Institute of Singapore, Singapore



- SPAIN**
 - CABIMER, University of Sevilla, Sevilla
 - Centre for Biological Studies, Madrid
 - Fundacion Instituto Valenciano de Oncologica (FIVO), Valencia
 - ICGC, Technical validation group and Ivo Gut, Barcelona
 - University of Lleida, Lleida
 - University of Valencia, Valencia
 - Universitat Politècnica de València, Valencia
 - Vall d'Hebron Institute of Oncology, Barcelona

- SWEDEN**
 - Karolinska Institutet and University of Stockholm, Stockholm
 - Lund University, Lund
 - The Sahlgrenska Academy at the University of Gothenburg, Gothenburg
 - Uppsala University Hospital, Uppsala

- SWITZERLAND**
 - University Hospital Zurich, Zurich

- THE NETHERLANDS**
 - Leiden University, Leiden
 - Netherlands Cancer Institute (NKI), Amsterdam
 - Radboud University Nijmegen, Nijmegen
 - The Netherlands Proteomics Centre, Utrecht
 - University Medical Center, Groningen
 - VU Medical Center, Amsterdam

- TUNISIA**
 - University of Tunis, Tunis

- UNITED KINGDOM**
 - Cambridge Cancer Institute, Cambridge
 - Hampshire Hospitals/Southampton University, Southampton
 - London Research Institute, The Francis Crick Institute, London
 - Royal National Orthopaedic Hospital, Stanmore, Middlesex
 - The Beatson Institute for Cancer Research, Glasgow
 - The European Bioinformatics Institute (EMBL-EBI), Hinxton
 - University College London Medical School, UCL, London
 - University of Cambridge, Cambridge
 - University of Liverpool, Liverpool

- University of Oxford, Oxford
 - Wellcome Sanger Institute, Hinxton

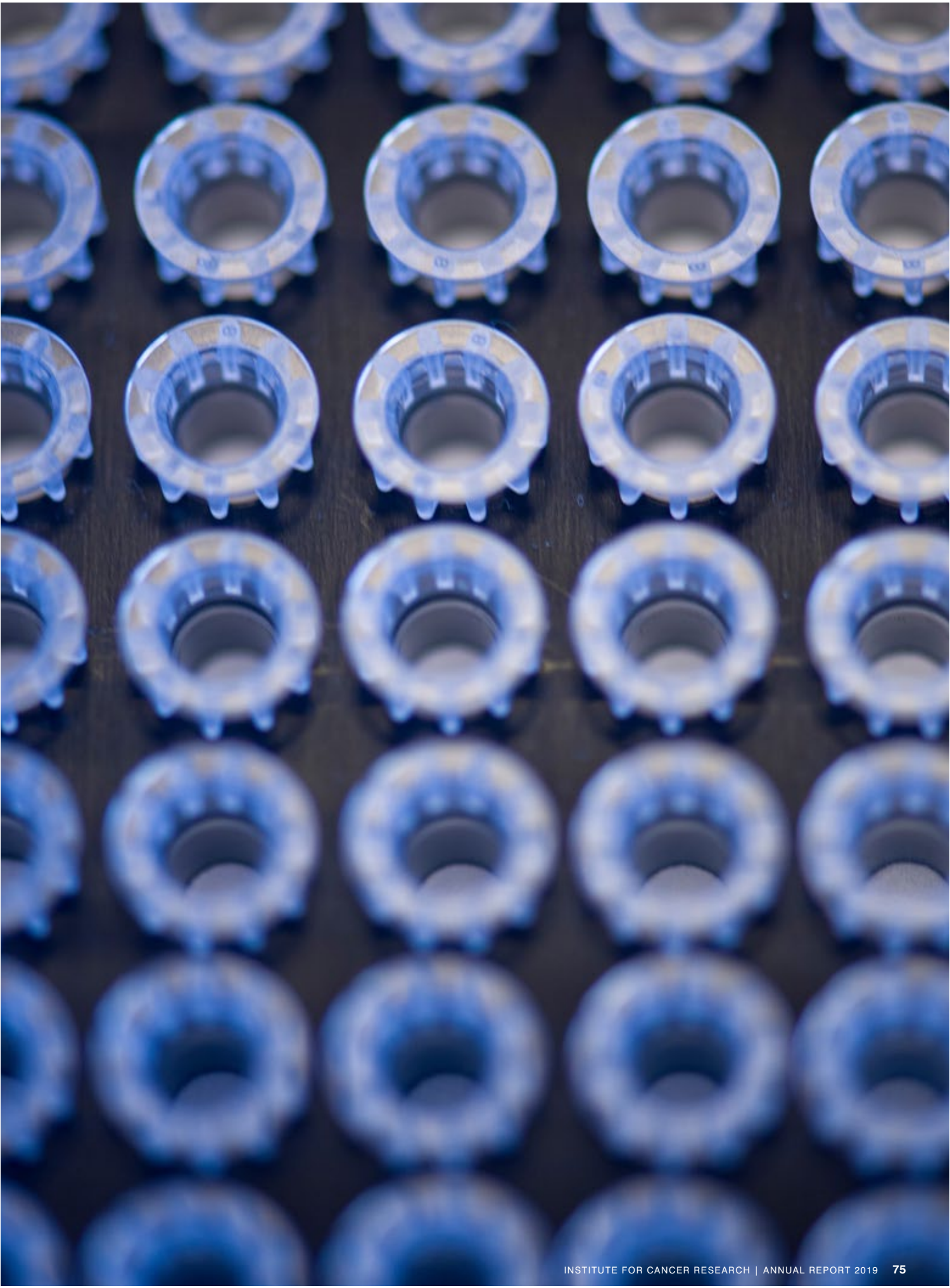
- USA**
 - Buck Institute for Research on Aging, Novato, California
 - Dana Farber Cancer Institute, Boston, Massachusetts
 - Dartmouth College, Hanover, New Hampshire
 - Duke University Medical Center, Durham, North Carolina
 - Fred Hutchinson Cancer Research Center, Seattle, Washington
 - Georgetown University, Washington DC
 - Harvard University, Boston, Massachusetts
 - Johns Hopkins Medicine, Baltimore, Maryland
 - Lawrence Berkeley National Laboratory, Berkeley, California
 - Lineberger Comprehensive Cancer Center, Chapel Hill, North Carolina
 - Masonic Cancer Center and University of Minnesota, Minneapolis
 - Massachusetts General Hospital, Boston, Massachusetts
 - MD Anderson Comprehensive Cancer Center, Houston, Texas
 - National Institutes of Health (NIH), Bethesda, Maryland
 - Oregon State University, Corvallis, Oregon
 - Princeton University, New Jersey
 - Rutgers Cancer Institute of New Jersey
 - Stanford University, California
 - The Mount Sinai Hospital, New York
 - The University of Kansas Hospital, Kansas
 - Tisch Cancer Institute, New York
 - UCSF, Helen Diller Family Cancer Centre, San Francisco, California
 - University of Albany, New York
 - University of California, Berkeley, California
 - University of Chicago, Illinois
 - University of Colorado, Denver, Colorado
 - University of Illinois, Champaign, Illinois
 - University of Washington, Seattle, Washington
 - Washington University, St Louis, Missouri
 - Weill Medical College of Cornell University, New York

Recent Innovations

Registered Disclosures of Invention (DOFIs),
Patent Applications and Granted Patents

Research group (inventor)	Department	DOFI #
Karl-Johan Malmberg	Cancer Immunology	19110
Kristian Berg	Radiation Biology	19075
Vessela Kristensen	Cancer Genetics	19074
Karl-Johan Malmberg	Cancer Immunology	19055
Kjersti Flatmark	Tumor Biology	19054
Therese Sørlie (Jens Henrik Norum)	Cancer Genetics	19050
Kjetil Taskén	Cancer Immunology	19038
Kristian Berg (Anette Weyergang)	Radiation Biology	19035
June Myklebust	Cancer Immunology	19021
Kjersti Flatmark	Tumor Biology	19033

Application type or granted patent	Research group(s)	Department	DOFI #
Priority Application	Kjetil Taskén	Cancer Immunology	19038
Priority Application	Kjersti Flatmark	Tumor Biology	17179
PCT Application	G. M. Mælandsmo, K. Flatmark, K. Sandvig	Tumor Biology, Molecular Cell Biology	18030
US (National) Application	Guro E. Lind	Molecular Oncology	17135
US (National) Application	Karl-Johan Malmberg	Cancer Immunology	16132
EP (Regional) Application	Karl-Johan Malmberg	Cancer Immunology	16132
US (National) Application	Kjersti Flatmark	Tumor Biology	18153
Granted EP patent (EP 3,069,138)	Johanna Olweus	Cancer Immunology	13003
Granted US patent (US 10,308,980)	Lothe R. A., Ågesen T.H., Sveen A., Lind G.E., Nesbakken A., Skotheim R.I.	Molecular Oncology	11057
Granted EP patent (EP 2,630,261)	Guro E. Lind	Molecular Oncology	53212
Granted EP patent (EP 3,243,077)	Kristin Austlid Taskén, Ingrid Jenny Guldvik	Tumor Biology	14040 15004
Granted EP patent (EP 2,861,568)	Kjetil Taskén	Cancer Immunology	12047



Publications

Publications 2019

Abrahamsson H, Porojnicu AC, Lindstrøm JC, Dueland S, **Flatmark K**, Hole KH, Seierstad T, Moan J, Redalen KR, Meltzer S, Ree AH (2019) **High level of circulating vitamin D during neoadjuvant therapy may lower risk of metastatic progression in high-risk rectal cancer** BMC Cancer, 19 (1), 488

Abravan A, **Eide HA**, Løndalen AM, **Helland Å**, Malinen E (2019) **Mapping Bone Marrow Response in the Vertebral Column by Positron Emission Tomography Following Radiotherapy and Erlotinib Therapy of Lung Cancer** Mol Imaging Biol, 21 (2), 391-398

Ackermann F, **Schink KO**, Bruns C, Izsvák Z, Hamra FK, Rosenmund C, Garner CC (2019) **Critical role for Piccolo in synaptic vesicle retrieval** Elife, 8

Aghayan DL, Fretland ÅA, Kazaryan AM, Sahakyan MA, **Dagenborg VJ**, Bjørnbeth BA, **Flatmark K**, Kristiansen R, Edwin B (2019) **Laparoscopic versus open liver resection in the posterosuperior segments: a sub-group analysis from the OSLO-COMET randomized controlled trial** HPB (Oxford), 21 (11), 1485-1490

Alcala N, Leblay N, Gabriel AAG, Mangiante L, Hervas D, Giffon T, Sertier AS, Ferrari A, Derks J, Ghantous A, Delhomme TM, Chabrier A, Cuenin C, Abedi-Ardekani B, Boland A, Olaso R, Meyer V, Altmüller J, Le Calvez-Kelm F, Durand G, Voegele C, Boyault S, Moonen L, Lemaître N, **Brustugun OT**, Lorimier P et al. (2019) **Integrative and comparative genomic analyses identify clinically relevant pulmonary carcinoid groups and unveil the supra-carcinoids** Nat Commun, 10 (1), 3407

Ali M, Foldvari Z, Giannakopoulou E, Böschen ML, Strønen E, Yang W, Toebes M, Schubert B, Kohlbacher O,

Schumacher TN, **Olweus J** (2019) **Induction of neoantigen-reactive T cells from healthy donors** Nat Protoc, 14 (6), 1926-1943

Anda S, Grallert B (2019) **Cell-Cycle-Dependent Regulation of Translation: New Interpretations of Old Observations in Light of New Approaches** Bioessays, 41 (8), e1900022

Andersen E, Chollet ME, Baroni M, Pinotti M, Bernardi F, **Skarpen E**, Sandset PM, Skretting G (2019) **The effect of the chemical chaperone 4-phenylbutyrate on secretion and activity of the p.Q160R mis-sense variant of coagulation factor FVII** Cell Biosci, 9, 69

Andersson Y, Inderberg EM, Kvalheim G, Herud TM, Engebraaten O, Flatmark K, Dueland S, Fodstad Ø (2019) **Immune stimulatory effect of anti-EpCAM immunotoxin - improved overall survival of metastatic colorectal cancer patients** Acta Oncol, 26:1-6

Bains SJ, Abrahamsson H, **Flatmark K**, Dueland S, Hole KH, Seierstad T, Redalen KR, Meltzer S, Ree AH (2019) **Immunogenic cell death by neoadjuvant oxaliplatin and radiation protects against metastatic failure in high-risk rectal cancer** Cancer Immunol Immunother (in press)

Barkovskaya A, Seip K, Hilmarsdottir B, Maelandsmo GM, Moestue SA, Itkonen HM (2019) **O-GlcNAc Transferase Inhibition Differentially Affects Breast Cancer Subtypes** Sci Rep, 9 (1), 5670

Bartolomé-Casado R, Landsverk OJB, Chauhan SK, Richter L, Phung D, Greiff V, Risnes LF, Yao Y, Neumann RS, Yaqub S, Øyen O, Horneland R, **Aandahl EM**, Paulsen V, Sollid LM, Qiao SW, Baekkeveld ES, Jahnsen FL (2019) **Resident memory CD8 T cells persist for years in human small intestine** J Exp Med, 216 (10), 2412-2426

Berg KCG, Sveen A, Høland M, Alagaratnam S, Berg M, Danielsen SA, Nesbakken A, Søreide K, **Lothe RA** (2019) **Gene expression profiles of CMS2-epithelial/canonical colorectal cancers are largely driven by DNA copy number gains** Oncogene, 38 (33), 6109-6122

Berge LAM, Andreassen BK, Stenehjem JS, Larsen IK, Furu K, **Juzeniene A**, Roscher I, Heir T, Green A, Veierød MB, Robsahm TE (2019) **Cardiovascular, antidepressant and immunosuppressive drug use in relation to risk of cutaneous melanoma: a protocol for a prospective case-control study** BMJ Open, 9 (2), e025246

Bergholtz H, Lien TG, Ursin G, Holmen MM, **Helland Å**, Sørlie T, Haakensen VD (2019) **A Longitudinal Study of the Association between Mammographic Density and Gene Expression in Normal Breast Tissue** J Mammary Gland Biol Neoplasia, 24 (2), 163-175

Blin M, Le Tallec B, **Nähse V**, Schmidt M, Brossas C, Millot GA, Prioleau MN, Debatisse M (2019) **Transcription-dependent regulation of replication dynamics modulates genome stability** Nature Structural & Molecular Biology 26, 58–66

Bowitz Lothe IM, Kleive D, Pomianowska E, Cvancarova M, **Kure E**, Dueland S, Gladhaug IP, Labori KJ (2019) **Clinical relevance of pancreaticobiliary and intestinal subtypes of ampullary and duodenal adenocarcinoma: Pattern of recurrence, chemotherapy, and survival after pancreatoduodenectomy** Pancreatology, 19 (2), 316-324

Boye E, Grallert B (2019) **eIF2 phosphorylation and the regulation of translation** Curr Genet (in press)

Brennecke P, Rasina D, Aubi O, Herzog K, Landskron J, Cautain B, Vicente F,

Quintana J, Mestres J, Stechmann B, Ellinger B, Brea J, Kolanowski JL, Pilarski R, Orzaez M, Pineda-Lucena A, Laraia L, Nami F, Zielenkiewicz P, Paruch K, Hansen E, von Kries JP, Neuenschwander M, Specker E, Bartunek P et al. (2019) **EU-OPENSCREEN: A Novel Collaborative Approach to Facilitate Chemical Biology** SLAS Discov, 24 (3), 398-413

Briem E, Ingthorsson S, Traustadottir GA, **Hilmarsdottir B**, Gudjonsson T (2019) **Application of the D492 Cell Lines to Explore Breast Morphogenesis, EMT and Cancer Progression in 3D Culture** J Mammary Gland Biol Neoplasia, 24 (2), 139-147

Brinkman AB, Nik-Zainal S, Simmer F, Rodríguez-González FG, Smid M, Alexandrov LB, Butler A, Martin S, Davies H, Glodzik D, Zou X, Ramakrishna M, Staaf J, Ringnér M, Sieuwerts A, Ferrari A, Morganella S, **Fleischer T**, Kristensen V, Gut M, van de Vijver MJ, **Børresen-Dale AL**, Richardson AL, Thomas G, Gut IG et al. (2019) **Partially methylated domains are hypervariable in breast cancer and fuel widespread CpG island hypermethylation** Nat Commun, 10 (1), 1749

Brison O, El-Hilali S, Azar D, Koundrioukoff S, Schmidt M, **Nähse V**, Jaszczyszyn Y, Lachages AM, Dutrillaux B, Thermes C, Debatisse M, Chen CL (2019) **Transcription-mediated organization of the replication initiation program across large genes sets common fragile sites genome-wide** Nat Commun, 10 (1), 5693

Brudvik KW, Jones RP, Giulianti F, Shindoh J, Passot G, Chung MH, Song J, Li L, **Dagenborg VJ**, Fretland ÅA, Røsok B, De Rose AM, Ardito F, Edwin B, Panettieri E, Larocca LM, Yamashita S, Conrad C, Aloia TA, Poston GJ, Bjørnbeth BA, Vauthey JN (2019) **RAS Mutation Clinical Risk Score to Predict Survival After Resection of Colorectal Liver Metastases** Ann Surg, 269 (1), 120-126

Brunsell TH, Cengija V, Sveen A, Bjørnbeth BA, Røsok BI, Brudvik KW, Guren MG, **Lothe RA**, Abildgaard A, Nesbakken A (2019)

Heterogeneous radiological response to neoadjuvant therapy is associated with poor prognosis after resection of colorectal liver metastases Eur J Surg Oncol, 45 (12), 2340-2346

Brunsell TH, Sveen A, Bjørnbeth BA, Røsok BI, **Danielsen SA**, Brudvik KW, **Berg KCG**, Johannessen B, Cengija V, Abildgaard A, Guren MG, Nesbakken A, **Lothe RA** (2019) **High Concordance and Negative Prognostic Impact of RAS/BRAF/PIK3CA Mutations in Multiple Resected Colorectal Liver Metastases** Clin Colorectal Cancer (in press)

Brynildsen J, Petäjä L, Myhre PL, Lyngbakken MN, **Nygård S**, Stridsberg M, Christensen G, Ottesen AH, Pettilä V, Omrand T, Røsjø H (2019) **Circulating Secretoneurin Concentrations After Cardiac Surgery: Data From the FINNish Acute Kidney Injury Heart Study** Crit Care Med, 47 (5), e412-e419

Braadland PR, Ramberg H, Grytli HH, Urbanucci A, Nielsen HK, Guldvik IJ, Engedal A, Ketola K, Wang W, Svindland A, Mills IG, Bjartell A, **Taskén KA** (2019) **The β₂-Adrenergic Receptor Is a Molecular Switch for Neuroendocrine Transdifferentiation of Prostate Cancer Cells** Mol Cancer Res, 17 (11), 2154-2168

Braadland PR, Urbanucci A (2019) **Chromatin reprogramming as an adaptation mechanism in advanced prostate cancer** Endocr Relat Cancer, 26 (4), R211-R235

Burocziova M, Burdova K, Martinikova AS, Kasperek P, Kleiblova P, **Danielsen SA**, Borecka M, Jenikova G, Janečková L, Pavel J, Zemankova P, Schneiderova M, Schwarzova L, Ticha I, Sun XF, Jiraskova K, Liska V, Vodickova L, Vodicka P, Sedlacek R, Kleibl Z, **Lothe RA**, Korinek V, Macurek L (2019) **Truncated PPM1D impairs stem cell response to genotoxic stress and promotes growth of APC-deficient tumors in the mouse colon** Cell Death Dis, 10 (11), 818

Calvete J, Larrinaga G, Errarte P, Martín AM, Dotor A, Esquinas C, **Nunes-Xavier CE**, Pulido R, López JI, Angulo JC (2019) **The coexpression of fibroblast acti-**

vation protein (FAP) and basal-type markers (CK 5/6 and CD44) predicts prognosis in high-grade invasive urothelial carcinoma of the bladder Hum Pathol, 91, 61-68

Camilio KA, Wang MY, Mauseth B, Waagene S, Kvalheim G, Rekdal Ø, Sveinbjørnsson B, **Mælandsmo GM** (2019) **Combining the oncolytic peptide LTX-315 with doxorubicin demonstrates therapeutic potential in a triple-negative breast cancer model** Breast Cancer Res, 21 (1), 9

Carm KT, Hoff AM, Bakken AC, Axcrona U, Axcrona K, **Lothe RA**, Skotheim RI, Løvf M (2019) **Interfocal heterogeneity challenges the clinical usefulness of molecular classification of primary prostate cancer** Sci Rep, 9 (1), 13579

Chellappa S, Kushekhar K, Munthe LA, Tjønnfjord GE, **Aandahl EM**, Okkenhaug K, **Taskén K** (2019) **The PI3K p110δ Isoform Inhibitor Idelalisib Preferentially Inhibits Human Regulatory T Cell Function** J Immunol, 202 (5), 1397-1405

Cornillet M, Jansson H, Schaffer M, Hertwig L, Berglin L, Zimmer CL, Johansson H, Ellis E, Isaksson B, Gonzalez-Galarza FF, Middleton D, **Malmberg KJ**, Sparrelid E, Björkström NK (2019) **Imbalance of Genes Encoding Natural Killer Immunoglobulin-Like Receptors and Human Leukocyte Antigen in Patients With Biliary Cancer** Gastroenterology, 157 (4), 1067-1080. e9

Cremaschi A, Argiento R, Shoemaker K, Peterson C, Vannucci M (2019) **Hierarchical Normalized Completely Random Measures for Robust Graphical Modeling** Bayesian Anal., 14 (4), 1271-1301

Crosbie EJ, Ryan NAJ, Arends MJ, Bosse T, Burn J, Cornes JM, Crawford R, Eccles D, Frayling IM, Ghaem-Maghani S, Hampel H, Kauff ND, Kitchener HC, Kitson SJ, Manchanda R, McMahon RFT, Monahan KJ, Menon U, **Møller P**, Möslein G, Rosenthal A, Sasieni P, Seif MW, Singh N, Skarrott P et al. (2019) **The Manchester International Consensus Group recommendations for the management of gynecological**

Publications

cancers in Lynch syndrome
Genet Med, 21 (10), 2390-2400

Debik J, Euceda LR, Lundgren S, Gythfeldt HVL, **Garred Ø, Borgen E, Engebraaten O**, Bathen TF, Giskeødegård GF (2019)
Assessing Treatment Response and Prognosis by Serum and Tissue Metabolomics in Breast Cancer Patients
J Proteome Res, 18 (10), 3649-3660

Della Valle A, Rossi BM, Palmero EI, Antelo M, Vaccaro CA, López-Kostner F, Alvarez K, Cruz-Correa M, Bruno LI, Forones NM, Mindiola JAR, Buleje J, Spirandelli F, Bohorquez M, Cock-Rada AM, Sulcahuan Y, Nascimento I, Abe-Sandes K, Lino-Silva LS, Petracchi F, Mampel A, Rodriguez Y, Rossi NT, Yañez CB, Rubio C et al. (2019)
A snapshot of current genetic testing practice in Lynch syndrome: The results of a representative survey of 33 Latin American existing centres/registries
Eur J Cancer, 119, 112-121

Dienstmann R, Villacampa G, **Sveen A**, Mason MJ, Niedzwiecki D, Nesbakken A, Moreno V, Warren RS, **Lothe RA**, Guinney J (2019)
Relative contribution of clinico-pathological variables, genomic markers, transcriptomic subtyping and microenvironment features for outcome prediction in stage II/III colorectal cancer
Ann Oncol, 30 (10), 1622-1629

Dominguez-Valentin M, Nakken S, Tubeuf H, **Vodak D, Ekstrøm PO**, Nissen AM, Morak M, Holinski-Feder E, Holth A, Capella G, Davidson B, Evans DG, Martins A, **Møller P, Hovig E** (2019)
Results of multigene panel testing in familial cancer cases without genetic cause demonstrated by single gene testing
Sci Rep, 9 (1), 18555

Dominguez-Valentin M, Sampson JR, Seppälä TT, Ten Broeke SW, Plazzer JP, **Nakken S**, Engel C, Aretz S, Jenkins MA, Sunde L, Bernstein I, Capella G, Balaguer F, Thomas H, Evans DG, Burn J, Greenblatt M, **Hovig E**, de Vos Tot Nederveen Cappel WH, Sijmons RH, Bertario L, Tibiletti MG, Cavestro GM, Lindblom A, Della Valle A et al. (2019)
Cancer risks by gene, age, and gender in 6350 carriers of pathogenic

mismatch repair variants: findings from the Prospective Lynch Syndrome Database
Genet Med, 22 (1), 15-25

Dominguez-Valentin M, Seppälä TT, Sampson JR, Macrae F, Winship I, Evans DG, Scott RJ, Burn J, Möslain G, Bernstein I, Pylvänäinen K, Renkonen-Sinisalo L, Lepistö A, Lindblom A, Plazzer JP, Tjandra D, Thomas H, Green K, Lalloo F, Crosbie EJ, Hill J, Capella G, Pineda M, Navarro M, Vidal JB et al. (2019)
Survival by colon cancer stage and screening interval in Lynch syndrome: a prospective Lynch syndrome database report
Hered Cancer Clin Pract, 17, 28

Dong X, Zhang R, He J, Lai L, Alolga RN, Shen S, Zhu Y, You D, Lin L, Chen C, Zhao Y, Duan W, Su L, Shafer A, Salama M, **Fleischer T, Bjaanæs MM**, Karlsson A, Planck M, Wang R, Staaf J, **Helland Å**, Esteller M, Wei Y, Chen F et al. (2019)
Trans-omics biomarker model improves prognostic prediction accuracy for early-stage lung adenocarcinoma
Aging (Albany NY), 11 (16), 6312-6335

Dörk T, Peterlongo P, Mannermaa A, Bolla MK, Wang Q, Dennis J, Ahearn T, Andrulis IL, Anton-Culver H, Arndt V, Aronson KJ, Augustinsson A, Freeman LEB, Beckmann MW, Beeghly-Fadiel A, Behrens S, Bermisheva M, Blomqvist C, Bogdanova NV, Bojesen SE, Brauch H, Brenner H, Burwinkel B, Canzian F, **Kristensen VN**, Chan TL, et al. (2019)
Two truncating variants in FANCC and breast cancer risk
Sci Rep, 9 (1), 12524

Eide PW, Eilertsen IA, Sveen A, Lothe RA (2019)
Long noncoding RNA MIR31HG is a bona fide prognostic marker with colorectal cancer cell-intrinsic properties
Int J Cancer, 144 (11), 2843-2853

Eilertsen IA, Sveen A, Strømme JM, Skotheim RI, Nesbakken A, **Lothe RA** (2018)
Alternative splicing expands the prognostic impact of KRAS in microsatellite stable primary colorectal cancer
Int J Cancer, 144 (4), 841-847

Eilertsen IA, Moosavi SH, Strømme

JM, Nesbakken A, **Johannessen B, Lothe RA, Sveen A** (2019) (Epub ahead of print)
Technical differences between sequencing and microarray platforms impact transcriptomic subtyping of colorectal cancer
Cancer Lett, 469, 246-255

Enqvist M, **Jacobs B**, Junlén HR, Schaffer M, Melén CM, Friberg D, Wahlin BE, **Malmberg KJ** (2019)
Systemic and Intra-Nodal Activation of NK Cells After Rituximab Monotherapy for Follicular Lymphoma
Front Immunol, 10, 2085

Escala-Garcia M, Abraham J, Andrulis IL, Anton-Culver H, Arndt V, Ashworth A, Auer PL, Auvinen P, Beckmann MW, Beesley J, Behrens S, Benitez J, Bermisheva M, Blomqvist C, Blot W, Bogdanova NV, Bojesen SE, Bolla MK, **Børresen-Dale AL**, Brauch H, Brenner H, Brucker SY, Burwinkel B, Caldas C, Canzian F et al. (2020)
A network analysis to identify mediators of germline-driven differences in breast cancer prognosis
Nat Commun, 11 (1), 312

Eriksson AG, Fallaas Dahl G, Nesbakken AJ, **Lund KV**, Amant F (2019)
Endometrial cancer during pregnancy: management strategies
Int J Gynecol Cancer, 29 (7), 1221-1224

Escala-Garcia M, Guo Q, Dörk T, Canisius S, Keeman R, Dennis J, Beesley J, Lecarpentier J, Bolla MK, Wang Q, Abraham J, Andrulis IL, Anton-Culver H, Arndt V, Auer PL, Beckmann MW, Behrens S, Benitez J, Bermisheva M, Bernstein L, Blomqvist C, Boeckx B, Bojesen SE, Bonanni B, **Børresen-Dale AL** et al. (2019)
Genome-wide association study of germline variants and breast cancer-specific mortality
Br J Cancer, 120 (6), 647-657

Fang EF, Hou Y, Lautrup S, Jensen MB, Yang B, SenGupta T, Caponio D, **Khezri R**, Demarest TG, Aman Y, Figueroa D, Morevati M, Lee HJ, Kato H, Kassahun H, Lee JH, Filippelli D, Okur MN, Mangerich A, Croteau DL, Maezawa Y, Lyssiotis CA, Tao J, Yokote K, **Rusten TE** et al. (2019)
NAD⁺ augmentation restores mitophagy and limits accelerated aging in Werner syndrome
Nat Commun, 10 (1), 5284

Farmer JR, **Foldvari Z**, Ujhazi B, De Ravin SS, Chen K, Bleesing JJH, Schuetz C, Al-Herz W, Abraham RS, Joshi AY, Costa-Carvalho BT, Buchbind-er D, Booth C, Reiff A, Ferguson PJ, Aghamohammadi A, Abolhassani H, Puck JM, Adeli M, Cancrini C, Palma P, Bertaina A, Locatelli F, Di Matteo G, Geha RS et al. (2019)
Outcomes and Treatment Strategies for Autoimmunity and Hyperinflammation in Patients with RAG Deficiency
J Allergy Clin Immunol Pract, 7 (6), 1970-1985.e4

Ferreira MA, Gamazon ER, Al-Ejeh F, Aittomäki K, Andrulis IL, Anton-Culver H, Arason A, Arndt V, Aronson KJ, Arun BK, Asseryanis E, Azzollini J, Balmaña J, Barnes DR, Barrowdale D, Beckmann MW, Behrens S, Benitez J, Bermisheva M, Białkowska K, Blomqvist C, Bogdanova NV, Bojesen SE, Bolla MK, **Kristensen VN** Borg A et al. (2019)
Genome-wide association and transcriptome studies identify target genes and risk loci for breast cancer
Nat Commun, 10 (1), 1741

Flem-Karlsen K, Fodstad Y, Nunes-Xavier CE (2019)
B7-H3 immune checkpoint protein in human cancer
Curr Med Chem (in press)

Flem-Karlsen K, McFadden E, **Omar N, Haugen MH, Øy GF**, Ryder T, Gullestad HP, Hermann R, **Mælandsmo GM**, Flørenes VA (2019)
Targeting AXL and the DNA damage response pathway as a novel therapeutic strategy in melanoma
Mol Cancer Ther (in press)

Flem-Karlsen K, Tekle C, Øyjord T, Flørenes VA, **Mælandsmo GM, Fodstad Ø, Nunes-Xavier CE** (2019)
p38 MAPK activation through B7-H3-mediated DUSP10 repression promotes chemoresistance
Sci Rep, 9 (1), 5839

Flørenes VA, Flem-Karlsen K, McFadden E, **Bergheim IR, Nygaard V, Nygård V**, Farstad IN, **Øy GF**, Emilsen E, **Giller-Fieten K**, Ree AH, **Flatmark K**, Gullestad HP, Hermann R, Ryder T, Wernhoff P, **Mælandsmo GM** (2019)
A Three-dimensional Ex Vivo Viability Assay Reveals a Strong Correlation Between Response to Targeted Inhibitors and Mutation Status in Melanoma Lymph Node Metastases

Transl Oncol, 12 (7), 951-958

Forthun RB, Hovland R, Schuster C, Puntervoll H, Brodal HP, **Namløs HM, Aasheim LB, Meza-Zepeda LA**, Gjertsen BT, Knappskog S, Straume O (2019)
ctDNA detected by ddPCR reveals changes in tumour load in metastatic malignant melanoma treated with bevacizumab
Sci Rep, 9 (1), 17471

Fougner C, Bergholtz H, Kuiper R, **Norum JH, Sørlie T** (2019)
Claudin-low-like mouse mammary tumors show distinct transcriptomic patterns uncoupled from genomic drivers
Breast Cancer Res, 21 (1), 85

Fretland ÅA, **Dagenborg VJ**, Waaler Bjørnelv GM, Aghayan DL, Kazaryan AM, Barkhatov L, Kristiansen R, Fagerland MW, Edwin B, Andersen MH (2019)
Quality of life from a randomized trial of laparoscopic or open liver resection for colorectal liver metastases
Br J Surg, 106 (10), 1372-1380

Frikstad KM, Molinari E, **Thoresen M**, Ramsbottom SA, Hughes F, Letteboer SJF, **Gilani S, Schink KO, Stokke T**, Geimer S, Pedersen LB, Giles RH, Akhmanova A, Roepman R, Sayer JA, **Patzke S** (2019)
A CEP104-CSPP1 Complex Is Required for Formation of Primary Cilia Competent in Hedgehog Signaling
Cell Rep, 28 (7), 1907-1922.e6

Frøysnes IS, Andersson Y, Larsen SG, Davidson B, **Øien JT**, Julsrud L, **Fodstad Ø**, Dueland S, **Flatmark K** (2019)
ImmunoPeCa trial: Long-term outcome following intraperitoneal MOC31PE immunotoxin treatment in colorectal peritoneal metastasis
Eur J Surg Oncol (in press)

Gaustad JV, Simonsen TG, Wegner CS, Rofstad EK (2019)
Vascularization, Oxygenation, and the Effect of Sunitinib Treatment in Pancreatic Ductal Adenocarcinoma Xenografts
Front Oncol, 9, 845

GBD 2016 Stroke Collaborators (2019)
Global, regional, and national burden of stroke, 1990-2016: a systematic analysis for the Global Burden

of Disease Study 2016
Lancet Neurol, 18 (5), 439-458

Gheorghe M, Sandve GK, Khan A, Chèneby J, Ballester B, **Mathelier A** (2019)
A map of direct TF-DNA interactions in the human genome
Nucleic Acids Res, 47 (4), e21

Glaire MA, Domingo E, **Sveen A, Bruun J**, Nesbakken A, Nicholson G, Novelli M, Lawson K, Oukrif D, Kildal W, Danielsen HE, Kerr R, Kerr D, Tomlinson I, **Lothe RA**, Church DN (2019)
Tumour-infiltrating CD8⁺ lymphocytes and colorectal cancer recurrence by tumour and nodal stage
Br J Cancer, 121 (6), 474-482

Global Burden of Disease Cancer Collaboration, Fitzmaurice C, Abate D, Abbasi N, Abbastabar H, Abd-Allah F, Abdel-Rahman O, Abdelalim A, Abdoli A, Abdollahpour I, Abdulle ASM, Abebe ND, Abraha HN, Abu-Raddad LJ, Abualhasan A, Adedeji IA, Advani SM, Afarideh M, Afshari M, Aghaali M, Agius D, Agrawal S, Ahmadi A, Ahmadian E, Ahmadpour E et al. (2019)
Global, Regional, and National Cancer Incidence, Mortality, Years of Life Lost, Years Lived With Disability, and Disability-Adjusted Life-Years for 29 Cancer Groups, 1990 to 2017: A Systematic Analysis for the Global Burden of Disease Study
JAMA Oncol (in press)

Goodridge JP, Jacobs B, Saeter-smoen ML, Clement D, Hammer Q, **Clancy T, Skarpen E, Brech A**, Landskron J, Grimm C, Pfefferle A, **Meza-Zepeda L, Lorenz S, Wiiger MT**, Louch WE, **Ask EH**, Liu LL, **Oei VYS**, Kjällquist U, Linnarsson S, Patel S, **Taskén K, Stenmark H, Malmberg KJ** (2019)
Remodeling of secretory lysosomes during education tunes functional potential in NK cells
Nat Commun, 10 (1), 514

Grigalavicius M, Mastrangelopoulou M, Berg K, Arous D, Ménard M, Raabe-Henriksen T, Brondz E, Siem S, Görgen A, Edin NFJ, Malinen E, **Theodossiou TA** (2019)
Proton-dynamic therapy following photosensitiser activation by accelerated protons demonstrated through fluorescence and singlet oxygen production
Nat Commun, 10 (1), 3986

Publications

Grinde MT, **Hilmarsdottir B**, Tunset HM, Henriksen IM, Kim J, **Haugen MH**, Rye MB, **Mælandsmo GM**, Moestue SA (2019)
Glutamine to proline conversion is associated with response to glutamine inhibition in breast cancer
Breast Cancer Res, 21 (1), 61

Ha TJ, Zhang PGY, Robert R, Yeung J, Swanson DJ, **Mathelier A**, Wasserman WW, Im S, Itoh M, Kawaji H, Lassmann T, Daub CO, Arner E, FANTOM Consortium, Carninci P, Hayashizaki Y, Forrest ARR, Goldowitz D (2019)
Identification of novel cerebellar developmental transcriptional regulators with motif activity analysis
BMC Genomics, 20 (1), 718

Hagberg G, Fure B, Thommessen B, Ihle-Hansen H, Øksengård AR, **Nygård S**, Pendlebury ST, Beyer MK, Wyller TB, Ihle-Hansen H (2019)
Predictors for Favorable Cognitive Outcome Post-Stroke: A-Seven-Year Follow-Up Study
Dement Geriatr Cogn Disord, 48 (1-2), 45-55

Halvorsen AR, Ragle Aure M, Öjlert ÅK, Brustugun OT, Solberg S, Nebdal D, Helland Å (2019)
Identification of microRNAs involved in pathways which characterize the expression subtypes of NSCLC
Mol Oncol, 13 (12), 2604-2615

Hamfjord J, Guren TK, Dajani O, Johansen JS, Glimelius B, Sorbye H, Pfeiffer P, **Lingjærde OC**, Tveit KM, **Kure EH**, Pallisgaard N, Spindler KG (2019)
Total circulating cell-free DNA as a prognostic biomarker in metastatic colorectal cancer before first-line oxaliplatin-based chemotherapy
Ann Oncol, 30 (7), 1088-1095

Handle F, Prekovic S, Helsen C, Van den Broeck T, Smeets E, Moris L, Eerlings R, Kharraz SE, **Urbanucci A**, Mills IG, Joniau S, Attard G, Claessens F (2019)
Drivers of AR indifferent anti-androgen resistance in prostate cancer cells
Sci Rep, 9 (1), 13786

Hanes R, Munthe E, Grad I, Han J, Karlsen I, McCormack E, **Meza-Zepeda LA, Stratford EW, Myklebost O** (2019)
Preclinical Evaluation of the Pan-FG-

FR Inhibitor LY2874455 in FRS2-Amplified Liposarcoma
Cells, 8 (2)

Hansem LMK, Huang R, Wegner CS, Simonsen TG, Gaustad JV, Hauge A, Rofstad EK (2019)
Intratumor Heterogeneity in Interstitial Fluid Pressure in Cervical and Pancreatic Carcinoma Xenografts
Transl Oncol, 12 (8), 1079-1085

Hansen EP, **Fromm B**, Andersen SD, Marcilla A, Andersen KL, Borup A, Williams AR, Jex AR, Gasser RB, Young ND, Hall RS, Stensballe A, Ovchinnikov V, Yan Y, Fredholm M, Thamsborg SM, Nejsum P (2019)
Exploration of extracellular vesicles from Ascaris suum provides evidence of parasite-host cross talk
J Extracell Vesicles, 8 (1), 1578116

Hauge A, Gaustad JV, Huang R, Simonsen TG, Wegner CS, Andersen LMK, Rofstad EK (2019)
DCE-MRI and Quantitative Histology Reveal Enhanced Vessel Maturation but Impaired Perfusion and Increased Hypoxia in Bevacizumab-Treated Cervical Carcinoma
Int J Radiat Oncol Biol Phys, 104 (3), 666-676

Hauge S, Macurek L, **Syljuåsen RG** (2019)
p21 limits S phase DNA damage caused by the Wee1 inhibitor MK1775
Cell Cycle, 18 (8), 834-847

Hausken J, Fretland ÅA, Edwin B, Andersen MH, **Dagenborg VJ**, Bjørnelv GMW, Kristiansen R, Røysland K, Kvarstein G, Tønnessen TI (2019)
Intravenous Patient-controlled Analgesia Versus Thoracic Epidural Analgesia After Open Liver Surgery: A Prospective, Randomized, Controlled, Noninferiority Trial
Ann Surg, 270 (2), 193-199

Hausott B, Förste A, Zach F, Mangger S, **Haugsten EM**, Klimaschewski L (2019)
Endocytosis and Transport of Growth Factor Receptors in Peripheral Axon Regeneration: Novel Lessons from Neurons Expressing Lysine-Deficient FGF Receptor Type 1 in vitro
Anat Rec (Hoboken), 302 (8), 1268-1275

Helgeland H, Sodeland M, Zoric N, Torgersen JS, Grammes F, von Lintig J, Moen T, Kjøglum S, Lien S, Våge DI (2019)
Genomic and functional gene studies suggest a key role of beta-carotene oxygenase 1 like (bco1l) gene in salmon flesh color
Sci Rep, 9 (1), 20061

Hoem G, Bowitz Larsen K, Øvervatn A, **Brech A**, Lamark T, Sjøttem E, Johansen T (2019)
The FMRpolyGlycine Protein Mediates Aggregate Formation and Toxicity Independent of the CGG mRNA Hairpin in a Cellular Model for FXTAS
Front Genet, 10, 249

Holdgaard SG, Cianfanelli V, Pupo E, Lambrughì M, Lubas M, Nielsen JC, Eibes S, Maiani E, Harder LM, Wesch N, Foged MM, Maeda K, Nazio F, de la Ballina LR, Dötsch V, **Brech A**, Frankel LB, Jäättelä M, Locatelli F, Barisic M, Andersen JS, Bekker-Jensen S, Lund AH, Rogov VV, Papaleo E et al. (2019)
Selective autophagy maintains centrosome integrity and accurate mitosis by turnover of centriolar satellites
Nat Commun, 10 (1), 4176

Huse K (2019)
Expanding the Clinical Cytometry Toolbox-Receptor Occupancy by Mass Cytometry
Cytometry A, 95 (10), 1046-1048

Huse K, Wogslund CE, Polikowsky HG, Diggins KE, **Smeland EB, Myklebust JH**, Irish JM (2019)
Human Germinal Center B Cells Differ from Naïve and Memory B Cells in CD40 Expression and CD40L-Induced Signaling Response
Cytometry A, 95 (4), 442-449

Ihle-Hansen H, Vigen T, Berge T, Hagberg G, Engedal K, Rønning OM, Thommessen B, Lyngbakken MN, **Nygård S**, Røsjø H, Tveit A, Ihle-Hansen H (2019)
Carotid Atherosclerosis and Cognitive Function in a General Population Aged 63-65 Years: Data from the Akershus Cardiac Examination (ACE) 1950 Study
J Alzheimers Dis, 70 (4), 1041-1049

Itkonen HM, **Urbanucci A**, Martin SE, Khan A, **Mathelier A**, Thiede B, Walker S, Mills IG (2019)
High OGT activity is essential for

MYC-driven proliferation of prostate cancer cells
Theranostics, 9 (8), 2183-2197

Jacobs B, Pfefferle A, **Clement D, Berg-Larsen A, Saetersmoen ML, Lorenz S, Wiiger MT, Goodridge JP, Malmberg KJ** (2018)
Induction of the BIM Short Splice Variant Sensitizes Proliferating NK Cells to IL-15 Withdrawal
J Immunol, 202 (3), 736-746

Jerjes W, Hamdoon Z, **Berg K**, Høgset A, Hopper C (2019)
Recurrent chondroblastic osteosarcoma of the right mandible subjected to photochemical internalization
Photodiagnosis Photodyn Ther, 27, 288-290

Jiang X, Finucane HK, Schumacher FR, Schmit SL, Tyrer JP, Han Y, Michailidou K, Lesseur C, Kuchenbaecker KB, Dennis J, Conti DV, Casey G, Gaudet MM, Huyghe JR, Albanes D, Aldrich MC, Andrew AS, Andrulis IL, Anton-Culver H, Antoniou AC, Antonenkova NN, Arnold SM, Aronson KJ, Arun BK, **Kristensen VN**, Bandera EV et al. (2019)
Shared heritability and functional enrichment across six solid cancers
Nat Commun, 10 (1), 431

Johansson HJ, Socciarelli F, Vacanti NM, **Haugen MH**, Zhu Y, Siavelis I, Fernandez-Woodbridge A, **Aure MR**, Sennblad B, Vesterlund M, Branca RM, Orre LM, Huss M, Fredlund E, Beraki E, Garred Ø, Boekel J, Sauer T, Zhao W, **Nord S, Höglander EK**, Jans DC, Brismar H, Haukaas TH, Bathen TF et al. (2019)
Breast cancer quantitative proteome and proteogenomic landscape
Nat Commun, 10 (1), 1600

Jonsson M, Fjeldbo CS, Holm R, **Stokke T**, Kristensen GB, **Lyng H** (2019)
Mitochondrial Function of CKS2 Oncoprotein Links Oxidative Phosphorylation with Cell Division in Chemoradioresistant Cervical Cancer
Neoplasia, 21 (4), 353-362

Josefsson SE, Beiske K, **Blaker YN**, Førsum MS, Holte H, Østenstad B, Kimby E, Köksal H, Wälchli S, **Bai B, Smeland EB**, Levy R, Kolstad A, **Huse K, Myklebust JH** (2019)
TIGIT and PD-1 Mark Intratumoral T Cells with Reduced Effector Function in B-cell Non-Hodgkin Lymphoma

Cancer Immunol Res, 7 (3), 355-362

Juraleviciute M, Pozniak J, Nsengimana J, Harland M, Randerson-Moor J, Werhoeff P, Bassarova A, **Øy GF**, Trøen G, Flørenes VA, Bishop DT, Herlyn M, Newton-Bishop J, Slipicevic A (2019)
MX 2 is a novel regulator of cell cycle in melanoma cells
Pigment Cell Melanoma Res (Epub ahead of print)

Kanduri C, Bock C, Gundersen S, **Hovig E**, Sandve GK (2019)
Colocalization analyses of genomic elements: approaches, recommendations and challenges
Bioinformatics, 35 (9), 1615-1624

Khezri R, Rusten TE (2019)
Autophagy and Tumorigenesis in Drosophila
Adv Exp Med Biol, 1167, 113-127

Kotsopoulos J, Lubinski J, Lynch HT, Tung N, Armel S, Senter L, Singer CF, Fruscio R, Couch F, Weitzel JN, Karlan B, Foulkes WD, **Moller P**, Eisen A, Ainsworth P, Neuhausen SL, Olopade O, Sun P, Gronwald J, Narod SA, Hereditary Breast Cancer Clinical Study Group (2019)
Oophorectomy and risk of contralateral breast cancer among BRCA1 and BRCA2 mutation carriers
Breast Cancer Res Treat, 175 (2), 443-449

Kraby MR, Opdahl S, **Russnes HG**, Bofin AM (2019)
Microvessel density in breast cancer: the impact of field area on prognostic informativeness
J Clin Pathol, 72 (4), 304-310

Kryeziu K, Bruun J, Guren TK, **Sveen A, Lothe RA** (2019)
Combination therapies with HSP90 inhibitors against colorectal cancer
Biochim Biophys Acta Rev Cancer, 1871 (2), 240-247

Kumar S, Gu Y, Abudu YP, Bruun JA, **Jain A**, Farzam F, Mudd M, Anonsen JH, **Rusten TE**, Kasof G, Ktistakis N, Lidke KA, Johansen T, Deretic V (2019)
Phosphorylation of Syntaxin 17 by TBK1 Controls Autophagy Initiation
Dev Cell, 49 (1), 130-144.e6

Kyte JA, Fåne A, Pule M, **Gaudernack G** (2019)
Transient redirection of T cells for adoptive cell therapy with telomer-

ase-specific T helper cell receptors isolated from long term survivors after cancer vaccination
Oncoimmunology, 8 (4), e1565236

Köksal H, Dillard P, **Josefsson SE**, Maggadottir SM, Pollmann S, Fåne A, **Blaker YN**, Beiske K, **Huse K**, Kolstad A, Holte H, Kvalheim G, **Smeland EB, Myklebust JH**, Inderberg EM, Wälchli S (2019)
Preclinical development of CD-37CAR T-cell therapy for treatment of B-cell lymphoma
Blood Adv, 3 (8), 1230-1243

Lai X, Geier OM, **Fleischer T**, Garred Ø, Borgen E, Funke SW, **Kumar S**, Rognes ME, Seierstad T, **Børresen-Dale AL, Kristensen VN, Engebraaten O**, Köhn-Luque A, Frigessi A (2019)
Toward Personalized Computer Simulation of Breast Cancer Treatment: A Multiscale Pharmacokinetic and Pharmacodynamic Model Informed by Multitype Patient Data
Cancer Res, 79 (16), 4293-4304

Lampignano R, Neumann MHD, Weber S, Klöten V, Herdean A, Voss T, Groelz D, Babayan A, Tibbesma M, Schlumpberger M, Chemi F, Rothwell DG, Wikman H, Galizzi JP, **Berghelm IR, Russnes H**, Mussolin B, Bonin S, Voigt C, Musa H, Pinzani P, Lianidou E, Brady G, Speicher MR, Pantel K et al. (2019)
Multicenter Evaluation of Circulating Cell-Free DNA Extraction and Downstream Analyses for the Development of Standardized (Pre)analytical Work Flows
Clin Chem (in press)

Landsverk HB, Sandquist LE, Sri-dhara SC, **Rødland GE**, Sabino JC, de Almeida SF, **Grallert B**, Trinkle-Mulcahy L, **Syljuåsen RG** (2019)
Regulation of ATR activity via the RNA polymerase II associated factors CDC73 and PNUTS-PP1
Nucleic Acids Res, 47 (4), 1797-1813

Larsen LK, **Lind GE**, Guldberg P, Dahl C (2019)
DNA-Methylation-Based Detection of Urological Cancer in Urine: Overview of Biomarkers and Considerations on Biomarker Design, Source of DNA, and Detection Technologies
Int J Mol Sci, 20 (11)

Lee YS, Krishnan A, Oughtred R, Rust J, Chang CS, Ryu J, **Kristensen VN**,

Publications

Dolinski K, Theesfeld CL, Troyanskaya OG (2019)

A Computational Framework for Genome-wide Characterization of the Human Disease Landscape Cell Syst, 8 (2), 152-162.e6

Lie-Jensen A, Ivanauskiene K, Malerød L, Jain A, Tan KW, Laerdahl JK, Liestøl K, Stenmark H, Haglund K (2019) **Centralspindlin Recruits ALIX to the Midbody during Cytokinetic Abcission in Drosophila via a Mechanism Analogous to Virus Budding** Curr Biol, 29 (20), 3538-3548.e7

Lien VT, Kristiansen MK, Pettersen S, **Haugen MH**, Olberg DE, Waaler J, Klaveness J (2019) **Towards dual inhibitors of the MET kinase and WNT signaling pathway; design, synthesis and biological evaluation** RSC Adv., 9 (63), 37092-37100

Lien VT, **Pettersen S, Haugen MH**, Olberg DE, **Maelandsmo GM**, Klaveness J (2019) **Design, synthesis and biological evaluation of 6-substituted quinolines derived from cabozantinib as c-Met inhibitors** Arch Pharm (Weinheim), 352 (9), e1900101

Lilleborge M, Falk RS, **Russnes H**, Sauer T, Ursin G, Hofvind S (2019) **Risk of breast cancer by prior screening results among women participating in BreastScreen Norway** Cancer, 125 (19), 3330-3337

Lindholm EM, Leivonen SK, **Undlien E, Nebdal D**, Git A, Caldas C, **Børresen-Dale AL, Kleivi K** (2019) **miR-342-5p as a Potential Regulator of HER2 Breast Cancer Cell Growth** Microna, 8 (2), 155-165

Lindholm EM, Ragle Aure M, Haugen MH, Kleivi Sahlberg K, Kristensen VN, Nebdal D, Børresen-Dale AL, Lingjaerde OC, Engebraaten O (2019) **miRNA expression changes during the course of neoadjuvant bevacizumab and chemotherapy treatment in breast cancer** Mol Oncol, 13 (10), 2278-2296

Lopes N, Bergsland CH, Bjørnslett M, Pellinen T, Svindland A, Nesbakken

A, Almeida R, **Lothe RA**, David L, **Bruun J** (2019 – Epub ahead of print) **Digital image analysis of multiplex fluorescence IHC in colorectal cancer recognizes the prognostic value of CDX2 and its negative correlation with SOX2** Lab Invest, 100 (1), 120-134

Luhr M, **Torgersen ML**, Szalai P, Hashim A, **Brech A**, Staerk J, Engedal N (2019) **The kinase PERK and the transcription factor ATF4 play distinct and essential roles in autophagy resulting from tunicamycin-induced ER stress** J Biol Chem, 294 (20), 8197-8217

Lund-Andersen C, Nakken S, Nygård S, Fromm B, Aasheim LB, Davidson B, Julsrud L, **Abrahamsen TW, Kristensen AT**, Dybdahl B, Larsen SG, **Hovig E, Flatmark K** (2019) **Integrative genomic analysis of peritoneal malignant mesothelioma: understanding a case with extraordinary chemotherapy response** Cold Spring Harb Mol Case Stud, 5 (2)

Lund KV, Simonsen TG, Kristensen GB, **Rofstad EK** (2019) **Pharmacokinetic analysis of DCE-MRI data of locally advanced cervical carcinoma with the Brix model** Acta Oncol, 58 (6), 828-837

Lunde NN, Gregersen I, Ueland T, Shetelig C, Holm S, Kong XY, Michelsen AE, Otterdal K, Yndestad A, Broch K, Gullestad L, **Nyman TA**, Bendz B, Eritsland J, Hoffmann P, Skagen K, Gonçalves I, Nilsson J, Grenegård M, Poreba M, Drag M, Seljeflot I, Sporshheim B, Espevik T, Skjelland M et al. (2019) **Legumain is upregulated in acute cardiovascular events and associated with improved outcome - potentially related to anti-inflammatory effects on macrophages** Atherosclerosis (in press)

Løvf M, Zhao S, Axcrona U, **Johannessen B, Bakken AC, Carm KT, Hoff AM, Myklebost O, Meza-Zepeda LA**, Lie AK, Axcrona K, **Lothe RA, Skotheim RI** (2018) **Multifocal Primary Prostate Cancer Exhibits High Degree of Genomic Heterogeneity** Eur Urol, 75 (3), 498-505

Madssen TS, Cao MD, **Pladsen AV, Ottestad L, Sahlberg KK**, Bathen TF,

Giskeødegård GF (2019) **Historical Biobanks in Breast Cancer Metabolomics- Challenges and Opportunities** Metabolites, 9 (11)

Malt EA, Juhasz K, Frengen A, Wangensteen T, Emilsen NM, Hansen B, **Aga-fonov O**, Nilsen HL (2019) **Neuropsychiatric phenotype in relation to gene variants in the hemizygous allele in 3q29 deletion carriers: A case series** Mol Genet Genomic Med, 7 (9), e889

Marcuello M, Vymetalkova V, Neves RPL, Duran-Sanchon S, **Vedeld HM**, Tham E, van Dalum G, Flügen G, Garcia-Barberan V, Fijneman RJ, Castells A, Vodicka P, **Lind GE**, Stoecklein NH, Heitzer E, Gironella M (2019) **Circulating biomarkers for early detection and clinical management of colorectal cancer** Mol Aspects Med, 69, 107-122

Mauseth B, **Camilio KA**, Shi J, Hammarström CL, Rekdal Ø, Sveinbjörnsson B, Line PD (2019) **The Novel Oncolytic Compound LTX-401 Induces Antitumor Immune Responses in Experimental Hepatocellular Carcinoma** Mol Ther Oncolytics, 14, 139-148

Mavaddat N, Michailidou K, Dennis J, Lush M, Fachal L, Lee A, Tyrer JP, Chen TH, Wang Q, Bolla MK, Yang X, Adank MA, Ahearn T, Aittomäki K, Allen J, Andrulis IL, Anton-Culver H, Antonenkova NN, Arndt V, Aronson KJ, Auer PL, Auvinen P, Barndahl M, Beane Freeman LE, **Børresen-Dale AB**, Beckmann MW et al. (2019)

Polygenic Risk Scores for Prediction of Breast Cancer and Breast Cancer Subtypes Am J Hum Genet, 104 (1), 21-34

Mehto S, Jena KK, Nath P, Chauhan S, Kolapalli SP, Das SK, Sahoo PK, **Jain A**, Taylor GA, Chauhan S (2019) **The Crohn’s Disease Risk Factor IRGM Limits NLRP3 Inflammasome Activation by Impeding Its Assembly and by Mediating Its Selective Autophagy** Mol Cell, 73 (3), 429-445.e7

Meltzer S, Bakke KM, Rød KL, Negård A, **Flatmark K**, Solbakken AM, **Kristensen AT**, Fuglestad AJ, Kersten C, Dueland S, Seierstad T, Hole KH, Lyckander LG, Larsen FO, Schou JV, **Patrick**

Brown D, Abrahamsson H, Redalen KR, Ree AH (2019) **Sex-related differences in primary metastatic site in rectal cancer; associated with hemodynamic factors?** Clin Transl Radiat Oncol, 21, 5-10

Meltzer S, Bjørnestrø T, Lyckander LG, **Flatmark K**, Dueland S, Samiappan R, Johansen C, Kalanxhi E, Ree AH, Redalen KR (2019) **Circulating Exosomal miR-141-3p and miR-375 in Metastatic Progression of Rectal Cancer** Transl Oncol, 12 (8), 1038-1044

Menden MP, Wang D, Mason MJ, Szalai B, Bulusu KC, Guan Y, Yu T, Kang J, Jeon M, Wolfinger R, Nguyen T, Zaslavskiy M, AstraZeneca-Sanger Drug Combination DREAM Consortium, Jang IS, Ghazoui Z, Ahsen ME, Vogel R, Neto EC, Norman T, Tang EKY, Garnett MJ, Veroli GYD, Fawell S, **Hovig E**, Stolvitzky G, Guinney J et al. (2019) **Community assessment to advance computational prediction of cancer drug combinations in a pharmacogenomic screen** Nat Commun, 10 (1), 2674

Menon U, Vedsted P, Zalounina Falborg A, Jensen H, Harrison S, Reguilon I, Barisic A, Bergin RJ, Brewster DH, Butler J, **Brustugun OT**, Bucher O, Cairnduff V, Gavin A, Grunfeld E, Harland E, Kalsi J, Knudsen AK, Lambe M, Law RJ, Lin Y, Malmberg M, Turner D, Neal RD, White V et al. (2019) **Time intervals and routes to diagnosis for lung cancer in 10 jurisdictions: cross-sectional study findings from the International Cancer Benchmarking Partnership (ICBP)** BMJ Open, 9 (11), e025895

Mensali N, Dillard P, Hebeisen M, **Lorenz S, Theodossiou T**, Myhre MR, Fåne A, **Gaudernack G**, Kvalheim G, **Myklebust JH**, Inderberg EM, Wälchli S (2019) **NK cells specifically TCR-dressed to kill cancer cells** EBioMedicine, 40, 106-117

Mensali N, Grenov A, Pati NB, Dillard P, Myhre MR, **Gaudernack G**, Kvalheim G, Inderberg EM, Bakke O, Wälchli S (2019) **Antigen-delivery through invariant chain (CD74) boosts CD8 and CD4 T cell immunity** Oncoimmunology, 8 (3), 1558663

Mensali N, Myhre MR, Dillard P, Pollmann S, **Gaudernack G**, Kvalheim G, Wälchli S, Inderberg EM (2019) **Preclinical assessment of transiently TCR redirected T cells for solid tumour immunotherapy** Cancer Immunol Immunother, 68 (8), 1235-1243

Mesquita P, Freire AF, **Lopes N**, Gomes R, Azevedo D, Barros R, Pereira B, Cavadas B, Pópulo H, Boaventura P, David L, Pereira L, Almeida R (2019) **Expression and Clinical Relevance of SOX9 in Gastric Cancer** Dis Markers, 2019, 8267021

Metcalfe K, Eisen A, Senter L, Armel S, Bordeleau L, Meschino WS, Pal T, Lynch HT, Tung NM, Kwong A, Ainsworth P, Karlan B, **Moller P**, Eng C, Weitzel JN, Sun P, Lubinski J, Narod SA, Hereditary Breast Cancer Clinical Study Group (2019) **International trends in the uptake of cancer risk reduction strategies in women with a BRCA1 or BRCA2 mutation** Br J Cancer, 121 (1), 15-21

Mingo J, Luna S, Gaafar A, **Nunes-Xavier CE**, Torices L, Mosteiro L, Ruiz R, Guerra I, Llaena R, Angulo JC, López JI, Pulido R (2019) **Precise definition of PTEN C-terminal epitopes and its implications in clinical oncology** NPJ Precis Oncol, 3, 11

Miranda A, Hamilton PT, Zhang AW, Pattnaik S, Becht E, Mezheyeuski A, **Bruun J**, Micke P, de Reynies A, Nelson BH (2019) **Cancer stemness, intratumoral heterogeneity, and immune response across cancers** Proc Natl Acad Sci U S A, 116 (18), 9020-9029

Moghimi SM, Simberg D, **Skotland T**, Yaghmur A, Hunter AC (2019) **The Interplay Between Blood Proteins, Complement, and Macrophages on Nanomedicine Performance and Responses** J Pharmacol Exp Ther, 370 (3), 581-592

Mohammadzadeh N, Lunde IG, Andenæs K, Strand ME, Aronsen JM, Skrbic B, Marstein HS, Bandlien C, **Nygård S**, Gorham J, Sjaastad I, Chakravarti S, Christensen G, Engbretsen KVT, Tønnessen T (2019) **The extracellular matrix proteogly-**

can lumican improves survival and counteracts cardiac dilatation and failure in mice subjected to pressure overload Sci Rep, 9 (1), 9206

Mukhopadhyay UK, Oturkar CC, Adams C, Wickramasekera N, Bansal S, Medisetty R, Miller A, Swetzig WM, **Silwal-Pandit L, Børresen-Dale AL**, Creighton CJ, Park JH, Konduri SD, Mukhopadhyay A, Caradori A, Omilian A, Bshara W, Kaipparattu BA, Das GM (2019) **TP53 Status as a Determinant of Pro- vs Anti-Tumorigenic Effects of Estrogen Receptor-Beta in Breast Cancer** J Natl Cancer Inst, 111 (11), 1202-1215

Møller P, Dominguez-Valentin M, Rødland EA, Hovig E (2019) **Causes for Frequent Pathogenic BRCA1 Variants Include Low Penetrance in Fertile Ages, Recurrent De-Novo Mutations and Genetic Drift** Cancers (Basel), 11 (2)

Namlos HM, Boye K, Meza-Zepeda LA (2019) **Cell-free DNA in blood as a non-invasive insight into the sarcoma genome** Mol Aspects Med, 100827 (in press)

Napoli E, Cessna JT, Fitzgerald R, Pibida L, Collé R, Laureano-Pérez L, Zimmerman BE, Bergeron DE (2019) **Primary standardization of ²²⁴Ra activity by liquid scintillation counting** Appl Radiat Isot, 155, 108933

Ney, A., Mahamed, I., Garcia-Sampedro, A., **Selbo, P. K.**, Sancho, P, MacRobert, A. J., Pereira, S. P., Acedo, P. (2019) **Combination light-based therapies to treat pancreatic cancer: a proof of concept.** Proc. SPIE, 11070.

Neckmann U, Wolowczyk C, Hall M, Almaas E, Ren J, **Zhao S, Johannessen B, Skotheim RI**, Bjørkøy G, Ten Dijke P, Holien T (2019) **GREM1 is associated with metastasis and predicts poor prognosis in ER-negative breast cancer patients** Cell Commun Signal, 17 (1), 140

Newman AM, **Steen CB**, Liu CL, Gentles AJ, Chaudhuri AA, Scherer F, Khodadoust MS, Esfahani MS, Luca BA, Steiner D, Diehn M, Alizadeh AA. (2019)

Publications

Determining cell type abundance and expression from bulk tissues with digital cytometry.
Nat Biotechnol. 2019 Jul;37(7):773-782

Nguéa P A, Robertson J, Herrera MC, Chymkowitch P, **Enserink JM** (2019) **Desumoylation of RNA polymerase III lies at the core of the Sumo stress response in yeast**
J Biol Chem, 294 (49), 18784-18795

Nik-Zainal S, Davies H, Staaf J, Ramakrishna M, Glodzik D, Zou X, Martincorena I, Alexandrov LB, Martin S, Wedge DC, Van Loo P, Ju YS, Smid M, Brinkman AB, Morganella S, **Aure MR, Lingjærde OC, Langerød A,** Ringnér M, Ahn SM, Boyault S, Brock JE, Broeks A, Butler A, Desmedt C et al. (2019) **Author Correction: Landscape of somatic mutations in 560 breast cancer whole-genome sequences**
Nature, 566 (7742), E1

Nilsen A, Jonsson M, Aarnes EK, Kristensen GB, **Lyng H** (2019) **Reference MicroRNAs for RT-qPCR Assays in Cervical Cancer Patients and Their Application to Studies of HPV16 and Hypoxia Biomarkers**
Transl Oncol, 12 (3), 576-584

Nilssen Y, **Brustugun OT,** Tandberg Eriksen M, Gulbrandsen J, Skaaheim Haug E, Naume B, Møller B (2019) **Decreasing waiting time for treatment before and during implementation of cancer patient pathways in Norway**
Cancer Epidemiol, 61, 59-69

Nissen-Meyer J, **Skotland T,** Østerud B, **Boye E** (2019) **Improving scientific practice in sports-associated drug testing**
FEBS J, 286 (14), 2664-2669

Njølstad PR, Andreassen OA, Brunak S, Børglum AD, Dillner J, Esko T, Franks PW, Freimer N, Groop L, Heimer H, Hougaard DM, **Hovig E,** Hveem K, Jalanko A, Kaprio J, Knudsen GP, Melbye M, Metspalu A, Mortensen PB, Palmgren J, Palotie A, Reed W, Stefánsson H, Stitzel NO, Sullivan PF et al. (2019) **Roadmap for a precision-medicine initiative in the Nordic region**
Nat Genet, 51 (6), 924-930

Nome ME, Euceda LR, Jabeen S, Debik J, Bathen TF, Giskeødegård GF, **Taskén KA, Maelandsmo GM,** Halvorsen

B, Yndestad A, Borgen E, Garred Ø, Aukrust P, Ueland T, **Engebraaten O, Kristensen VN, Tekpli X** (2019) **Serum levels of inflammation-related markers and metabolites predict response to neoadjuvant chemotherapy with and without bevacizumab in breast cancers**
Int J Cancer, 146 (1), 223-235

Norum JH, Frings O, Kasper M, **Bergholtz H, Zell Thime H,** Bergström Å, Andersson A, Kuiper R, Fredlund E, **Sørlie T,** Tøftgård R (2019) **GLI1-induced mammary gland tumours are transplantable and maintain major molecular features**
Int J Cancer, 146 (4), 1125-1138

Nunes-Xavier CE, Angulo JC, Pulido R, López JI (2019) **A Critical Insight into the Clinical Translation of PD-1/PD-L1 Blockade Therapy in Clear Cell Renal Cell Carcinoma**
Curr Urol Rep, 20 (1), 1

Nunes-Xavier CE, Aurtenetxe O, Zaldumbide L, López-Almaraz R, Erramuzpe A, Cortés JM, López JI, Pulido R (2019) **Protein tyrosine phosphatase PTPN1 modulates cell growth and associates with poor outcome in human neuroblastoma**
Diagn Pathol, 14 (1), 134

Nunes-Xavier CE, Zaldumbide L, Aurtenetxe O, López-Almaraz R, López JI, Pulido R (2019) **Dual-Specificity Phosphatases in Neuroblastoma Cell Growth and Differentiation**
Int J Mol Sci, 20 (5)

Nyakas M, Aamdal E, Jacobsen KD, Guren TK, Aamdal S, Hagene KT, Brunsvig P, Yndestad A, Halvorsen B, **Tasken KA,** Aukrust P, **Maelandsmo GM,** Ueland T (2019) **Prognostic biomarkers for immunotherapy with ipilimumab in metastatic melanoma**
Clin Exp Immunol, 197 (1), 74-82

Nygård S, Lingjærde OC, Caldas C, **Hovig E, Børresen-Dale AL, Helland Å, Haakensen VD** (2019) **PathTracer: High-sensitivity detection of differential pathway activity in tumours**
Sci Rep, 9 (1), 16332

Olsen CE, Cheung LH, **Weyergang A,**

Berg K, Vallera DA, Rosenblum MG, **Selbo PK** (2019) **Design, Characterization, and Evaluation of scFvCD133/rGelonin: A CD133-Targeting Recombinant Immunotoxin for Use in Combination with Photochemical Internalization**
J Clin Med, 9 (1)

Olsvik HL, Svenning S, Abudu YP, **Brech A, Stenmark H,** Johansen T, Mejlvang J (2019) **Endosomal microautophagy is an integrated part of the autophagic response to amino acid starvation**
Autophagy, 15 (1), 182-183

Olweus J, Lund-Johansen F (2019) **Finding Neo (antigens, that is)**
Blood, 134 (2), 108-109

Overå KS, Garcia-Garcia J, Bhujabal Z, **Jain A,** Øvervatn A, Larsen KB, Deretic V, Johansen T, Lamark T, Sjøttem E (2019) **TRIM32, but not its muscular dystrophy-associated mutant, positively regulates and is targeted to autophagic degradation by p62/SQSTM1**
J Cell Sci, 132 (23)

Palomo-Guerrero M, Fadó R, Casas M, Pérez-Montero M, Baena M, Helmer PO, Domínguez JL, Roig A, Serra D, Hayen H, **Stenmark H, Raiborg C,** Casals N (2019) **Sensing of nutrients by CPT1C regulates late endosome/lysosome anterograde transport and axon growth**
Elife, 8

Panagopoulos I, Brunetti M, Stoltenberg M, Strandabø RAU, Staurseth J, Andersen K, Kostolomov I, Hveem TS, **Lorenz S,** Nystad TA, Flægstad T, Micci F, Heim S (2019) **Novel GTF2I-PDGFRB and IKZF1-TYW1 fusions in pediatric leukemia with normal karyotype**
Exp Hematol Oncol, 8, 12

Pandya AD, Jäger E, **Bagheri Fam S,** Höcherl A, Jäger A, Sincari V, Nyström B, Štěpánek P, **Skotland T, Sandvig K,** Hrubý M, **Mælandsmo GM** (2019) **Paclitaxel-loaded biodegradable ROS-sensitive nanoparticles for cancer therapy**
Int J Nanomedicine, 14, 6269-6285

Parashar D, Geethadevi A, **Aure MR,** Mishra J, George J, Chen C, Mishra MK, Tahiri A, Zhao W, Nair B, Lu Y, Mangala

LS, Rodriguez-Aguayo C, Lopez-Berestein G, Camara AKS, Liang M, Rader JS, Ramchandran R, You M, Sood AK, **Kristensen VN,** Mills GB, Pradeep S, Chaluvally-Raghavan P (2019) **miRNA551b-3p Activates an Oncostatin Signaling Module for the Progression of Triple-Negative Breast Cancer**
Cell Rep, 29 (13), 4389-4406.e10

Pashov A, Shivarov V, Hadzhieva M, Kostov V, Ferdinandov D, **Heintz KM,** Pashova S, Todorova M, Vassilev T, Kieber-Emmons T, **Meza-Zepeda LA, Hovig E** (2019) **Diagnostic Profiling of the Human Public IgM Repertoire With Scalable Mimotope Libraries**
Front Immunol, 10, 2796

Pfefferle A, **Jacobs B, Netskar H, Ask EH, Lorenz S, Clancy T, Goodridge JP,** Sohlberg E, **Malmberg KJ** (2019) **Intra-lineage Plasticity and Functional Reprogramming Maintain Natural Killer Cell Repertoire Diversity**
Cell Rep, 29 (8), 2284-2294.e4

Pichard A, Marcatili S, Karam J, Constanzo J, Ladjohounlou R, Courteau A, Jarlier M, Bonnefoy N, **Patzke S,** Stenberg V, Coopman P, Cartron G, Navarro-Teulon I, Repetto-Llamazares A, Heyerdahl H, Dahle J, Bardiès M, Pouget JP (2019) **The therapeutic effectiveness of ¹⁷⁷Lu-lilotomab in B-cell non-Hodgkin lymphoma involves modulation of G2/M cell cycle arrest**
Leukemia (in press)

Pielke R, **Boye E** (2019) **Scientific integrity and anti-doping regulation**
Int. J. Sport Policy Polit., 11 (2), 295-313

Pielke R, Tucker R, **Boye E** (2019) **Scientific integrity and the IAAF testosterone regulations**
Int. Sports Law J., 19 (1-2), 18-26

Pitman KE, Alluri SR, **Kristian A, Aarnes EK, Lyng H,** Riss PJ, Malinen E (2019) **Influx rate of ¹⁸F-fluoroaminosuberic acid reflects cystine/glutamate antiporter expression in tumour xenografts**
Eur J Nucl Med Mol Imaging, 46 (10), 2190-2198

Pitman KE, Bakke KM, **Kristian A,** Malinen E (2019)

Ultra-early changes in vascular parameters from dynamic contrast enhanced MRI of breast cancer xenografts following systemic therapy with doxorubicin and liver X receptor agonist
Cancer Imaging, 19 (1), 88

Prikrylova T, Robertson J, Ferrucci F, Konorska D, Aanes H, Manaf A, **Zhang B,** Vågbø CB, Kuśnierczyk A, Gilljam KM, Løvkvam-Køster C, Otterlei M, Dahl JA, **Enserink J,** Klungland A, Robertson AB (2019) **5-hydroxymethylcytosine Marks Mammalian Origins Acting as a Barrier to Replication**
Sci Rep, 9 (1), 11065

Pulido R, Mingo J, Gaafar A, **Nunes-Xavier CE,** Luna S, Torices L, Angulo JC, López JI (2019) **Precise Immunodetection of PTEN Protein in Human Neoplasia**
Cold Spring Harb Perspect Med, 9 (12)

Pällmann N, Livgård M, Tesikova M, Zeynep Nenseth H, Akkus E, Sikkeland J, **Jin Y,** Koc D, Kuzu OF, Pradhan M, Danielsen HE, Kahraman N, Mokhlis HM, Ozpolat B, Banerjee PP, Uren A, Fazli L, Rennie PS, Jin Y, Saatcioglu F (2019) **Regulation of the unfolded protein response through ATF4 and FAM129A in prostate cancer**
Oncogene, 38 (35), 6301-6318

Pölönen P, Jawahar Deen A, Leinonen HM, Jyrkkänen HK, Kuosmanen S, Mononen M, **Jain A,** Tuomainen T, Pasonen-Seppänen S, Hartikainen JM, Mannermaa A, Nykter M, Tavi P, Johansen T, Heinäniemi M, Levonen AL (2019) **Nrf2 and SQSTM1/p62 jointly contribute to mesenchymal transition and invasion in glioblastoma**
Oncogene, 38 (50), 7473-7490

Ree AH, **Nygaard V, Russnes HG,** Heinrich D, **Nygaard V,** Johansen C, **Bergheim IR, Hovig E,** Beiske K, Negård A, **Børresen-Dale AL, Flatmark K, Mælandsmo GM** (2019) **Responsiveness to PD-1 Blockade in End-Stage Colon Cancer with Gene Locus 9p24.1 Copy-Number Gain**
Cancer Immunol Res, 7 (5), 701-706

Richardsen E, Andersen S, Al-Saad S, Rakaee M, Nordby Y, Pedersen MI, Ness N, Ingebriktsen LM, Fassina A, **Taskén KA,** Mills IG, Donnem T,

Bremnes RM, Busund LT (2019) **Low Expression of miR-424-3p is Highly Correlated with Clinical Failure in Prostate Cancer**
Sci Rep, 9 (1), 10662

Ruoff P, **Agafonov O,** Tveit DM, Thorsen K, Drengstig T (2019) **Homeostatic controllers compensating for growth and perturbations**
PLoS One, 14 (8), e0207831

Russell AE, Sneider A, Witwer KW, Bergese P, Bhattacharyya SN, Cocks A, Cocucci E, Erdbrügger U, Falcon-Perez JM, Freeman DW, Gallagher TM, Hu S, Huang Y, Jay SM, Kano SI, Lavieu G, Leszczynska A, **Llorente AM,** Lu Q, Mahairaki V, Muth DC, Noren Hooten N, Ostrowski M, Prada I, Sahoo S et al. (2019) **Biological membranes in EV biogenesis, stability, uptake, and cargo transfer: an ISEV position paper arising from the ISEV membranes and EVs workshop**
J Extracell Vesicles, 8 (1), 1684862

Rødland GE, Melhus K, Generalov R, **Gilani S,** Bertoni F, Dahle J, **Syljuåsen RG, Patzke S** (2019) **The Dual Cell Cycle Kinase Inhibitor JNJ-7706621 Reverses Resistance to CD37-Targeted Radioimmunotherapy in Activated B Cell Like Diffuse Large B Cell Lymphoma Cell Lines**
Front Oncol, 9, 1301

Røsok BI, **Høst-Brunsell T,** Brudvik KW, Carling U, Dorenberg E, Björnsson B, **Lothe RA,** Bjørnbeth BA, Sandström P (2019) **Characterization of early recurrences following liver resection by ALPPS and two stage hepatectomy in patients with colorectal liver metastases and small future liver remnants; a translational substudy of the LIGRO-RCT**
HPB (Oxford), 21 (8), 1017-1023

Saeednejad Zanjani L, Madjd Z, Rasti A, Asgari M, Abolhasani M, Tam KJ, Roudi R, **Mælandsmo GM, Fodstad Ø, Andersson Y** (2019) **Spheroid-Derived Cells From Renal Adenocarcinoma Have Low Telomerase Activity and High Stem-Like and Invasive Characteristics**
Front Oncol, 9, 1302

Salvatore S, Dagestad Rand K, Grytten I, Ferkingstad E, Domanska D, Holden L, Gheorghe M, **Mathelier A,** Glad I,

Publications

Kjetil Sandve G (2019) **Beware the Jaccard: the choice of similarity measure is important and non-trivial in genomic colocalisation analysis** Brief Bioinform (in press)

Sandhu V, Manem VSK, Mer AS, **Kure EH**, Haibe-Kains B, (2019) **Applications of Computational Systems Biology in Cancer Signaling Pathways** Springer, Singapore, pp 513-537

Sandhu V, Labori KJ, Borgida A, Lungu I, Bartlett J, Hafezi-Bakhtiari S, Denroche RE, Jang GH, Pasternack D, Mbaabali F, Watson M, Wilson J, **Kure EH**, Gallinger S, Haibe-Kains B (2019) **Meta-Analysis of 1,200 Transcriptional Profiles Identifies a Prognostic Model for Pancreatic Ductal Adenocarcinoma** JCO Clin Cancer Inform, 3, 1-16

Scorrano L, De Matteis MA, Emr S, Giordano F, Hajnóczky G, Kornmann B, Lackner LL, Levine TP, Pellegrini L, Reinisch K, Rizzuto R, Simmen T, **Stenmark H**, Ungermann C, Schuldiner M, (2019) **Coming together to define membrane contact sites** Nature Communications 10, 1287

Selbo, P. K., Janetzki, S., Welters, M. J. P., **Håkerud, M., Nedberg, A. G., Edwards, V. T.**, Olivecrona, H., van der Burg, S. H., Otterhaug, T., Hogset, A. (2019) **109P Phase I clinical study for validation of fimaporfin-based photochemical internalisation: A novel technology for enhancing cellular immune responses important for therapeutic effect of peptide-and protein-based vaccines** Ann Oncol, 30, Suppl. 11: xi40-xi41

Seppälä TT, Ahadova A, **Dominguez-Valentin M**, Macrae F, Evans DG, Therkildsen C, Sampson J, Scott R, Burn J, Möslin G, Bernstein I, Holinski-Feder E, Pylvänäinen K, Renkonen-Sinisalo L, Lepistö A, Lautrup CK, Lindblom A, Plazzer JP, Winship I, Tjandra D, Katz LH, Aretz S, Hüneburg R, Holzapfel S, Heinimann K et al. (2019) **Lack of association between screening interval and cancer stage in Lynch syndrome may be accounted for by over-diagnosis; a prospective Lynch syndrome database report** Hered Cancer Clin Pract, 17, 8

Shao B, **Bjaanæs MM, Helland Å**, Schütte C, Conrad T (2019) **EMT network-based feature selection improves prognosis prediction in lung adenocarcinoma** PLoS One, 14 (1), e0204186

Shin D, Nguyen L, T Le M, Ju D, N Le J, **Berg K**, Hirschberg H (2019) **The effects of low irradiance long duration photochemical internalization on glioma spheroids** Photodiagnosis Photodyn Ther, 26, 442-447

Shoemaker CJ, Huang TQ, Weir NR, Polyakov NJ, **Schultz SW**, Denic V (2019) **CRISPR screening using an expanded toolkit of autophagy reporters identifies TMEM41B as a novel autophagy factor** PLoS Biol, 17 (4), e2007044

Shu X, Wu L, Khankari NK, Shu XO, Wang TJ, Michailidou K, Bolla MK, Wang Q, Dennis J, Milne RL, Schmidt MK, Pharoah PDP, Andrulis IL, Hunter DJ, Simard J, Easton DF, Zheng W, Breast Cancer Association Consortium (2019) **Associations of obesity and circulating insulin and glucose with breast cancer risk: a Mendelian randomization analysis** Int J Epidemiol, 48 (3), 795-806

Silva A, Bassim S, Sarkozy C, Mottok A, Lackraj T, Jurinovic V, **Brodtkorb M, Lingjaerde OC**, Sehn LH, Gascoyne RD, Weigert O, Steidl C, Kridel R (2019) **Convergence of risk prediction models in follicular lymphoma** Haematologica, 104 (6), e252-e255

Sioud M (2019) **Releasing the Immune System Brakes Using siRNAs Enhances Cancer Immunotherapy** Cancers (Basel), 11 (2)

Sioud M (2019) **Phage Display Libraries: From Binders to Targeted Drug Delivery and Human Therapeutics** Mol Biotechnol, 61 (4), 286-303

Sioud M, Pettersen S, Ailte I, Fløisand Y (2019) **Targeted Killing of Monocytes/Macrophages and Myeloid Leukemia Cells with Pro-Apoptotic Peptides** Cancers (Basel), 11 (8)

Skjesol A, Yurchenko M, Bösl K, Gravastrand C, Nilsen KE, Grøvdal LM, Agliano F, Patane F, Lentini G, Kim H, Teti G, Kumar Sharma A, Kandasamy RK, Sporsheim B, Starheim KK, Golenbock DT, **Stenmark H**, McCaffrey M, Espevik T, Husebye H (2019) **The TLR4 adaptor TRAM controls the phagocytosis of Gram-negative bacteria by interacting with the Rab11-family interacting protein 2** PLoS Pathog, 15 (3), e1007684

Skotland T, Sandvig K (2019) **The role of PS 18:0/18:1 in membrane function** Nat Commun, 10 (1), 2752

Skånland SS, Cie Iar-Pobuda A (2019) **Off-label uses of drugs for depression** Eur J Pharmacol, 865, 172732

Skånland SS, Cremaschi A, Bendiksen H, Hermansen JU, **Thimiri Govinda Raj DB**, Munthe LA, Tjønnfjord GE, **Taskén K** (2019) **An in vitro assay for biomarker discovery and dose prediction applied to ibrutinib plus venetoclax treatment of CLL** Leukemia, 34 (2), 478-487

Skånland SS, Taskén K (2019) **Carboxyl-Terminal Src Kinase Binds CD28 upon Activation and Mutes Downstream Signaling** J Immunol, 203 (4), 1055-1063

Smeby J, Sveen A, Bergsland CH, Eilertsen IA, Danielsen SA, Eide PW, Hektoen M, Guren MG, Nesbakken A, **Bruun J, Lothe RA** (2019) **Exploratory analyses of consensus molecular subtype-dependent associations of TP53 mutations with immunomodulation and prognosis in colorectal cancer** ESMO Open, 4 (3), e000523

Smeby J, Sveen A, Eilertsen IA, Danielsen SA, Hoff AM, Eide PW, Johannessen B, Hektoen M, Skotheim RI, Guren MG, Nesbakken A, **Lothe RA** (2019) **Transcriptional and functional consequences of TP53 splice mutations in colorectal cancer** Oncogenesis, 8 (6), 35

Smid M, Wilting SM, Uhr K, Rodríguez-González FG, de Weerd V, Prager-Van der Smissen WJC, van der Vlugt-

Daane M, van Galen A, **Børresen-Dale AL**, Nik-Zainal S, Butler A, Martin S, Davies HR, Staaf J, van de Vijver MJ, Richardson AL, MacGrogan G, Salgado R, van den Eynden GGGM, Purdie CA, Thompson AM, Caldas C, Span PN, Sweep FCGJ, Simpson PT, Lakhani SR et al. (2019) **The circular RNome of primary breast cancer** Genome Res, 29 (3), 356-366

Sneeggen M, Pedersen NM, Campsteijn C, **Haugsten EM, Stenmark H, Schink KO** (2019) **WDFY2 restrains matrix metalloproteinase secretion and cell invasion by controlling VAMP3-dependent recycling** Nat Commun, 10 (1), 2850

Sneeggen M, Schink KO, Stenmark H (2019) **Tumor suppression by control of matrix metalloproteinase recycling** Mol Cell Oncol, 6 (6), e1646606

Sowa-Rogozińska N, Sominka H, Nowakowska-Gołacka J, **Sandvig K**, Słomińska-Wojewódzka M (2019) **Intracellular Transport and Cytotoxicity of the Protein Toxin Ricin** Toxins (Basel), 11 (6)

Solberg S, Nilssen Y, **Brustugun OT**, Grimsrud TK, Haram PM, Helbekkmo N, **Helland Å**, Hjelde HH, Jakobsen B, Møller B, Petersen M, Strand TE, Wahl SGF, Aanerud M, Fjellbirkeland L (2019) **Increase in curative treatment and survival of lung cancer in Norway 2001-2016** Eur J Epidemiol, 34 (10), 951-955

Stankovic B, Bjørhovde HAK, Skarshaug R, Aamodt H, Frafjord A, Müller E, Hammarström C, Beraki K, Bækkevold ES, Woldbæk PR, **Helland Å, Brustugun OT**, Øynebråten I, Corthay A (2019) **Immune Cell Composition in Human Non-small Cell Lung Cancer** Front Immunol, 9, 3101

Steen CB, Leich E, **Myklebust JH**, Lockmer S, **Wise JF**, Wahlin BE, Østenstad B, Liestøl K, Kimby E, Rosenwald A, **Smeland EB**, Holte H, **Lingjærde OC, Brodtkorb M** (2019) **A clinico-molecular predictor identifies follicular lymphoma patients at risk of early transformation after first-line immunotherapy** Haematologica, 104 (10), e460-e464

Stiksrud B, Aass HCD, **Lorvik KB**, Ueland T, Trøseid M, Dyrhol-Riise AM (2019) **Activated dendritic cells and monocytes in HIV immunological nonresponders: HIV-induced interferon-inducible protein-10 correlates with low future CD4+ recovery** AIDS, 33 (7), 1117-1129

Sukonina V, Ma H, Zhang W, Bartesaghi S, Subhash S, Heglind M, **Foyt H**, Betz MJ, Nilsson D, Lidell ME, Naumann J, Haufs-Brusberg S, Palmgren H, Mondal T, Beg M, Jedrychowski MP, **Taskén K**, Pfeifer A, Peng XR, Kanduri C, Enerbäck S (2019) **FOXK1 and FOXK2 regulate aerobic glycolysis** Nature, 566 (7743), 279-283

Sun CH, **Berg K**, Hirschberg H (2019) **Photochemical Internalization Enhanced Nonviral Suicide Gene Therapy** Methods Mol Biol, 1895, 165-176

Sveen A, Cremolini C, Dienstmann R (2019) **Predictive modeling in colorectal cancer: time to move beyond consensus molecular subtypes** Ann Oncol, 30 (11), 1682-1685

Sveen A, Kopetz S, Lothe RA (2019) **Biomarker-guided therapy for colorectal cancer: strength in complexity** Nat Rev Clin Oncol, 17 (1), 11-32

Swanson DM, **Lien T, Bergholtz H, Sørlie T**, Frigessi A (2019) **A Bayesian two-way latent structure model for genomic data integration reveals few pan-genomic cluster subtypes in a breast cancer cohort** Bioinformatics, 35 (23), 4886-4897

Switlyk MD, **Salberg UB**, Geier OM, Vlatkovic L, Lilleby W, **Lyng H**, Seierstad T (2019) **PTEN Expression in Prostate Cancer: Relationship With Clinicopathologic Features and Multiparametric MRI Findings** AJR Am J Roentgenol, 212 (6), 1206-1214

Szwed M, Sønstevoid T, Øverbye A, Engedal N, **Grallert B**, Mørch Y, Sulheim E, **Iversen TG, Skotland T, Sandvig K, Torgersen ML** (2019) **Small variations in nanoparticle structure dictate differential cellular**

stress responses and mode of cell death Nanotoxicology, 13 (6), 761-782

Szybowska P, Kostas M, Wesche J, Wiedlocha A, Haugsten EM (2019) **Cancer Mutations in FGFR2 Prevent a Negative Feedback Loop Mediated by the ERK1/2 Pathway** Cells, 8 (6)

Saetersmoen ML, Hammer Q, Valamehr B, Kaufman DS, **Malmberg KJ** (2019) **Off-the-shelf cell therapy with induced pluripotent stem cell-derived natural killer cells** Semin Immunopathol, 41 (1), 59-68

Sørensen O, Andersen AM, Larsen SG, Giercksky KE, **Flatmark K** (2019) **Intraperitoneal mitomycin C improves survival compared to cytoreductive surgery alone in an experimental model of high-grade pseudomyxoma peritonei** Clin Exp Metastasis, 36 (6), 511-518

Taylor-King JP, Baratchart E, Dhawan A, Coker EA, **Rye IH, Russnes H**, Chapman SJ, Basanta D, Marusyk A (2019) **Simulated ablation for detection of cells impacting paracrine signalling in histology analysis** Math Med Biol, 36 (1), 93-112

Tekpli X, Lien T, Røssevold AH, Nebdal D, Borgen E, Ohnstad HO, **Kyte JA**, Vallon-Christersson J, **Fongaard M, Due EU**, Svartdal LG, Sveli MAT, Garred Ø, OSBREAC, Frigessi A, **Sahlberg KK, Sørlie T, Russnes HG**, Naume B, **Kristensen VN** (2019) **An independent poor-prognosis subtype of breast cancer defined by a distinct tumor immune microenvironment** Nat Commun, 10 (1), 5499

Ten Broeke SW, Rodríguez-Girondo M, Suerink M, Aretz S, Bernstein I, Capellá G, Engel C, Gomez-García EB, van Hest LP, von Knebel Doeberitz M, Lagerstedt-Robinson K, Letteboer TGW, **Moller P**, van Os TA, Pineda M, Rahner N, Olderode-Berends MJW, von Salomé J, Schackert HK, Spruijt L, Steinke-Lange V, Wagner A, Tops CMJ, Nielsen M (2019) **The Apparent Genetic Anticipation in PMS2-Associated Lynch Syndrome Families Is Explained by Birth-cohort Effect** Cancer Epidemiol Biomarkers Prev, 28 (6), 1010-1014

Publications

Theodossiou TA, Ali M, Grigalavicius M, Grallert B, Dillard P, Schink KO, Olsen CE, Wäichli S, Inderberg EM, Kubin A, Peng Q, Berg K (2019) **Simultaneous defeat of MCF7 and MDA-MB-231 resistances by a hypericin PDT-tamoxifen hybrid therapy** NPJ Breast Cancer, 5, 13

Thorsen T, Solheim JM, Labori KJ, Line PD, **Aandahl EM** (2019) **Liver transplantation as a lifesaving procedure for postthepatectomy liver failure and iatrogenic liver injuries** Langenbecks Arch Surg, 404 (3), 301-308

Tolios A, De Las Rivas J, **Hovig E**, Trouillas P, Scorilas A, Mohr T (2019) **Computational approaches in cancer multidrug resistance research: Identification of potential biomarkers, drug targets and drug-target interactions** Drug Resist Updat, 48, 100662

Tomescu-Baciu A, Johansen JN, Holmøy T, Greiff V, **Stensland M, de Souza GA**, Vartdal F, Lossius A (2019) **Persistence of intrathecal oligoclonal B cells and IgG in multiple sclerosis** J Neuroimmunol, 333, 576966

Totland MZ, Rasmussen NL, Knudsen LM, Leithe E (2019) **Regulation of gap junction intercellular communication by connexin ubiquitination: physiological and pathophysiological implications** Cell Mol Life Sci (in press)

Umu SU, Langseth H, Keller A, Meese E, **Helland Å**, Lyle R, Rounge TB (2020) **A 10-year prediagnostic follow-up study shows that serum RNA signals are highly dynamic in lung carcinogenesis** Mol Oncol, 14 (2), 235-247

Vaccaro CA, López-Kostner F, Adriana DV, Palmero EI, Rossi BM, Antelo M, Solano A, Carraro DM, Forones NM, Bohorquez M, Lino-Silva LS, Buleje J, Spirandelli F, Abe-Sandes K, Nascimento I, Sullcahuaman Y, Saroca C, Gonzalez ML, Herrando AI, Alvarez K, Neffa F, Galvão HC, Esperon P, Golubicki M, **Hovig E**, Cisterna D et al. (2019) **From colorectal cancer pattern to the characterization of individuals at risk: Picture for genetic research in Latin America** Int J Cancer, 145 (2), 318-326

Vaclavikova R, Klajic J, Brynychova V, Elsnerova K, **Alnaes GIG**, Tost J, **Kristensen VN**, Rob L, Kodet R, Skapa P, Mrhalova M, Soucek P (2019) **Development of highresolution melting analysis for ABCB1 promoter methylation: Clinical consequences in breast and ovarian carcinoma** Oncol Rep, 42 (2), 763-774

Vietri M, Radulovic M, Stenmark H (2019) **The many functions of ESCRTs** Nat Rev Mol Cell Biol, 21 (1), 25-42

Webb M, Manley K, Olivan M, **Guldvik I**, Palczynska M, Hurst R, Connell SP, Mills IG, Brewer DS, Mills R, Cooper CS, Clark J (2019) **Methodology for the at-home collection of urine samples for prostate cancer detection** Biotechniques (in press)

Wise JF, Lawrence MS. (2019) **Huge whole-genome study of human metastatic cancers.** Nature. Nov;575(7781):60-61

Woldemariam NT, **Agafonov O**, Høyheim B, Houston RD, Taggart JB, Andreasen R (2019) **Expanding the miRNA Repertoire in Atlantic Salmon; Discovery of IsomiRs and miRNAs Highly Expressed in Different Tissues and Developmental Stages** Cells, 8 (1)

Zakrzewska M, Opalinski L, **Haugsten EM**, Otlewski J, **Wiedlocha A** (2019) **Crosstalk between p38 and Erk 1/2 in Downregulation of FGF1-Induced Signaling** Int J Mol Sci, 20 (8)

Zhang R, Lai L, Dong X, He J, You D, Chen C, Lin L, Zhu Y, Huang H, Shen S, Wei L, Chen X, Guo Y, Liu L, Su L, Shafer A, Moran S, **Fleischer T, Bjaanæs MM**, Karlsson A, Planck M, Staaf J, **Helland Å**, Esteller M, Wei Y et al. (2019) **SIPA1L3 methylation modifies the benefit of smoking cessation on lung adenocarcinoma survival: an epigenomic-smoking interaction analysis** Mol Oncol, 13 (5), 1235-1248

Zhang R, Lai L, He J, Chen C, You D, Duan W, Dong X, Zhu Y, Lin L, Shen S, Guo Y, Su L, Shafer A, Moran S, **Fleischer T, Bjaanæs MM**, Karlsson

A, Planck M, Staaf J, **Helland Å**, Esteller M, Wei Y, Chen F, Christiani DC (2019) **EGLN2 DNA methylation and expression interact with HIF1A to affect survival of early-stage NSCLC** Epigenetics, 14 (2), 118-129

Zhen Y, Spangenberg H, Munson MJ, **Brech A, Schink KO, Tan KW, Sørensen V, Wenzel EM, Radulovic M**, Engedal N, Simonsen A, **Raiborg C, Stenmark H** (2019) **ESCRT-mediated phagophore sealing during mitophagy** Autophagy

Zhu Q, **Tekpli X**, Troyanskaya OG, **Kristensen VN** (2019) **Subtype-Specific Transcriptional Regulators in Breast Tumors Subjected to Genetic and Epigenetic Alterations** Bioinformatics (in press)

Öjlert ÅK, Halvorsen AR, Nebdal D, Lund-Iversen M, Solberg S, **Brustugun OT, Lingjaerde OC, Helland Å** (2019) **The immune microenvironment in non-small cell lung cancer is predictive of prognosis after surgery** Mol Oncol, 13 (5), 1166-1179

Aasen T, **Leithe E**, Graham SV, Kameritsch P, Mayán MD, Mesnil M, Pogoda K, Tabernero A (2019) **Connexins in cancer: bridging the gap to the clinic** Oncogene, 38 (23), 4429-4451

Publications 2020 and in press

Bai B, Myklebust JH, Wäichli S (2020) **Gene Editing in B-Lymphoma Cell Lines Using CRISPR/Cas9 Technology** Methods Mol Biol, 2115, 445-454

Casey NP, Kyte JA, Fujiwara H (2020) **Use of RNA Interference with TCR Transfer to Enhance Safety and Efficiency** Methods Mol Biol, 2115, 327-349

Chu DT, Phuong TNT, Tien NLB, Tran DK, Thanh VV, Quang TL, Truong DT, Pham VH, Ngoc VTN, Chu-Dinh T, **Kushekhar K** (2020) **An Update on the Progress of Isola-**

tion, Culture, Storage, and Clinical Application of Human Bone Marrow Mesenchymal Stem/Stromal Cells Int J Mol Sci, 21 (3)

Dal NK, Kocere A, Wohlmann J, Van Herck S, Bauer TA, Resseguier J, **Bagherifam S**, Hyldmo H, Barz M, De Geest BG, Fenaroli F (2020) **Zebrafish Embryos Allow Prediction of Nanoparticle Circulation Times in Mice and Facilitate Quantification of Nanoparticle-Cell Interactions** Small, 16 (5), e1906719

Darvekar S, **Juzenas P, Oksvold M**, Kleinauskas A, Holien T, Christensen E, **Stokke T, Sioud M**, Peng Q (2020) **Selective Killing of Activated T Cells by 5-Aminolevulinic Acid Mediated Photodynamic Effect: Potential Improvement of Extracorporeal Photopheresis** Cancers 2020, 12(2), 377

Escala-Garcia M, Abraham J, Andrulis IL, Anton-Culver H, Arndt V, Ashworth A, Auer PL, Auvinen P, Beckmann MW, Beesley J, Behrens S, Benitez J, Bermisheva M, Blomqvist C, Blot W, Bogdanova NV, Bojesen SE, Bolla MK, **Børresen-Dale AL**, Brauch H, Brenner H, Brucker SY, Burwinkel B, Caldas C, Canzian F et al. (2020)

A network analysis to identify mediators of germline-driven differences in breast cancer prognosis Nat Commun, 11 (1), 312

Fachal L, Aschard H, Beesley J, Barnes DR, Allen J, Kar S, Pooley KA, Dennis J, Michailidou K, Turman C, Soucy P, Lemaçon A, Lush M, Tyrer JP, Ghous-saini M, Moradi Marjaneh M, Jiang X, Agata S, Aittomäki K, Alonso MR, Andrulis IL, Anton-Culver H, Antonenkova NN, Arason A, Arndt V et al. (2020) **Fine-mapping of 150 breast cancer risk regions identifies 191 likely target genes** Nat Genet, 52 (1), 56-73

Eilertsen IA, Moosavi SH, Strømme JM, Nesbakken A, **Johannessen B, Lothe RA, Sveen A** (2020) **Technical differences between sequencing and microarray platforms impact transcriptomic subtyping of colorectal cancer** Cancer Lett, 469, 246-255

Flem-Karlsen K, **Nyakas M**, Farstad IN, McFadden E, Wernhoff P, Jacobsen KD, Flørenes VA, **Mælandsmo GM** (2020)

Soluble AXL as a marker of disease progression and survival in melanoma PLoS One, 15 (1), e0227187

Fornes O, Castro-Mondragon JA, Khan A, van der Lee R, Zhang X, Richmond PA, Modi BP, Correard S, Gheorghe M, Baranašić D, Santana-Garcia W, Tan G, Chèneby J, Ballester B, Parcy F, Sandelin A, Lenhard B, Wasserman WW, **Mathelier A** (2020) **JASPAR 2020: update of the open-access database of transcription factor binding profiles** Nucleic Acids Res, 48 (D1), D87-D92

Fromm B, Domanska D, **Høye E**, Ovchinnikov V, Kang W, Aparicio-Puerta E, Johansen M, **Flatmark K, Mathelier A, Hovig E**, Hackenberg M, Friedländer MR, Peterson KJ (2020) **MirGeneDB 2.0: the metazoan microRNA complement** Nucleic Acids Res, 48 (D1), D132-D141

Fromm B, Domanska D, **Høye E**, Ovchinnikov V, Kang W, Aparicio-Puerta E, **Johansen M, Flatmark K, Mathelier A, Hovig E**, Hackenberg M, Friedländer MR, Peterson KJ (2020) **MirGeneDB 2.0: the metazoan microRNA complement** Nucleic Acids Res, 48 (D1), D1172

Georgiesh T, Boye K, Bjerkehagen B (2020) **A novel risk score to predict early and late recurrence in solitary fibrous tumour** Histopathology (in press)

Goleva-Fjellset S, Bjurholt AM, **Kure EH**, Larsen IK, Støren Ø, Sæbø M (2020) **Distribution of allele frequencies for genes associated with physical activity and/or physical capacity in a homogenous Norwegian cohort- a cross-sectional study** BMC Genet, 21 (1), 8

Hovda T, Holen ÅS, Lång K, Albertsen JL, Bjørndal H, Brandal SHB, **Sahlberg KK**, Skaane P, Suhrke P, Hofvind S (2020) **Interval and Consecutive Round Breast Cancer after Digital Breast Tomosynthesis and Synthetic 2D Mammography versus Standard 2D Digital Mammography in Breast-Screen Norway** Radiology, 294 (2), 256-264

Iversen PO, **Sioud M** (2020)

Harnessing the Antiviral-Type Responses Induced by Immunostimulatory siRNAs for Cancer Immunotherapy Methods Mol Biol, 2115, 281-287

Jakobi AJ, Huber ST, Mortensen SA, **Schultz SW**, Palara A, Kuhm T, Shrestha BK, Lamark T, Hagen WJH, Wilmanns M, Johansen T, **Brech A**, Sachse C (2020) **Structural basis of p62/SQSTM1 helical filaments and their role in cellular cargo uptake** Nat Commun, 11 (1), 440

Jakobsen LH, Ellin F, Smeland KB, Wåsterlid T, Christensen JH, Jørgensen JM, Josefsson PL, Øvlisen AK, Holte H, **Blaker YN**, Grauslund JH, Bjørn J, Molin D, Lagerlöf I, Smedby KE, Colvin K, Thanarajasingam G, Maurer MJ, Habermann TM, Song KW, Zhu KY, Gerrie AS, Cheah CY, El-Galaly TC (2020) **Minimal relapse risk and early normalization of survival for patients with Burkitt lymphoma treated with intensive immunochemotherapy: an international study of 264 real-world patients** Br J Haematol (in press)

Jerjes W, Hamdoon Z, **Berg K**, Høgset A, Hopper C (2020) **Apparent Complete Response of a Treatment Refractory and Recurrent Squamous Cell Carcinoma Lesion to Photochemical Internalization: A Clinical Case Study** Photochem Photobiol (in press)

Knutsen E, Lellahi SM, **Aure MR, Nord S**, Fismen S, Larsen KB, Gabriel MT, Hedberg A, **Bjørklund SS**, Oslo Breast Cancer Research Consortium (OSBRE-AC), Bofin AM, **Mælandsmo GM, Sør-lie T**, Mortensen ES, Perander M (2020) **The expression of the long NEAT1_2 isoform is associated with human epidermal growth factor receptor 2-positive breast cancers** Sci Rep, 10 (1), 1277

Lopes N, Bergsland CH, Bjørnslett M, Pellinen T, Svindland A, Nesbakken A, Almeida R, **Lothe RA**, David L, **Brun J** (2020) **Digital image analysis of multiplex fluorescence IHC in colorectal cancer recognizes the prognostic value of CDX2 and its negative correlation with SOX2** Lab Invest, 100 (1), 120-134

Publications

Malenge MM, **Patzke S**, Ree AH, **Stokke T**, Ceuppens P, Dahle J, Repetto-Llamazares AHV. (2020) ¹⁷⁷Lu-lilotomab satetraxetan has the potential to counteract resistance to rituximab in rituximab resistant non-Hodgkin lymphoma. J Nuclear Medicine (in press)

Mensali N, Myhre MR, Dillard P, Pollmann S, **Gaudernack G**, Kvalheim G, Wälchli S, Inderberg EM (2020) **Correction to: Preclinical assessment of transiently TCR redirected T cells for solid tumour immunotherapy** Cancer Immunol Immunother, 69 (1), 159-161

Meås HZ, Haug M, Beckwith MS, Louet C, Ryan L, Hu Z, Landskron J, Nordbø SA, **Taskén K**, Yin H, Damås JK, Flo TH (2020) **Sensing of HIV-1 by TLR8 activates human T cells and reverses latency** Nat Commun, 11 (1), 147

Mobergslie A, Sioud M (2020) **Exploring 5’-Biotinylation of the Sense Strand to Improve siRNA Specificity and Potency** Methods Mol Biol, 2115, 163-170

Nakstad ER, Stær-Jensen H, Wimmer H, Henriksen J, Alteheld LH, Reichenbach A, Drægni T, Šaltytė-Benth J, Wilson JA, Etholm L, **Øijordsbakken M**, Eritsland J, Seljeflot I, Jacobsen D, Andersen GØ, Lundqvist C, Sunde K (2020) **Late awakening, prognostic factors and long-term outcome in out-of-hospital cardiac arrest - results of the prospective Norwegian Cardio-Respiratory Arrest Study (NORCAST)** Resuscitation (in press)

Omsland M, Andresen V, Gullaksen SE, **Ayuda-Durán P**, Popa M, Hovland R, Brendehaug A, **Enserink J**, McCormack E, Gjertsen BT (2020) **Tyrosine kinase inhibitors and interferon-α increase tunneling nanotube (TNT) formation and cell adhesion in chronic myeloid leukemia (CML) cell lines** FASEB J 34(3), 3773-3791

Peng W, de Bruijn HS, Ten Hagen TLM, **Berg K**, Roodenburg JLN, van Dam GM, Witjes MJH, Robinson DJ (2020) **In-Vivo Optical Monitoring of the Efficacy of Epidermal Growth Factor**

Receptor Targeted Photodynamic Therapy: The Effect of Fluence Rate Cancers (Basel), 12 (1)

Rein ID, Notø HØ, Bostad M, Huse K, Stokke T (2020) **Cell Cycle Analysis and Relevance for Single-Cell Gating in Mass Cytometry** Cytometry A

Simonsen TG, Gaustad JV, Rofstad EK (2020) **Bevacizumab treatment of meningeal melanoma metastases** J Transl Med, 18 (1), 13

Sioud M (2020) **Unleashing the Therapeutic Potential of Dendritic and T Cell Therapies Using RNA Interference** Methods Mol Biol, 2115, 259-280

Sioud M (2020) **RNA and CRISPR Interferences: Past, Present, and Future Perspectives** Methods Mol Biol, 2115, 1-22

Sioud M (2020) **Optimized siRNA Delivery into Primary Immune Cells Using Electroporation** Methods Mol Biol, 2115, 119-131

Šošić L, **Selbo PK**, Kotkowska ZK, Kündig TM, Høgset A, Johansen P (2020) **Photochemical Internalization: Light Paves Way for New Cancer Chemotherapies and Vaccines** Cancers (Basel), 12 (1)

Stenberg VY, Juzeniene A, Chen Q, Yang X, Bruland ØS, Larsen RH (2020) **Preparation of the alpha-emitting prostate-specific membrane antigen targeted radioligand [²¹² Pb]Pb-NG001 for prostate cancer** J Labelled Comp Radiopharm

Szwed M, Torgersen ML, Kumari RV, Yadava SK, **Pust S, Iversen TG, Skotland T**, Giri J, **Sandvig K** (2020) **Biological response and cytotoxicity induced by lipid nanocapsules** J Nanobiotechnology, 18 (1), 5

Sæbøe-Larssen S, Sioud M (2020) **Improving Dendritic Cell Cancer Vaccine Potency Using RNA Interference** Methods Mol Biol, 2115, 249-258

Umu SU, Langseth H, Keller A, Meese E, **Helland Å**, Lyle R, Rounge TB (2020) **A 10-year prediagnostic follow-up study shows that serum RNA signals are highly dynamic in lung carcinogenesis** Mol Oncol, 14 (2), 235-247

Wälchli S, **Sioud M** (2020) **Next Generation of Adoptive T Cell Therapy Using CRISPR/Cas9 Technology: Universal or Boosted?** Methods Mol Biol, 2115, 407-417

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