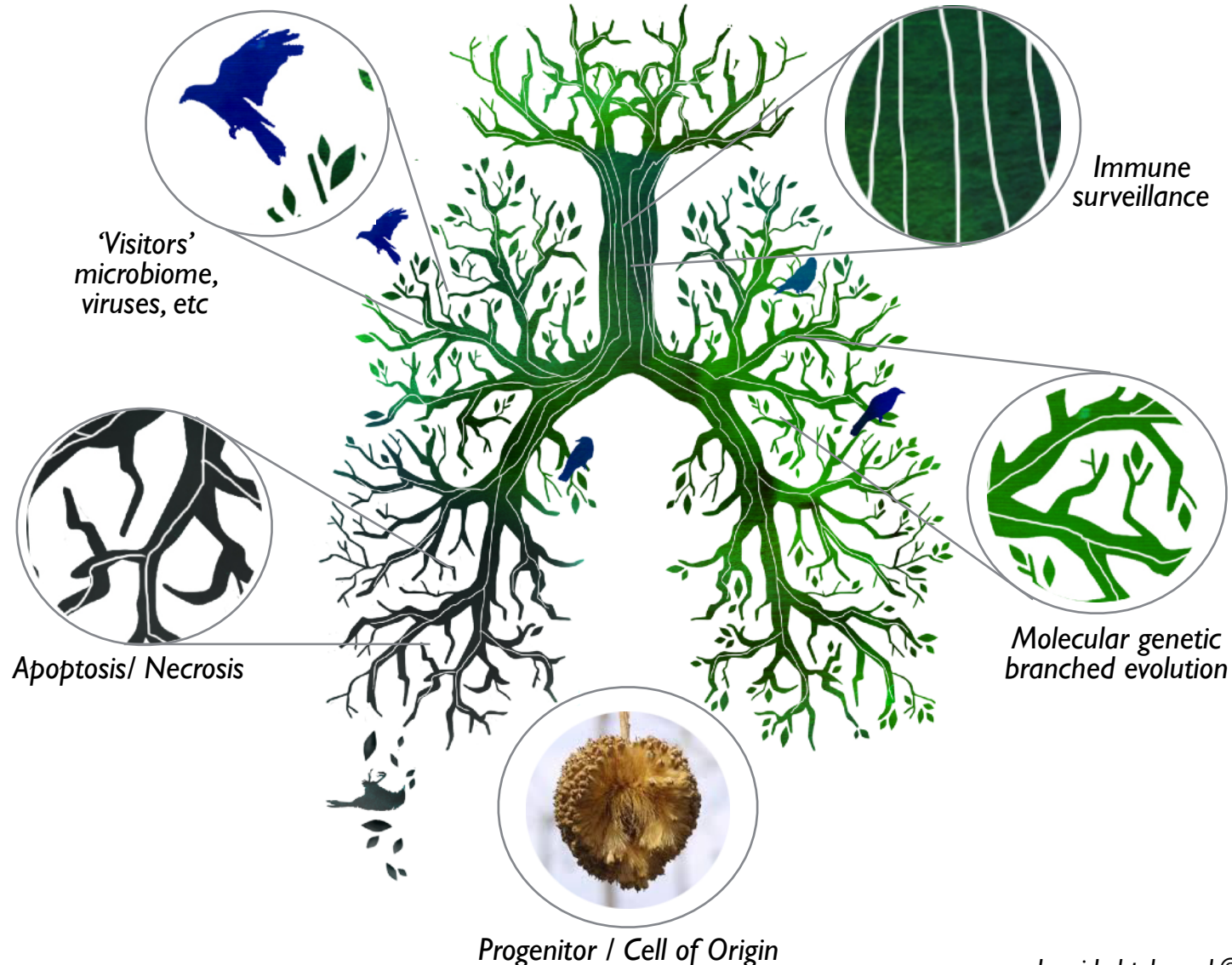




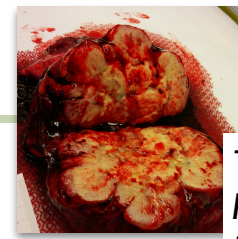
Personalised Phenotypic Diagnostics to Benefit Lung Cancer Patients

Emmy W. Verschuren, PhD
Academy of Finland R'Life Kick-off meeting
November 27 2020

Cancers are Ecosystems that Evolve via a Sequence of Dynamically Changing Environments

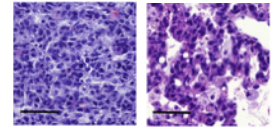


Questions in Translational Cancer Research



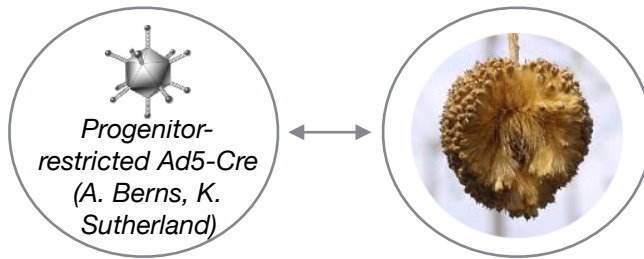
Tissue context ?
Passenger vs driver mutation ?
Intra-tumour heterogeneity ?
Lifetime mutagenic load ?

- ❖ Do **phenotypes** converge as targetable disease vulnerabilities - *histopathology*
- ❖ What can we learn from **primary cell & tissue** analysis *ex vivo* - *IMI-PREDECT*
- ❖ How can **preclinical models** inform on **clinical diagnostic advances** - *Ac. of Finland*

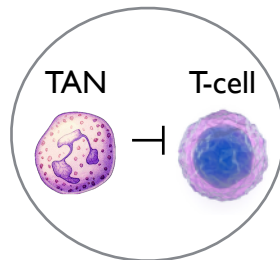


Meilahti University of Helsinki Medical Campus

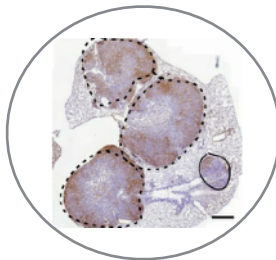




Progenitor cell and genetic drivers
cooperatively define NSCLC
histopathology spectrum



Histopathology-selective phenotypes:
immune suppression, signalling, metastatic propensity



Signalling activities align more with
histopathology than genotype, and show
significant **spatial heterogeneity**



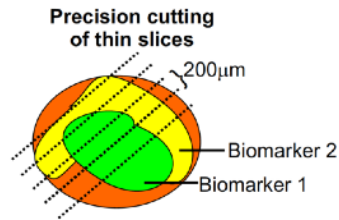
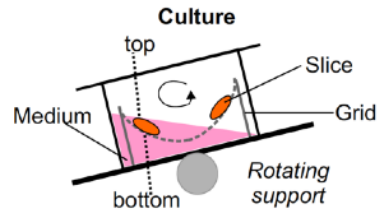
Kras^{G12D} ; *Lkb1*^{fl}
Drivers in appr. 30% human NSCLC
Wide histopathology spectrum

Take home messages
preclinical studies

Translational studies to consider
histopathology-selective phenotypes
in addition to driver mutations

Nagaraj & Lahtela et al, Cell Rep '17
Närhi et al, J Path '18
Bao et al, Sci Rep '19

Leica VT1200S



Take home messages organotypic tissue slice studies

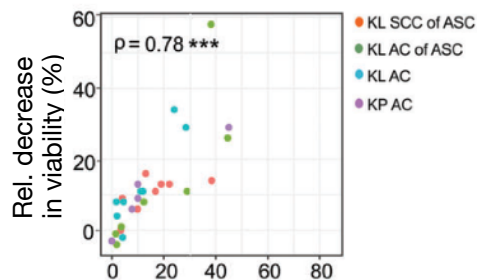
Oxygen and culture supports are required for tissue survival

Combination drug sensitivity in tissue slices relates to spatial signalling activities of targeted pathways

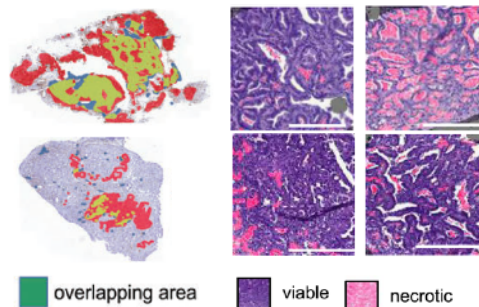
Combination selumetinib (MEKi) + dactolisib (PI3K/mTORCi) treatment

Overlay pERK and p4EBP1

DMSO sel +dact



Overlapping expression area (%) of pERK and p4EBP1



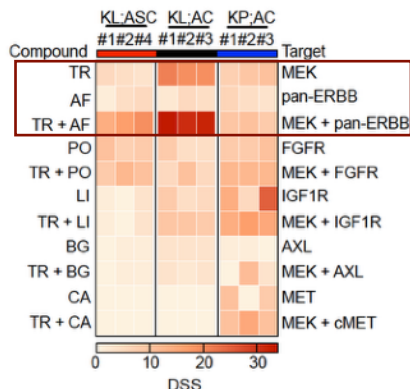
Tissue slice studies/protocols:

Davies et al., Sci. Rep. '15

De Hoogt et al., Sci. Data '17

Nagaraj et al, J Vis Exp '18

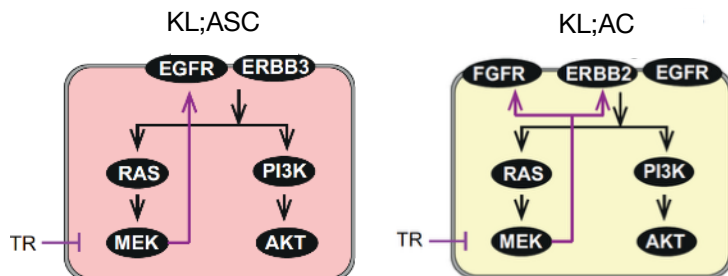
Närhi et al, J. Path '18



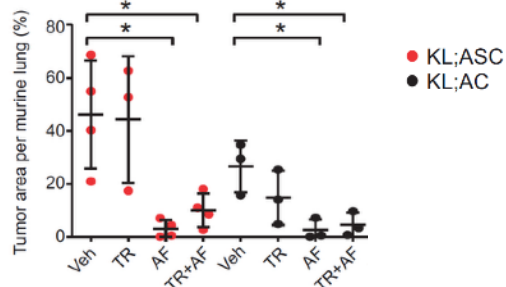
Trametinib (MEKi) + afatinib (pan-ERBBi) combination treatment selective for Kras;Lkb1 NSCLC

Combination drug sensitivity

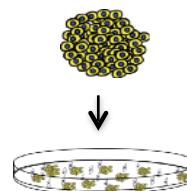
- Relates to **signalling networks selective for NSCLC histotypes**



- Is **validated in vivo**, where increased single pan-ERBBi response corresponds with increased ERBB biomarker activity



Can primary cultures be used to identify and predict drug sensitivities reflective of the native *in vivo* tumour tissue?



Early passage primary epithelial cultures (conditionally reprogrammed cells; CRC/PDC; Schlegel et al)

- ❖ Only half of all resected clinical NSCLC tumours are sliceable, yielding limited numbers of short-lived already necrotic tissue slices, compromising robust study
- ❖ Primary epithelial cultures are established at low success rates (10%), and this takes 2-3 months, possibly leading to genetic and phenotypic drift
- ❖ Surgically resected tissue is not the disease entity to treat; the majority (>70%) of patients are diagnosed with metastatic stage disease -> assays to be adapted to biopsies or pleural effusions

Towards research translation

Challenges of functional diagnostic modelling with clinical samples

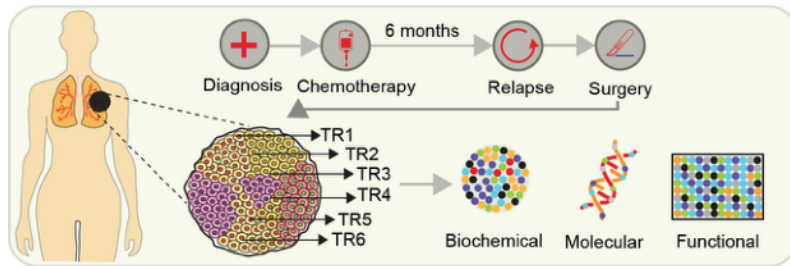


NSCLC - adenocarcinoma

EML4-ALKv3 fusion (driver in 4-7% NSCLC); TP53 mutation (R175H)

chemoresistant (cisplatin + pemetrexed)

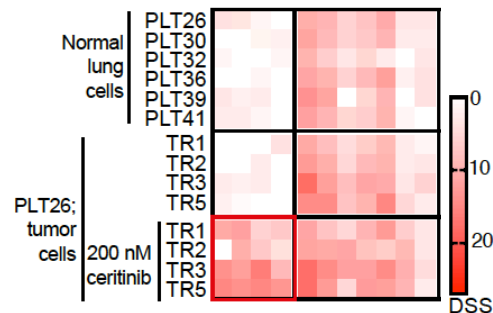
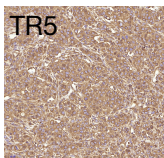
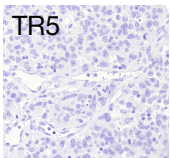
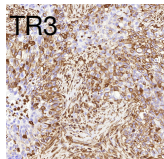
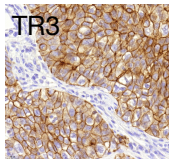
Female never smoker



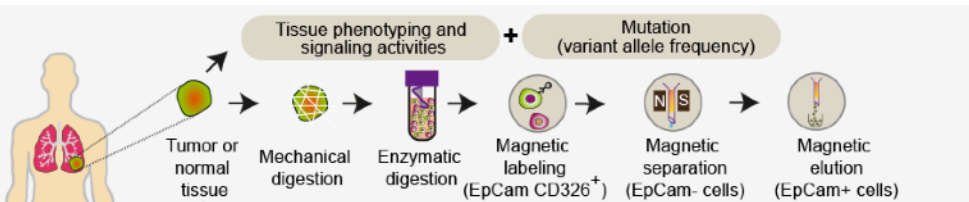
Towards research translation a case study

**Target inhibition enhances Alki (ceritinib) sensitivity,
without affecting normal epithelial cells**

TR3: epithelial (E-cadherin) TR5: EMT/ mesenchymal (CK18)



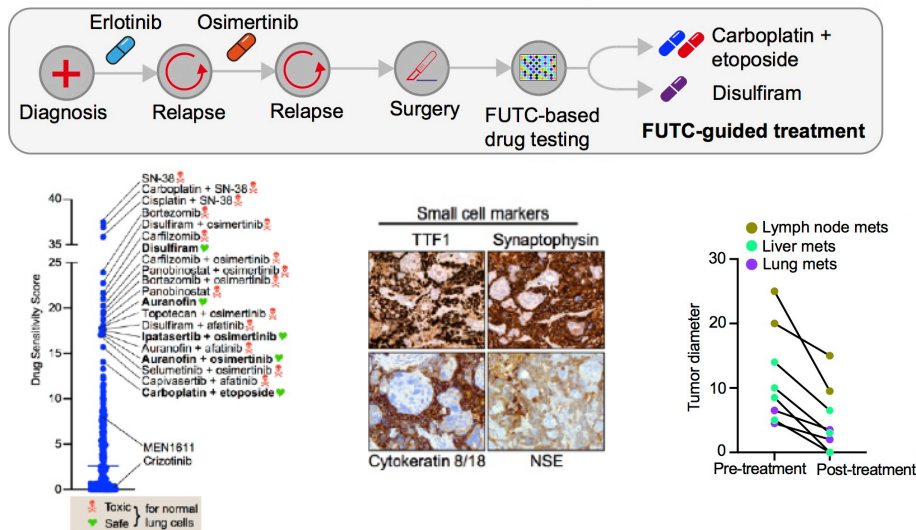
Functional diagnostic profiling using **Fresh Uncultured Tumour Cells (FUTCs)**



- ❖ Normal and tumour **epithelial (EpCam⁺) and stromal (EpCAM⁻ cells)**
- ❖ Drug sensitivity profiling **within three days**
- ❖ **Validation:** mimic of pharmacological and adaptive signalling profiles of murine histopathology subtype-matched cultured cells
- ❖ **Clinical sample-derived FUTCs show drug response matched to driver mutations (EGFR, ALK, MET, KRAS) in 18/19 cases**

Towards research translation *circumventing culture challenges*

FUTC profiling-guided compassionate treatment of a chemorefractory metastatic EGFRmut NSCLC patient



- ❖ Patient was scheduled to receive pemetrexed + carboplatin
- ❖ FUTC profiling showed **etoposide + carboplatin** as superior combination
- ❖ Treatment was adjusted, **small-cell histotype conversion** was confirmed 2 wks later
- ❖ Patient shows response to 4 cycles of treatment over period of one year

- ❖ Develop NSCLC organoid cultures (in collaboration with HUB, Utrecht)
- ❖ Compare FUTC responses to those seen in primary organoids
- ❖ Functionally interrogate adaptive signalling networks to understand primary/acquired resistance mechanisms
- ❖ Extend diagnostic applications to biopsies & pleural effusions

Future directions in R'Life

Thanks to the Team !

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PhD

Nora Linnavirta

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Ashwini Nagaraj, PhD
Katja Närhi, PhD
Jennifer Devlin, PhD
Elina Parri, BSc
Barun Pradhan, BSc
Annabrita Hemmes

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Elissar Al Kazzi (ITN student)
Bassel Alsaed (MSc student)



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Krister Wennerberg - *Drug sensitivities*

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Kaisa Salmenkivi, HUS - *Pathology*

Jari Räsänen, HUCH - *Surgical tumours*

Aija Knuuttila, HUCH - *Clinical oncology*

Jon Lømo, Oslo Uni Hospital - *Pathology*

Lars Søråas, Oslo - *Compassionate care case*

Johan Lundin - *Digital pathology, IMI-PREDECT*

Riku Turkki - *Necrosis analysis, IMI-PREDECT*

Sami Blom - *Systems Pathology, IMI-PREDECT*

Teijo Pellinen - *Multiplexed staining, IMI-PREDECT*

Astrid Murumägi - *CRC methodology*

Simon Anders - *Bioinformatics*

Peter Horvath - *Imaging software*

Preclinical and clinical patients

IMI-PREDECT colleagues

John Hickman, Servier

Wolfgang Sommergruber, Boehringer Ingelheim

Emma Davies & Simon Barry, AstraZeneca

Heiko van der Kuip & Meng Dong, RBMF

Jan Trapman & Petra van Duijn, EMC

Julia Schuler, Oncotest/Charles River

Wytske van Weerden

& Hanneke van Zoggel, EMC

FIMM TC & HTB unit Support

Pekka Ellonen - *DNA seq*

Matti Kankainen - *Bioinformatics*

Swapnil Potdar - *DSRT analysis*

Laura Turunen - *Drug plates*

Jani Saarela - *Assay development*

Current



Previous

