

NCMM

Centre for Molecular Medicine Norway

Nordic EMBL partnership for Molecular Medicine

From disease mechanisms to clinical practice

”

Hvis det ikke var for stor variabilitet mellom pasientene, ville medisin vært en eksakt vitenskap og ikke en kunst.

Sir William Osler
Johns Hopkins School of Medicine
Baltimore 1892

Human genome – what does our genes do?





GENOMICS

Understanding human diversity

David B. Goldstein and Gianpiero L. Cavalleri

Variation : 1 per mille or 6-10 millions small differences
that together constitute our
individual genetic "make-up"

“The normal is the rarest thing in the world ”

- Vi er alle unike

- ~10 millioner SNPer skiller 2 individer fra hverandre
- 30 – 80 SNPer er tilstrekkelig til å identifisere et individ
- SNPer arves i blokker som kalles haplotyper

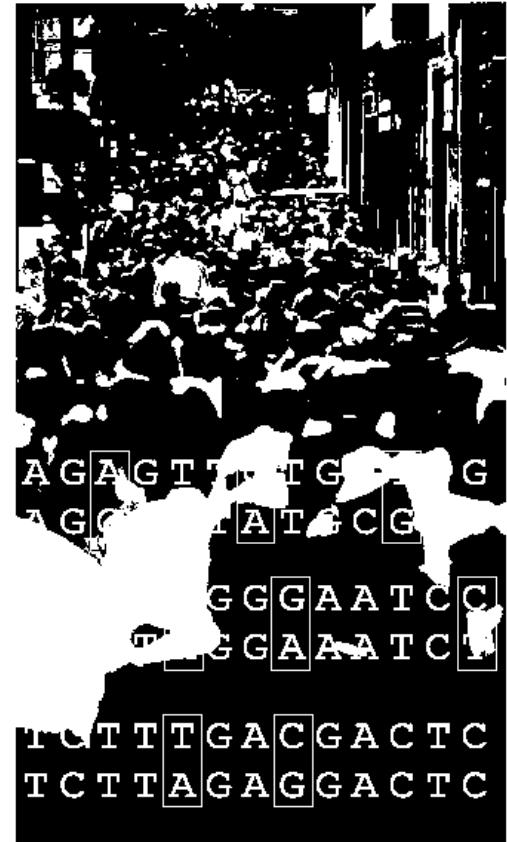
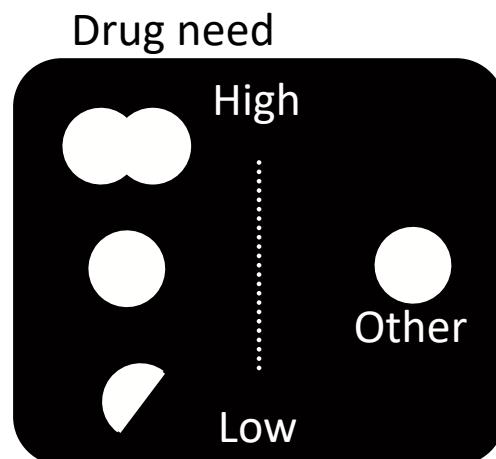
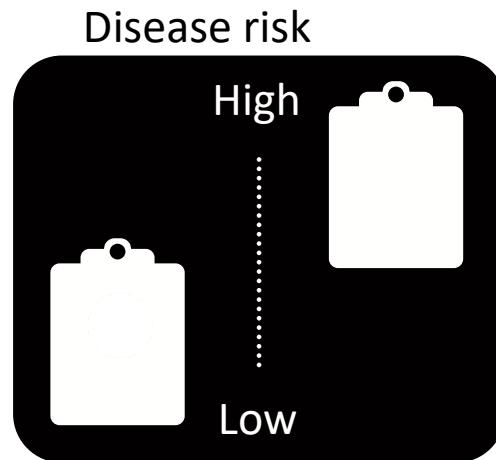
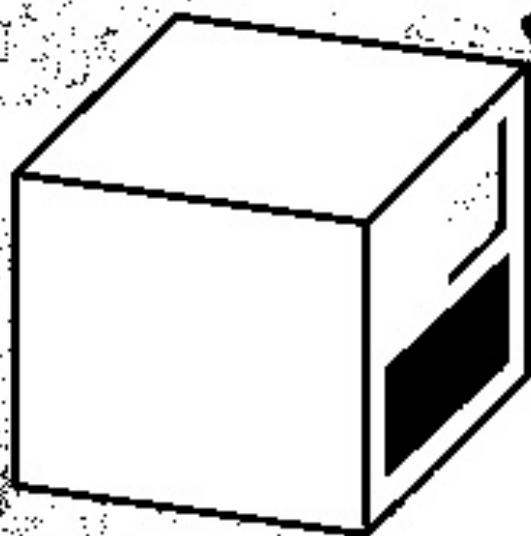


Figure 1 The most common sources of variation between humans are single nucleotide polymorphisms (SNPs) — single base differences between genome sequences. Fragments of two sequences, with eight SNPs, are shown.



GENOMIC SEQUENCING



Reduced!

Evidence-based vs tailored medicine

Access to Precision Medicine

N-of-One works with you and your medical team to create a diagnostic and treatment strategy tailored to YOUR specific tumor type and molecular profile
[Learn More](#)



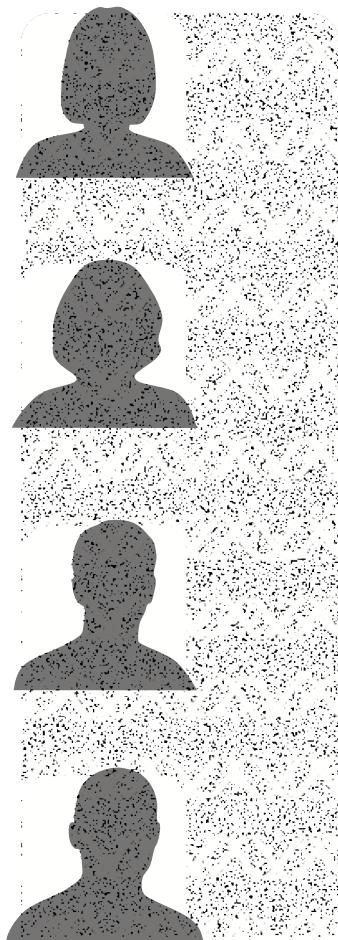
N-of-1 Trials: A New Future?

For reprint orders, please contact: reprints@futuremedicine.com

Eric B. Larson, MD, MPH

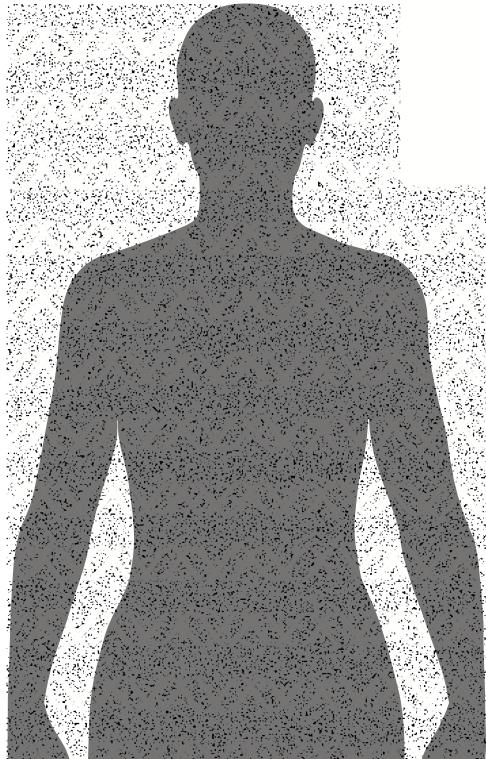
Group Health Research Institute, Seattle, WA, USA.

Personalized medicine: a competitor of evidence-based medicine?



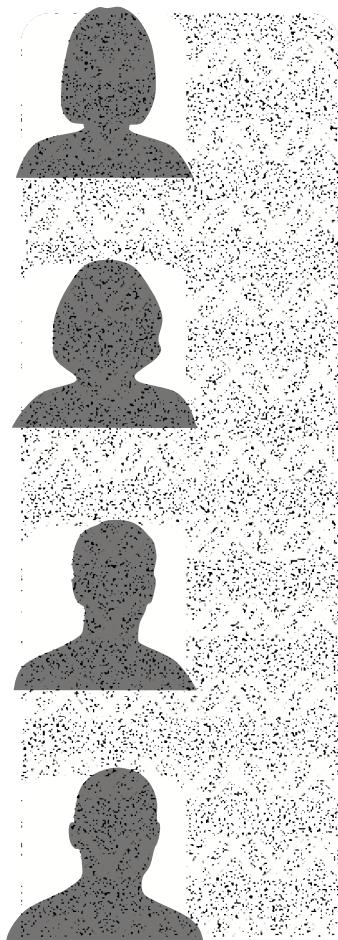
Utfordring med individualisert medisin

- Hver pasient sin sykdom
- En sykdom i hver pasient
- Alle observations er n=1
- Utfordrer sykdomsklassifikasjon og diagnosesystemer



Framtid i molekylærmedisin

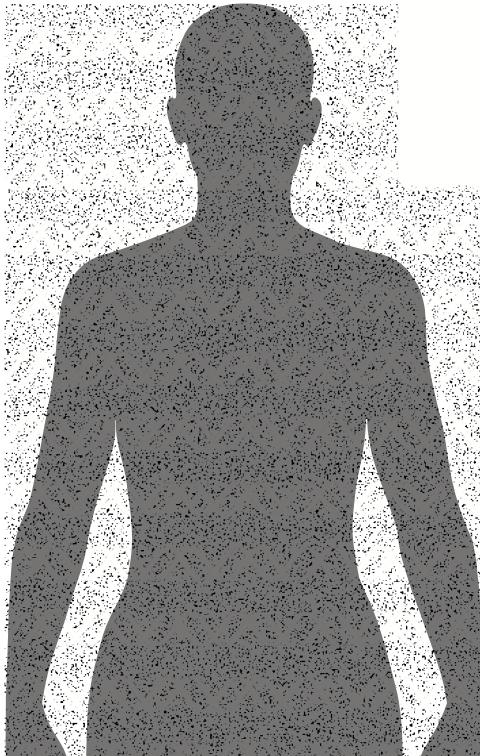
- Kjenne alle sykdomsgener
- Kunne diagnostisere feil i alle disse (dog uten at det nødvendigvis finnes behandling)
- Ha mulighet til å iverksette forebyggende tiltak
- Koble spesifikk diagnostikk med målstyrt og individuell behandling
- DETTE HAR ALLEREDE STARTET I KREFTBEHANDLING



Dyrt og eksklusivt?

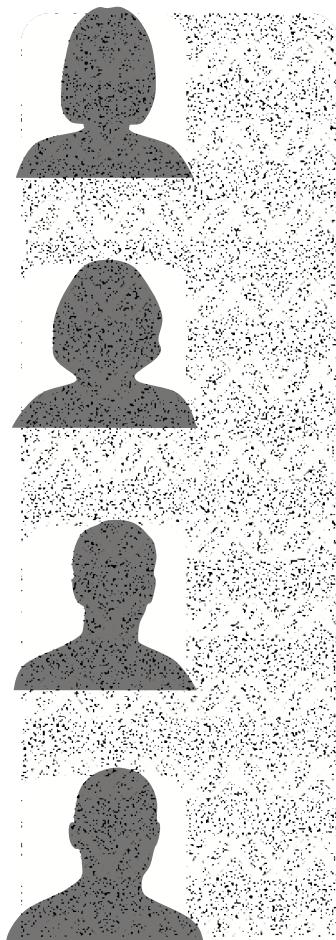
- Internasjonal utvikling
- Forventning
- Kan bli dyrt, eksklusivt og bare tilgjengelig i rike land
- Gradvis og asynkront
- Betalingsvilje og prioriteringsspørsmål
- Presendens

..Eller investering i fremtidig pasient-nytte og bedre helseøkonomi?



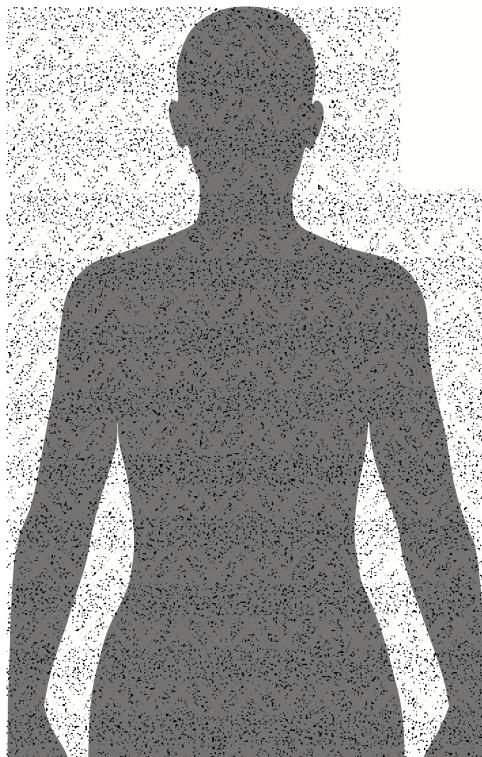
- Behandling rettet mot årsak i hver enkelt pasient
- God behandlingseffekt hos de man behandler og dermed god effekt av ”investeringen”
- Ikke behandle pasienter som man vet ikke vil ha effekt - unngår bivirkninger, falske håp og sparer kostnader
- Kostnadseffektive intervensjoner i livsstil i riskogrupper
- Forskning og utvikling som foregår i dag kan gi fremtidig pasient-nytte og mulig helseøkonomisk gevinst

Investeringer i skreddersydd medisin?



- Internasjonal utvikling – ønske om nasjonale tilbud
- Krever infrastruktur og ekspertise
- Kunnskap om særskilte genetiske risikofaktorer for norsk befolkning
- Prising viktig faktor –Pris per pasient opp når mengden pasienter går ned pga stratifisering, men god effekt på de som behandles. Samtidig ikke ta ut hele innsparingen i økt pris, dvs balanse.
- ”Utgifter” til medisinisk forskning innen dette fagområdet kan sees som investeringskostnader i et helseøkonomisk perspektiv

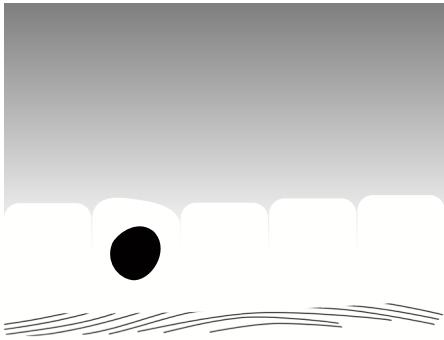
Hva kan gjøres nå?



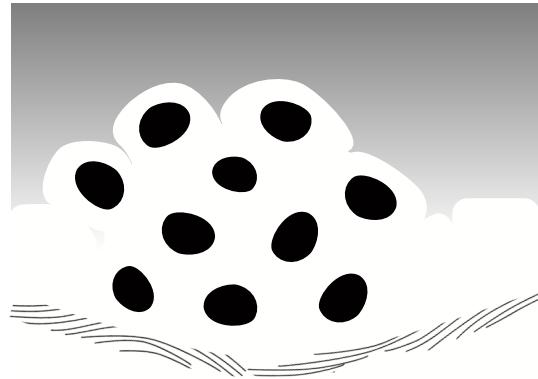
- Industrien:
 - "One-drug-fits-all" forlates
 - Companion diagnostics
 - Stratifisering
- Offentlig sektor:
 - Standarder for individbasert behandling
 - Bekreftende undersøkelser og helseøkonomi
 - Biobankmateriale og helsedata - rask validering av nye tester
 - Kvalifiserte avgjørelser med bakgrunn i klinisk effekt, pasientnytte og god samfunns- og helseøkonomi

Skreddersydd
medisin og individ-
tilpasset
kreftbehandling

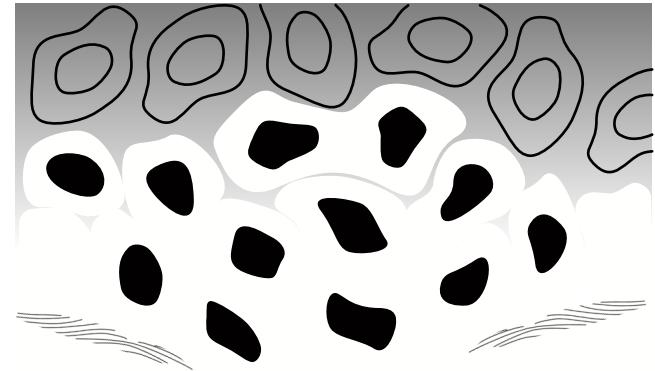




Mutation

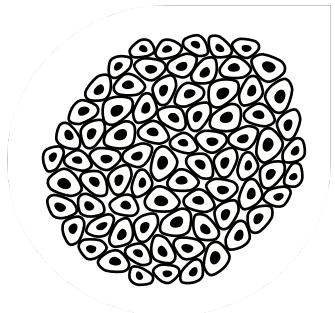


Uncontrolled cell growth



Breach of boundaries

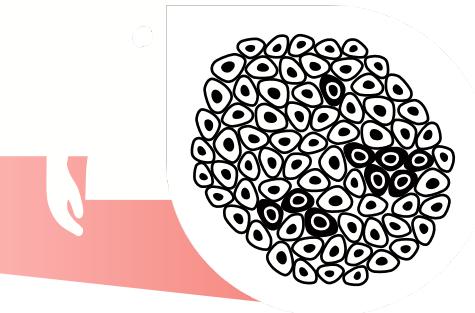




XX

Few mutations

Heterogeneity



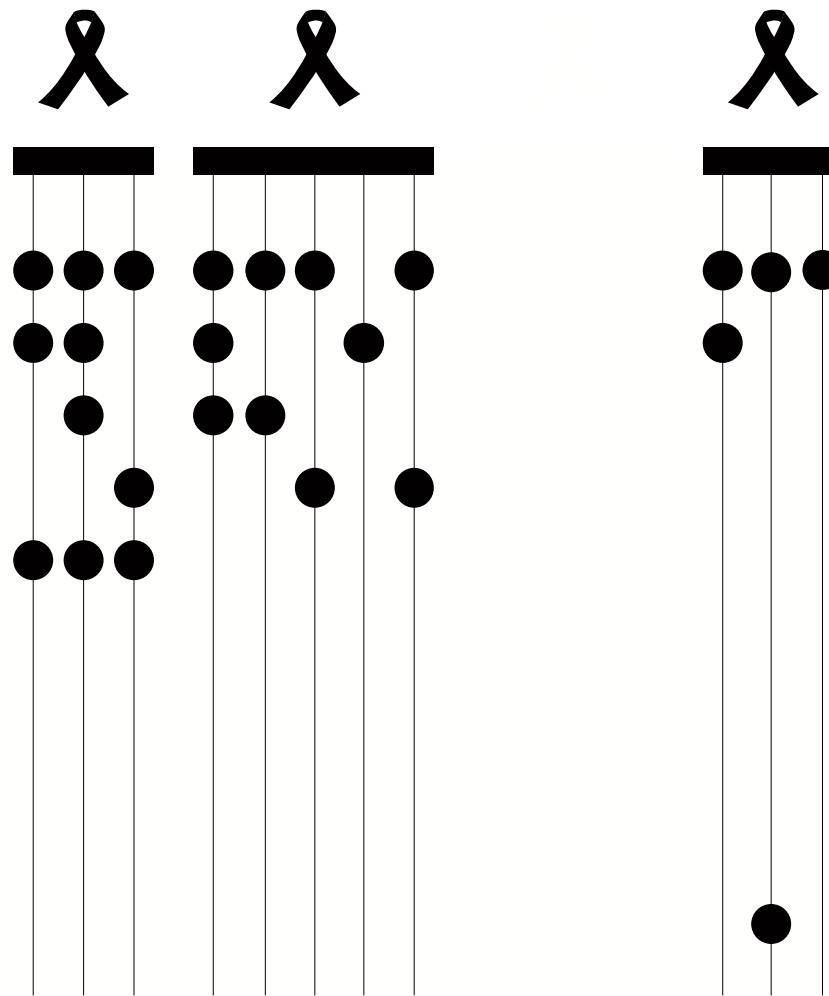
XX XX XX
XX XX XX

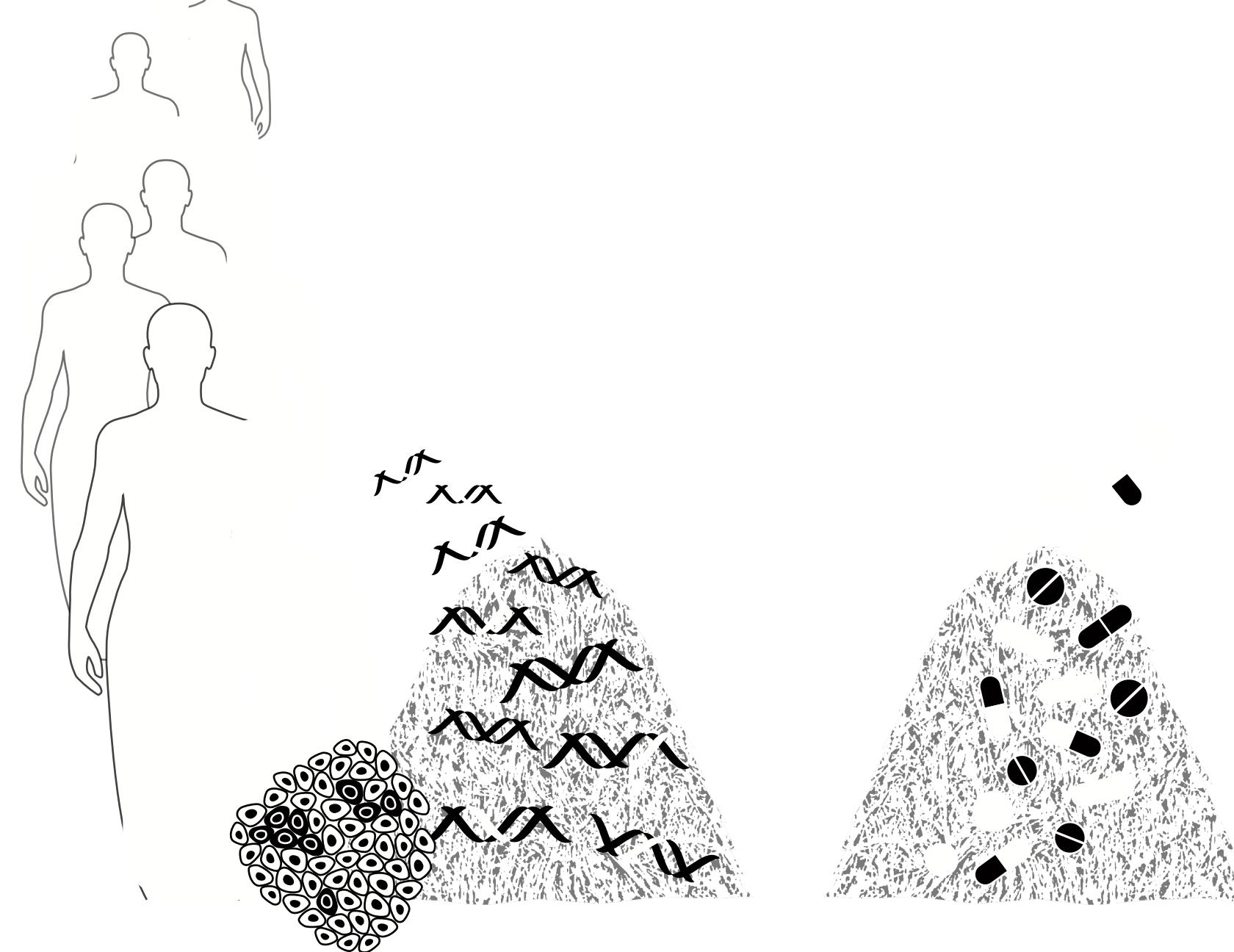
Many mutations

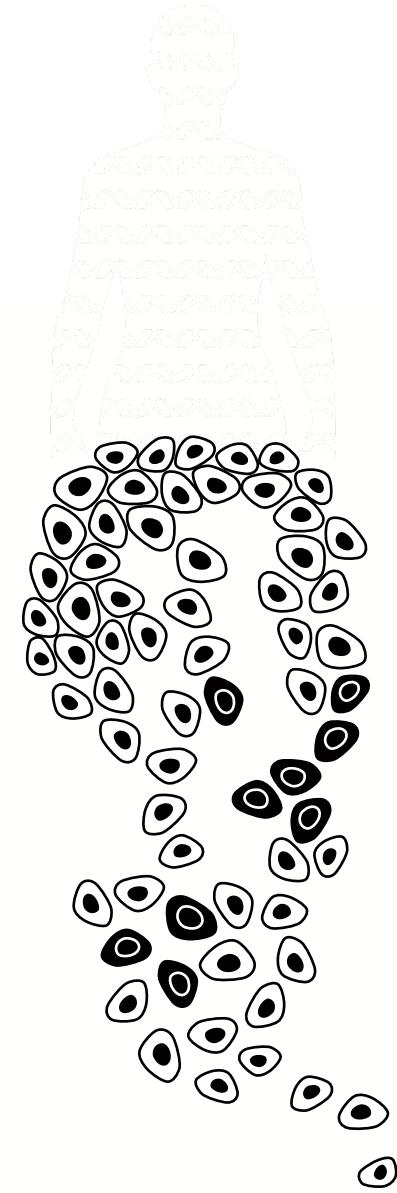
Mutation



Cancer type

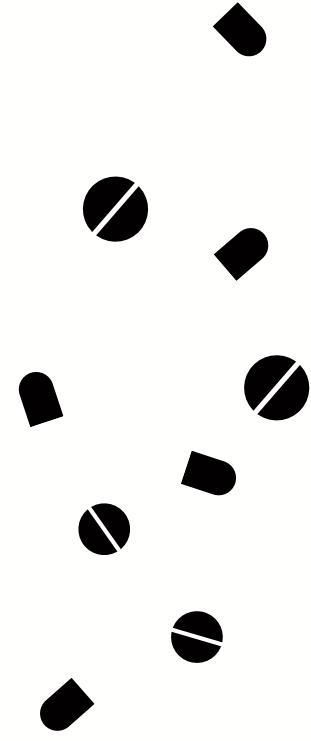






Taskén, Munthe, Tjønnfjord, Schjesvold, Frigessi

Hematology pipeline project: Taskén / Munthe / Tjønnfjord
& Schjesvold / Frigessi groups / NOR-OPENSCREEN

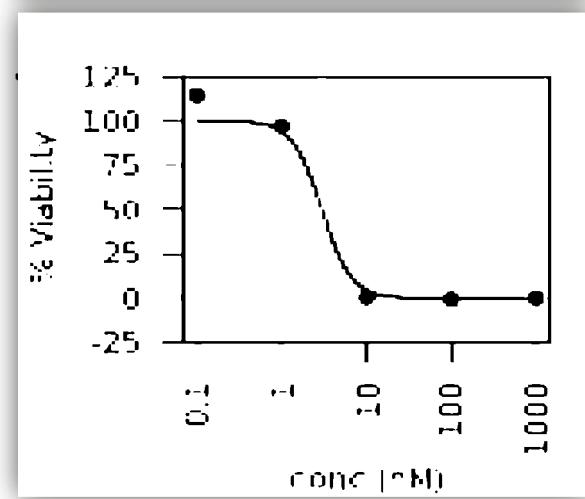
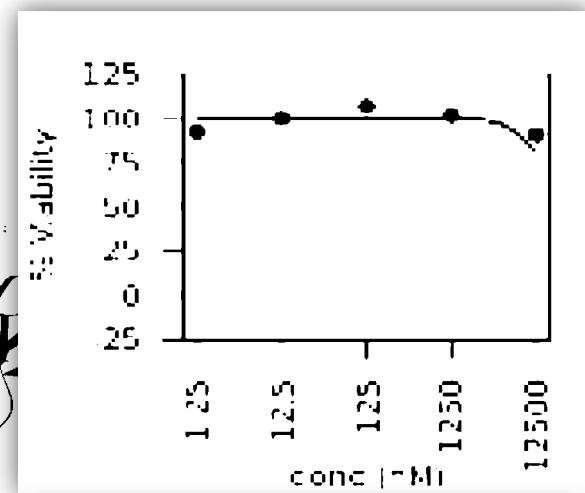


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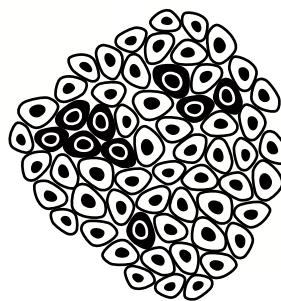
Oslo
University Hospital

Hematology pipeline project: Taskén / Munthe / Tjønnfjord
& Schjesvold / Frigessi groups / NOR-OPENSCREEN

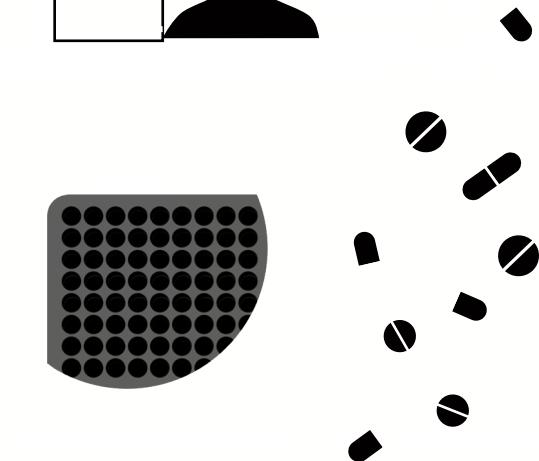
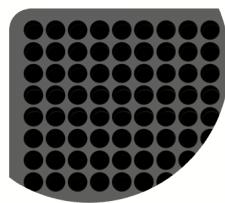


Single drug or combination

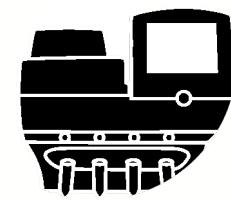
Personalized
therapy



Sample

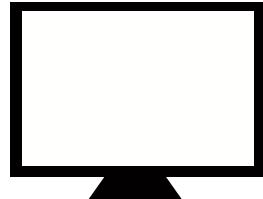


Omics



(...)

New
therapy



Iterative
learning

www.openscreen.no



Latest news

Welcome to NOR-OPENS SCREEN

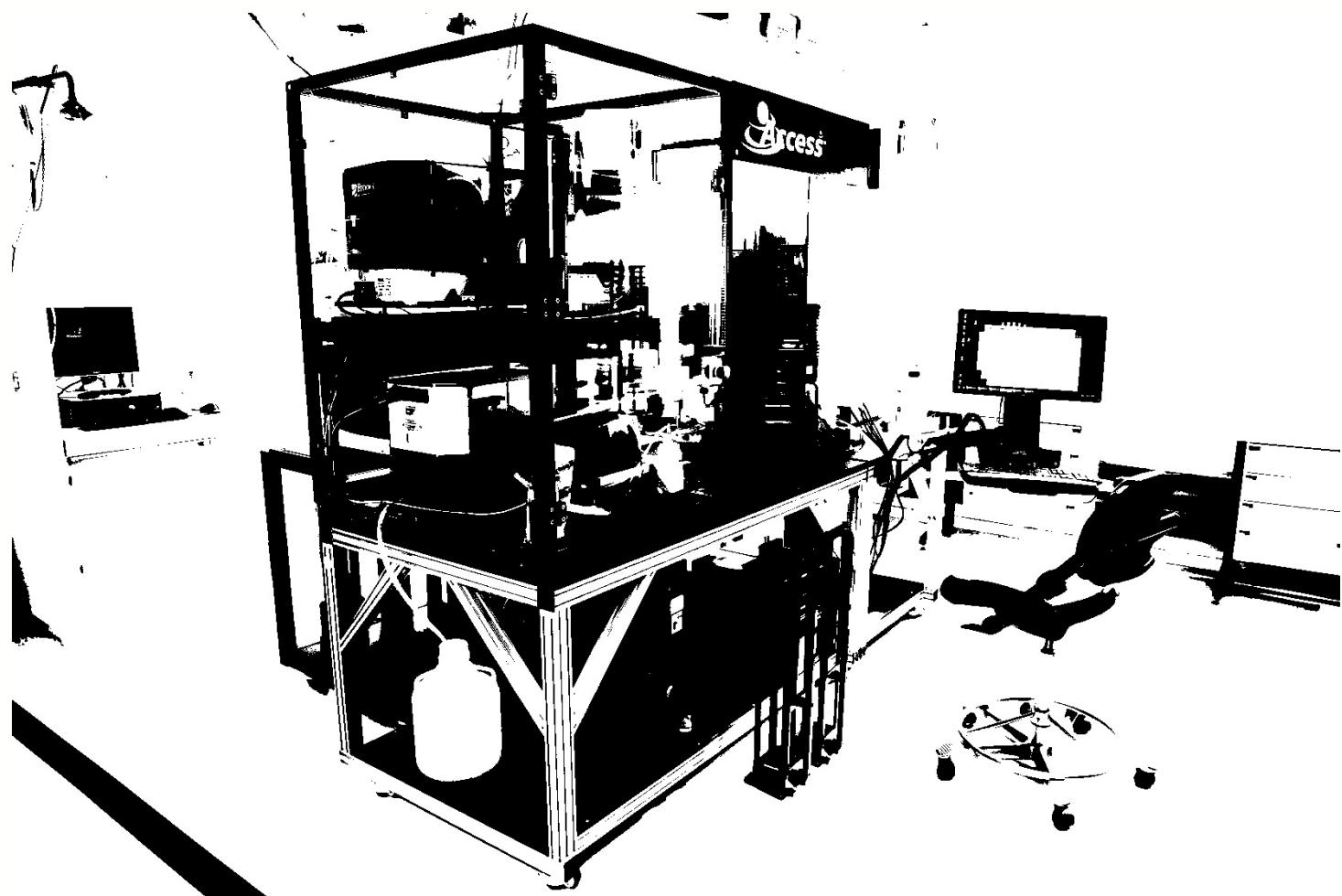
ABOUT US

NOR-OpenScreen successful application

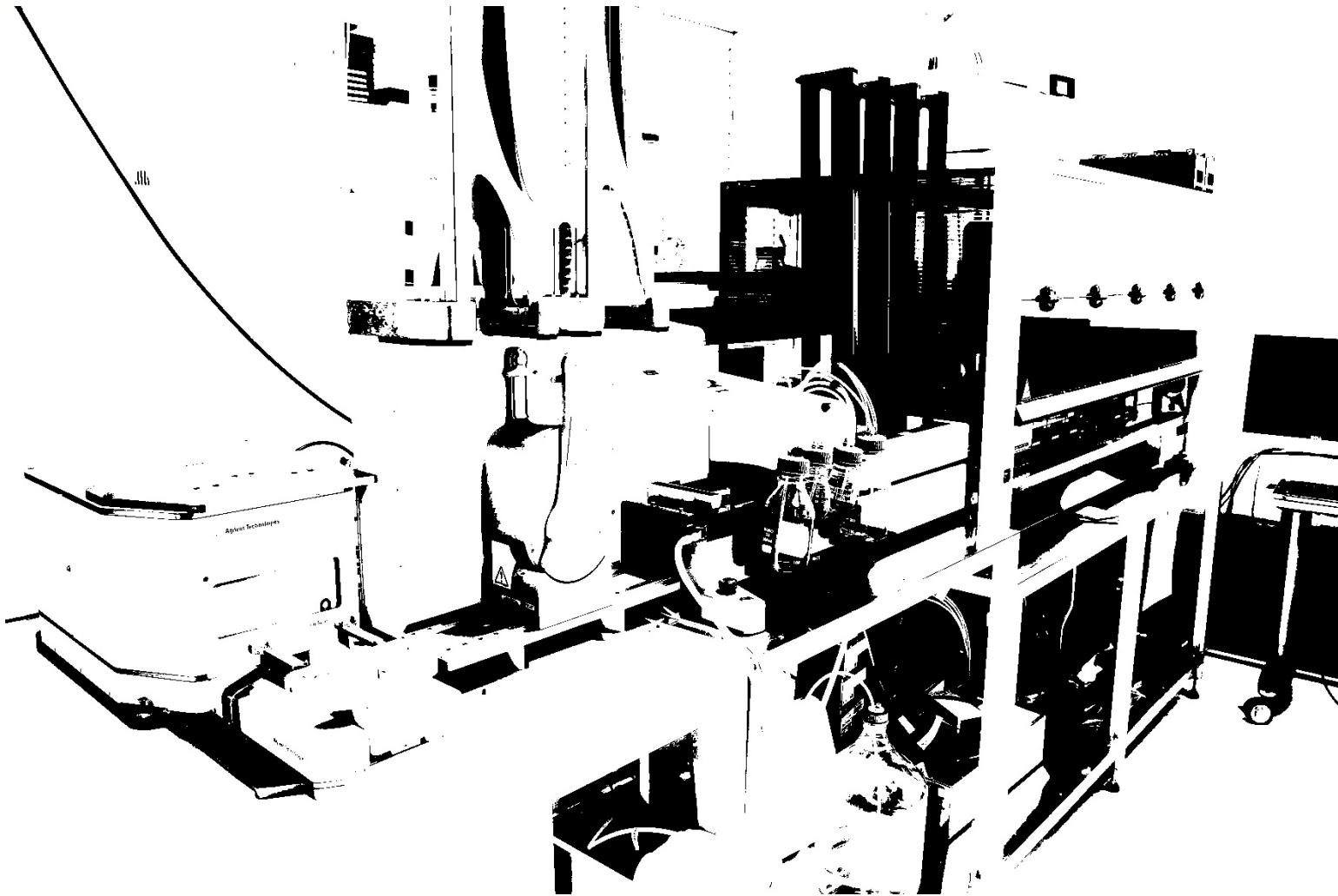
The award of major research funds for "Novel ion chemistry, biology and materials processing by use of E-beam technology" Centre of Excellence Professor Jørgen Tiedje and Øystein Hæg特朗es the highest level of commitment - and forward to the reputation of being one of the best users of existing new technologies and to join in future in the major European initiatives. When EU-OPENS SCREEN is transformed from a research and innovation centre EITP-project to a Eu-research Infrastructure (EFRI) in 2013, the award will still be valid.

Publications

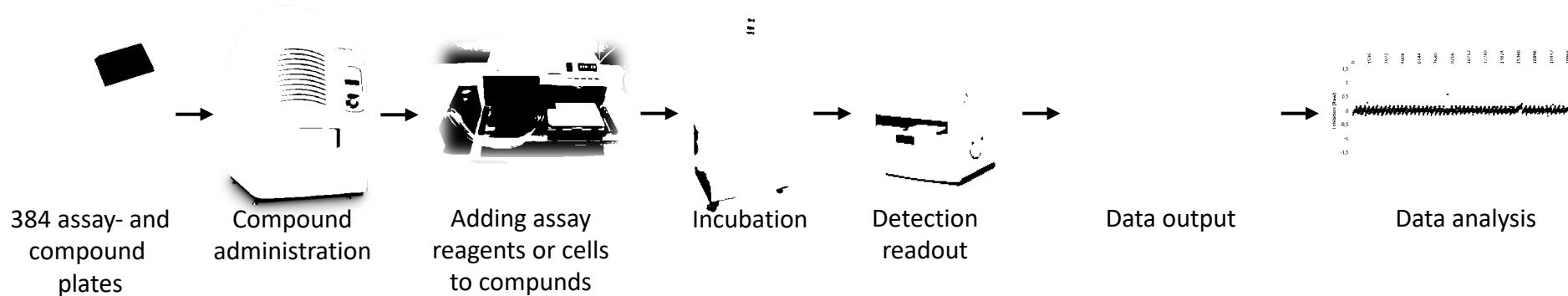
Automated workstation_1



Automated workstation_2



Method – drug sensitivity screens



Sigma LOPAC ¹²⁸⁰	1 280	Library of pharmacologically active compounds.
Prestwick	1 280	Collection of FDA approved drugs
SelleckChem	384	Cambridge Cancer Compound Library, Collection of cancer-targeted compounds
SelleckChem*	193	Known kinase inhibitors
Biomol*	80	Known kinase inhibitors
Biomol*	80	Bioactives
Enzo	477	Target and Pathway Libraries
Tocris Biosciences	1 120	Tocris Tocriscreen Mini, selection of biologically active compounds
SelleckChem	1 650	Bioactive Compound Library
PPI library	~1200	Protein protein disruptors
ChemBioNet	~17000	Diversity collection
Enamine	~28500	Diversity collection
Chembridge	~18000	Diversity collection

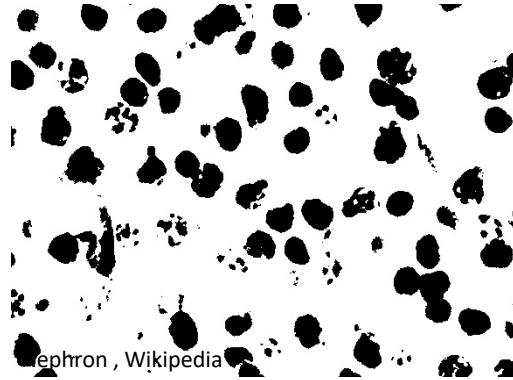


Pipeline for individually tailoring new treatments for relapsed, intractable chronic lymphocytic leukemia and multiple myeloma



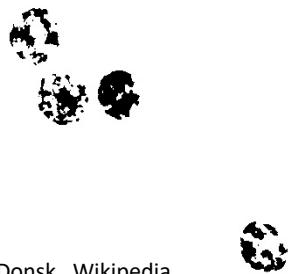
We will develop a fast approach to assist clinical decisions in individualized cancer therapy

Multiple Myeloma (MM)



- Plasma cell neoplasm
- Accounts for 10% of all haematologic malignancies
- Median age at diagnosis: 66 years
- Improved 5 year survival, but still incurable

Chronic lymphocytic leukemia (CLL)



- B cell cancer
- Most common leukemia in the Western world
 - accounts for ~40% of all adult leukemias
- Median age at diagnosis: 72 years
- A slowly progressive disease

A. CLL treatment at present

<70 years and fit no p53-dysfunction	<70 years and fit p53-dysfunction	>70 years and fit no p53-dysfunction	>70 years, unfit or comorbidity no p53-dysfunction	>70 years p53-dysfunction
Chemo-immuno- therapy FCR	Chemoimmuno- therapy HDMP + alemtuzumab Allogeneic stem cell tx.	Chemo-immuno- therapy BR	Chemoimmuno- therapy CR	?? Signal pathway inhibitors (+ mAb)
<2 years	Signal pathway inhibitors + mAb (?) Allogeneic stem cell tx.	Signal pathway inhibitors + mAb (?) Allogeneic stem cell tx.	Signal pathway inhibitors (+ mAb)	?? Signal pathway inhibitors (+ mAb)
>2 years	Retreatment	?? Signal pathway inhibitors (+ mAb)	Retreatment	Retreatment ?? Signal pathway inhibitors (+ mAb)
	??	??	??	??

B. Multiple myeloma treatment at present

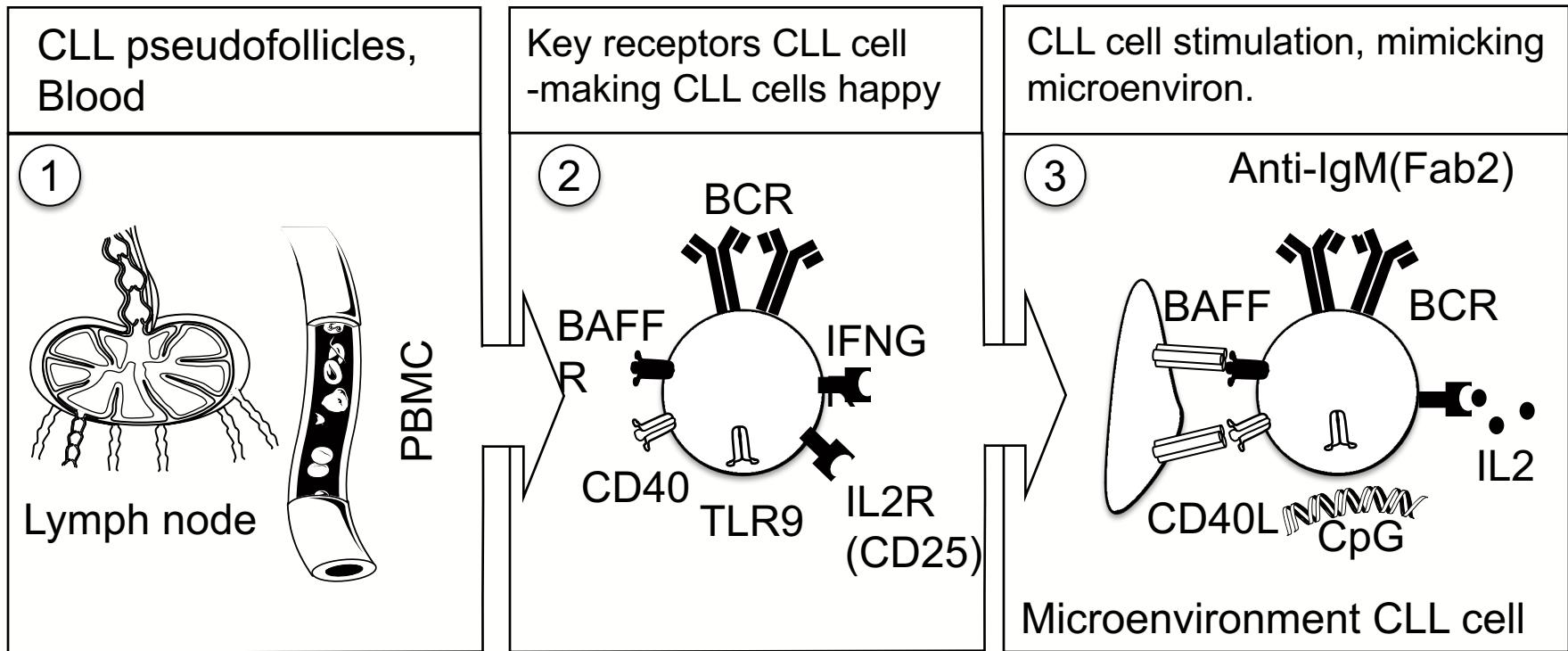
<65 years of age	>65 years of age
Induction with Cy/Vel/Dex, HMAS consolidation	MPT
Retreatment <1 year	Thal/Dex, Len/Dex
Retreatment >1 year	Reinduction HMAS consolidation
	Retreatment
	??
	??

Cy: cyclophosphamide; Vel: Velcade; Dex: Dexamethasone
MPT: Melphalan, prednisone, thalidomide
Thal: Thalidomide; Dex: Dexamethasone; Len: Lenalidomide

Chemoimmuno-therapy. FCR = fludarabin + cyclophosphamide, BR = bendamustine + rituximab, CR = chlorambucil + rituximab
Signal pathway inhibitors = ibrutinib (Bruton's kinase inhibitor), idelalisib (PI3K inhibitor), fostamatinib (SYK inhibitor)
 mAbs = anti-CD20 antibodies rituximab, ofatumumab, obinutuzumab and anti-CD52 antibody alemtuzumab

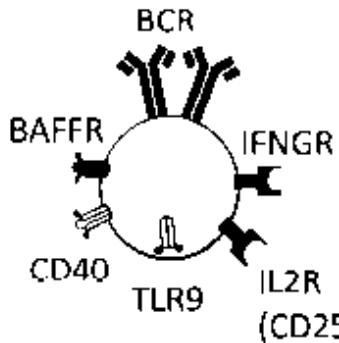
Figure 3. Treatment options for MM and CLL at present

WP1. Bioassays and Drug Discovery. CLL

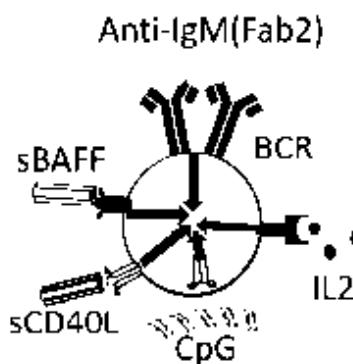


Ludvig A. Munthe

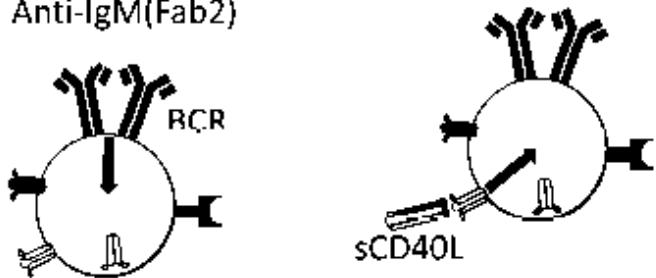
A) Unstimulated. Relevant Receptors for initiation of signaling pathways CLL



B) Global activation



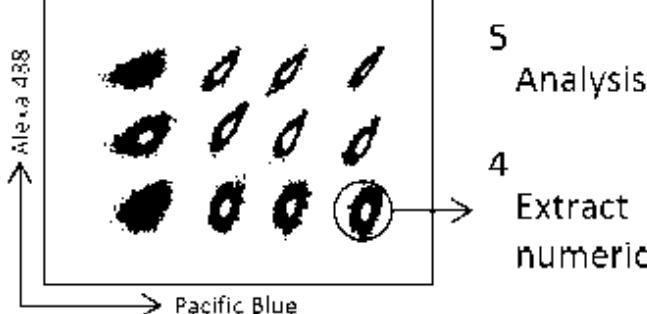
C. Specific pathways
Anti-IgM(Fab2)



1 Purify and stimulate cells

2 Fix, barcode and stain with Abs

3 Analyze by FACS

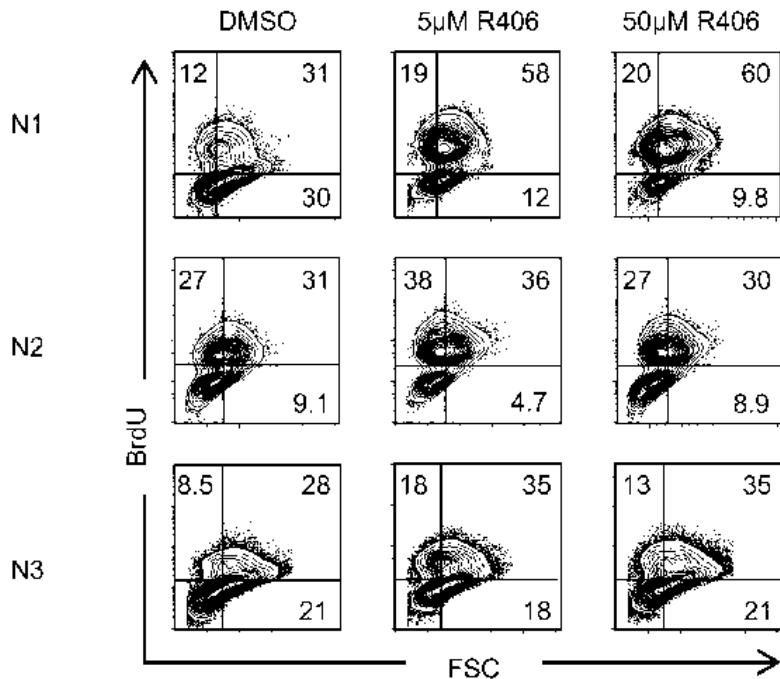


5 Analysis

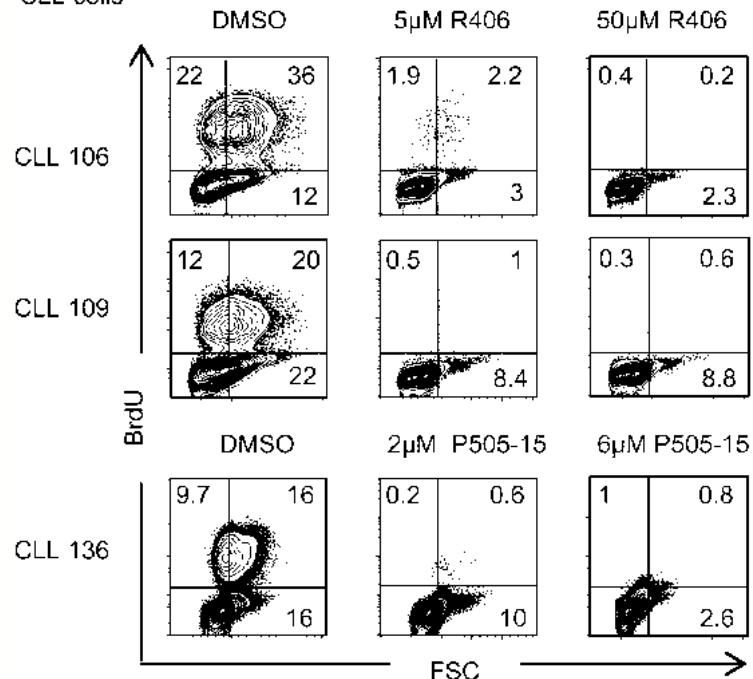
4 Extract numerical data

SYK Inhibitors Reduce CLL Proliferation

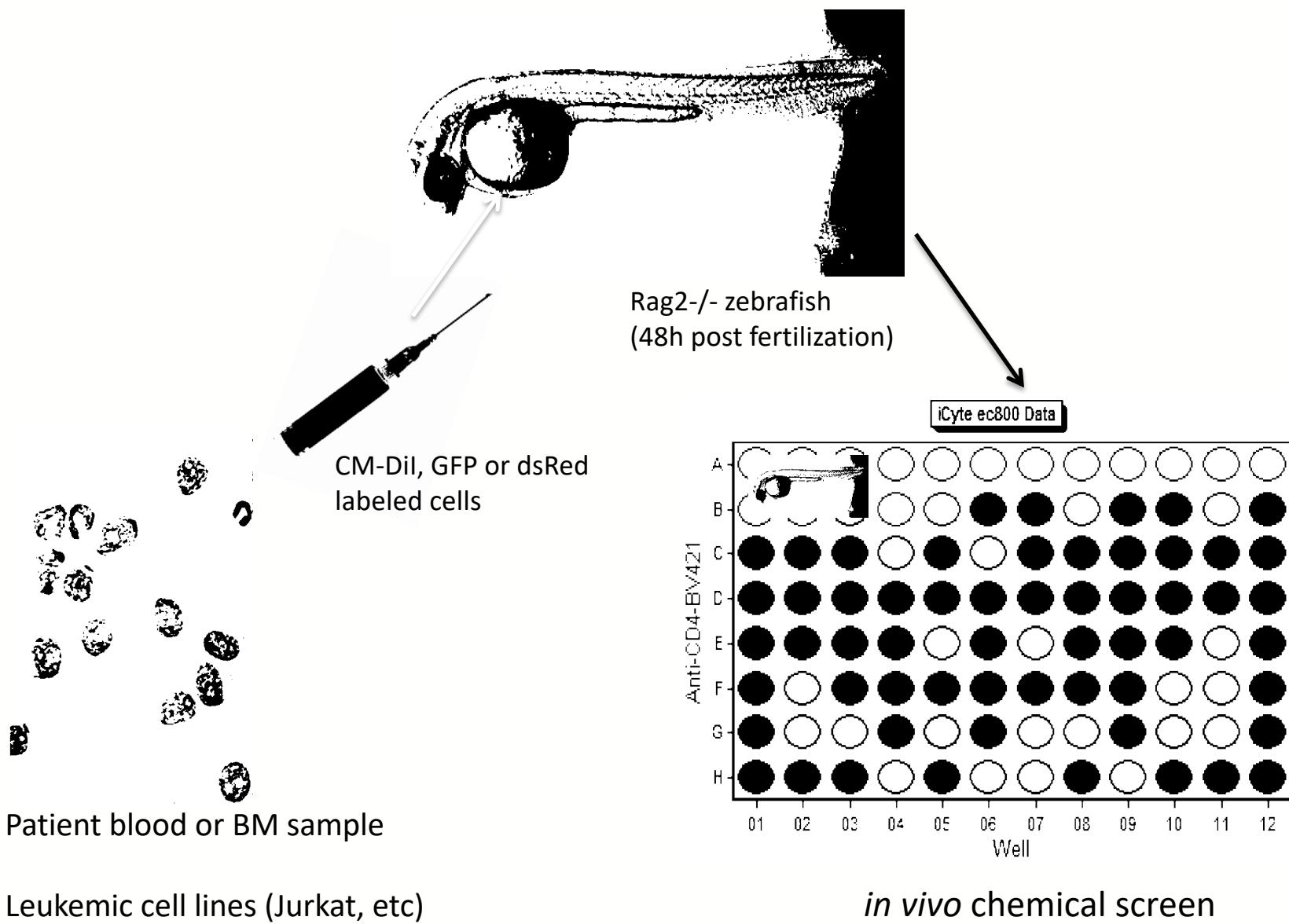
Normal B cells



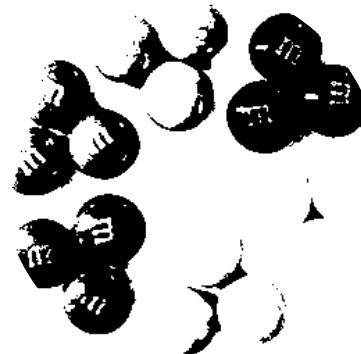
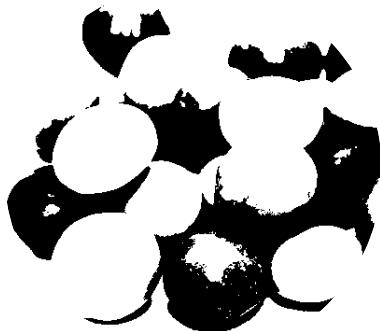
CLL cells



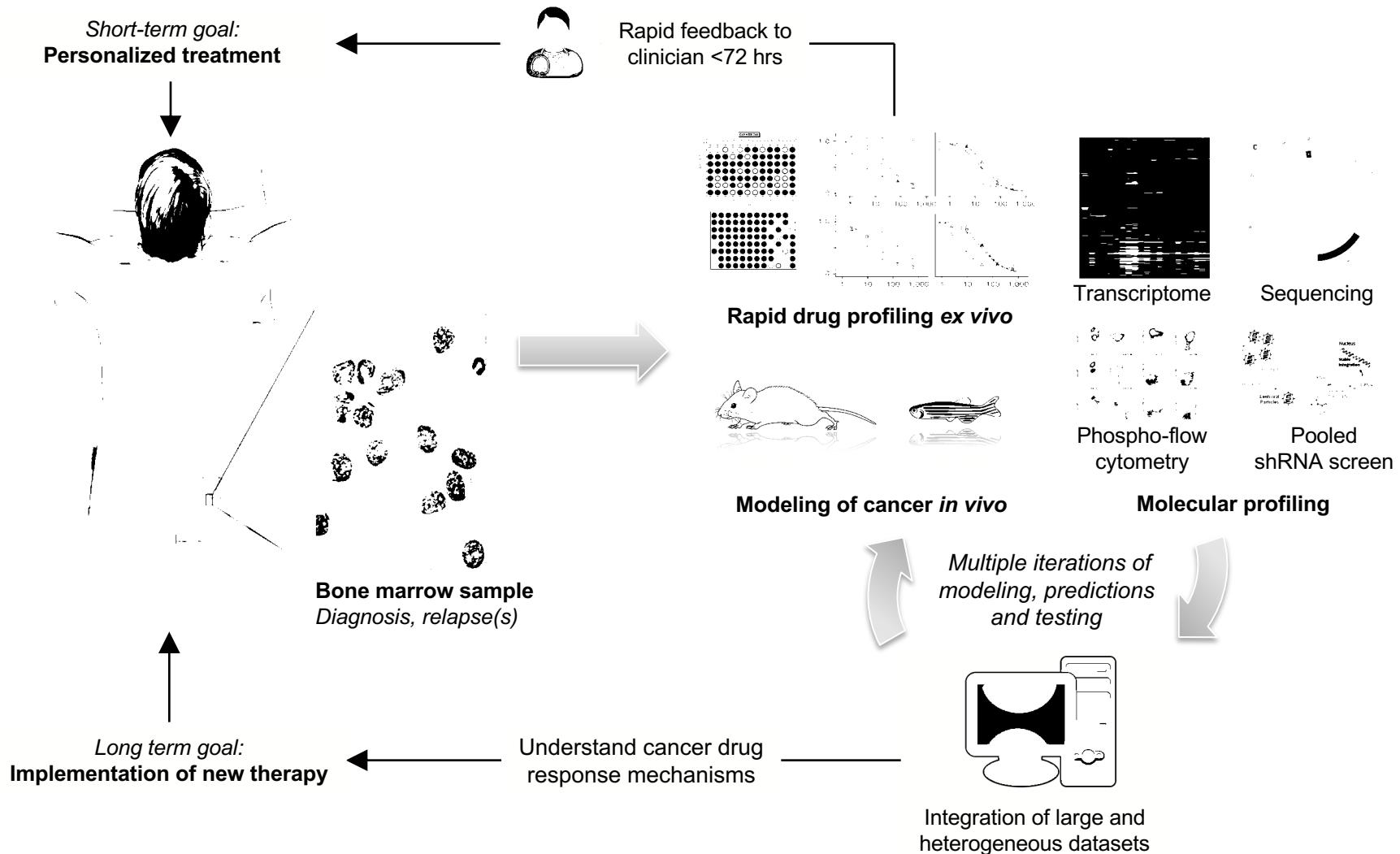
Zebrafish xenotransplantation as *in vivo* screening platform for hematopoietic malignancies



“ Integrate heterogeneous data from
pFLOW analyses
and drug sensitivity screens
in order to predict patient responses



Improving cancer treatment through personalized medicine and iterative learning



Protokoll – individuell behandling

A PHASE 1 MULTICENTRE OPEN-LABEL STUDY OF
CANCER CELL SCREENING IN PERSONALIZED
MALIGNANCIES

Forespørsel om dokumentasjon

Nr beta

TITTEL: En fase I multisenter, åpen, studie med individuell behandling av B-celle kreft

[A phase 1 multicentre open-label study of personalized cancer cell screening in individualized malignancies]

Alkylerende midler:

Cyklofosfamid, klorambucil er prototypene

Bendamustin er mye brukt nå

Ifosfamid, trofosfamid, melfalan, busulfan, treosulfan kan være aktuelle

Antracykliner:

Doksorubicin er prototypen

Epirubicin, idarubicin, daunorubicin og mitoksantron kan være aktuelle

Folsyreantagonister:

Metotreksat

Purinanaloger:

Merkaptopurin, fludarabin, cladribin, pentostatin

Vinkalkaloider:

Vinkristin, vinblastin

Kortikosteroider:

Prednison, metylprednisolon

Immunmodulerende medikamenter:

Lenalidomid

Signalveisemhemmere:

Brutonkinase hemmere; ibrutinib

PI3K-hemmere; idelalisib

SYK-hemmere; fostamatinib, P505-15, entospletinib

Protein-kinase hemmere; flavopiridol

BCL-2 hemmere; oblimersen, obatoclax, ABT-263

Monoklonale antistoffer:

Anti-CD20 antistoffer; rituximab, ofatumumab, obinutuzumab

Anti-CD52 antistoffer; alemtuzumab

Anti-CD23 antistoffer; lumiliximab

Anti-CD38 antistoffer; daratumumab

Anti-CD40 antistoffer; lucatumumab

Pipeline for treatments in *Heterogeneous disease pathways and test sensitivity screening support clinical development*

Professor Kjetil Taskén
Centre for Molecular Medicine Norway

Oslo, 10th of October 2016

RE: User-participation in project on development of individualised therapies for hematological cancer

This letter to confirm that Blodkreftforeningen (The Blood Cancer Society in Norway) is informed about the project application from Professors Kjetil Taskén, Jorrit Enserink and Arnoldo Frigessi and colleagues entitled "*Pipeline for individually tailoring new treatments in hematological cancers*" to the Norwegian Research Council.

We have reviewed the project synopsis and understand that this proposal will gather several ongoing research projects that directly screen patient samples for drug sensitivity and will aim to provide tools to predict and find optimal combinations of drugs for each patient to assist clinicians in individualizing therapies for patients that have exploited other treatment opportunities.

We are enthusiastic about these plans as we foresee potential future patient benefit for Norwegian patients. We have been asked to participate in user activities relating to the project (steering committee and user discussion panels). We will be happy to participate and help setting up such activities as the project develops.

Best regards
Blodkreftforeningen



Eddy Grønset
Head of secretary

Cancer Drug Sensitivity Screening - Utfordringer

- Liste med legemidler som kan virke, evt kombinasjoner som kan gi synergi
- Støtte kliniske beslutninger om individualiserte behandlingsvalg
- Lister med legemidler som kan være fra 20 til 400 legemidler i bruk eller i utvikling
- “Off-label” bruk, ukjente interaksjoner
- Randomiserte studier vil teste algoritmen for utvalg av legemidler, ikke de enkelte legemidler
- Internasjonale konsortier for å aggregere data om hvert enkelt legemiddel? – “Basket design?

SKREDDERSYDD MEDISIN vil føre til en redefinisjon av begrepet "*evidens-basert medisin*" siden det ikke bare vil være mulig å bestemme *om* en behandling har effekt men også *hvem* som har nytte av behandlingen.

Kjetil Taskén group

NCMM, UiO/OUH

NCMM



The Research Council
of Norway



Oslo
University Hospital



SEVENTH FRAMEWORK
PROGRAMME

HELSE



norden

NordForsk

• SØR-ØST

EMBL .



NORWEGIAN CANCER SOCIETY



K.G. Jebsen Inflammation
Research Centre



KG JEBSEN CENTER
for Cancer Immunotherapy

novo nordisk fonden

Prosjekt-team



NCMM, UiO/OUS

Kjetil Taskén

Andrea Cremaschi

Johanne Uthus Hermansen

Deepak Balaji Thimiri

Govinda Raj

Sigrid Skånlund

Andre partnere:

Bjørn Tore Gjertsen

Jorrit Enserink



**Oslo
University Hospital**

Inst. for Immunologi, OUS

Ludvig A. Munthe

Dong Wang

Avd for Blodsykdommer, OUS

Geir E. Tjønnfjord

Fredrik Schjesvold

***Oslo senter for biostatistikk
og epidemiologi, UiO/OUS***

Arnoldo Frigessi

Manuela Zucknick

NCMM

Centre for Molecular Medicine Norway

Nordic EMBL partnership for Molecular Medicine

From disease mechanisms to clinical practice