Cellular responses to low dose radiation

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Outline

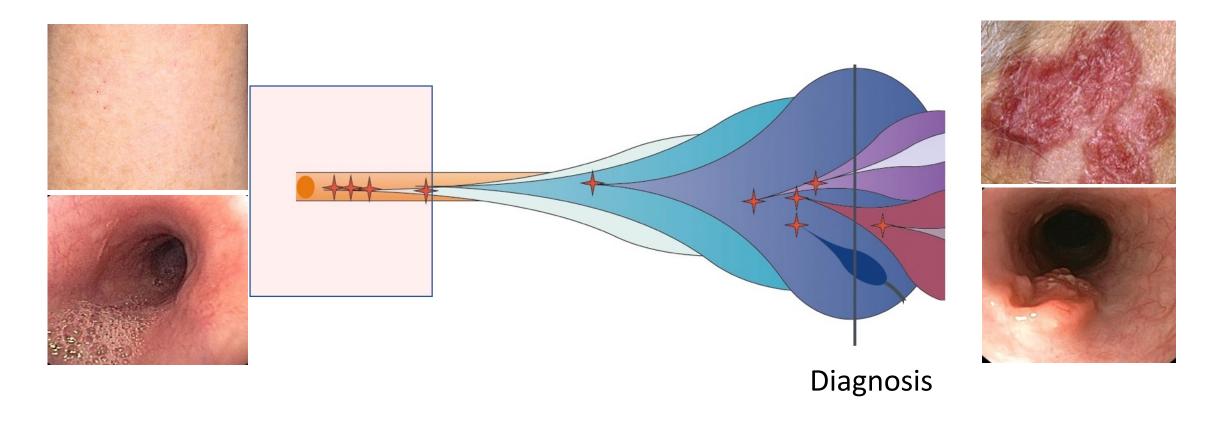
Mapping somatic mutant cells in normal tissues

Application to epidemiology studies?

Somatic mutants in low dose radiation research

Research priorities

Cancer origins: mutant clones in normal tissue

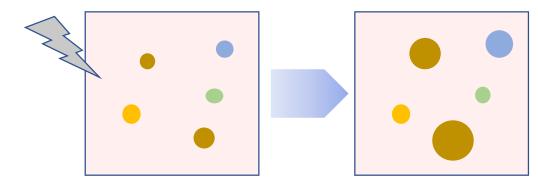


Models of low dose radiation & cancer risk

Damage

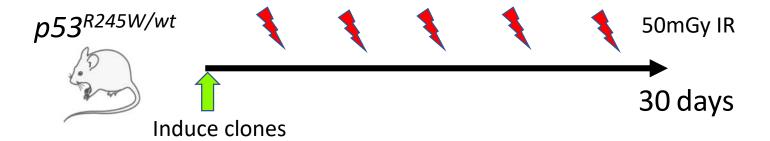
Genome damage induces new clones

Selection

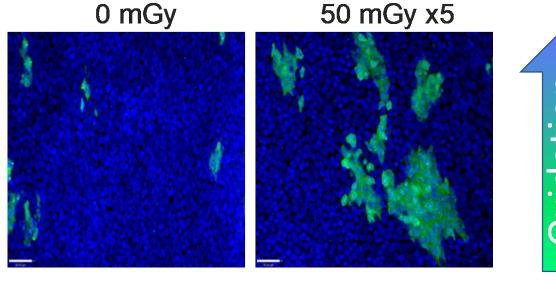


Environmental stress expands pre-existing clones

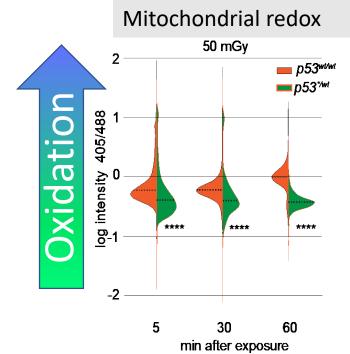
mutant p53 & low dose radiation in mouse oesophagus



Top down views esophagus p53 mutant clones

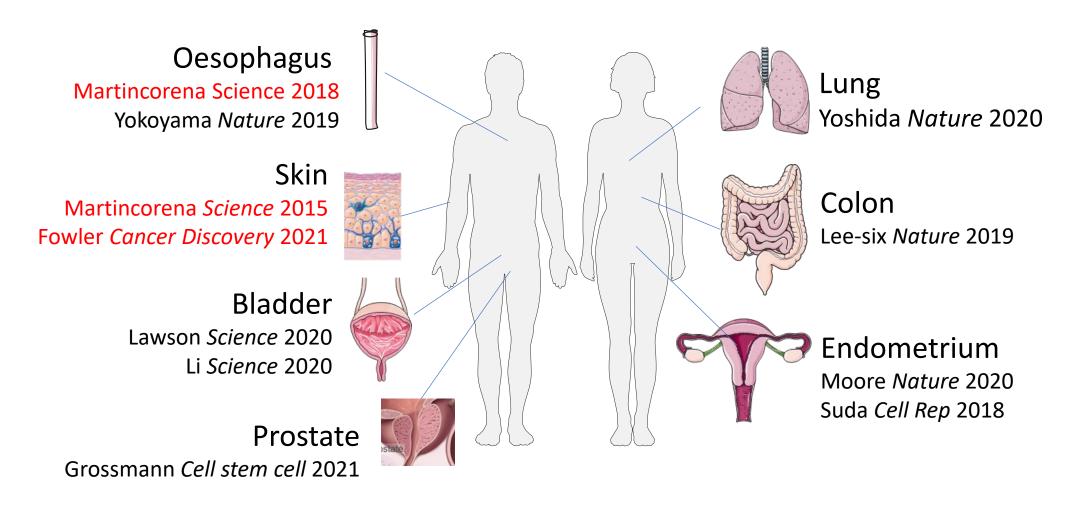


p53 mutation is a gatekeeper to esophageal cancer



Fernandez Antouran Cell Stem Cell 2019

Normal aging epithelia carry somatic mutants....



.....many are linked to cancer

Mapping mutants in normal tissues



Micro dissect clonal units

Whole Genome Sequence Mutational burden
Mutational signatures
Copy number alterations
Mutant prevalence & selection

Lee-six *Nature* 2019 Moore *Nature* 2020 Lawson *Science* 2020



Small area of epithelium

Deep targeted sequence Spatial map
Mutant prevalence & selection
Mutational signatures
(Mutational burden)
Limited copy no. information

Martincorena Science 2015 Martincorena Science 2018



Micro-biopsy

Nanoseq

Mutational burden Mutational signatures

Abscal Nature 2021

Mapping mutants in normal tissues



Micro dissect clonal units

Whole Genome Sequence Laborious
Expensive
High quality data, but few clones

Lee-six *Nature* 2019 Moore *Nature* 2020 Lawson *Science* 2020



Sheet of epithelium

Deep targeted sequence Epithelial sheets only
Cheap
Many 1000's of mutants
Limited clonal & copy no. info
Very sensitive at detecting selection
& changes in mutant prevalence

Martincorena Science 2015 Martincorena Science 2018



Micro-biopsy

Nanoseq

Single molecule resolution Burden/signatures only Works on small samples

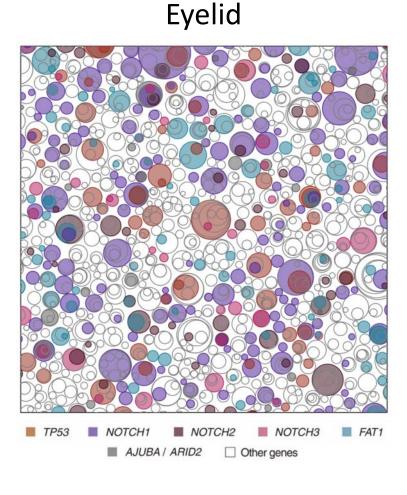
Abscal Nature 2021

Examples: middle aged skin & esophagus

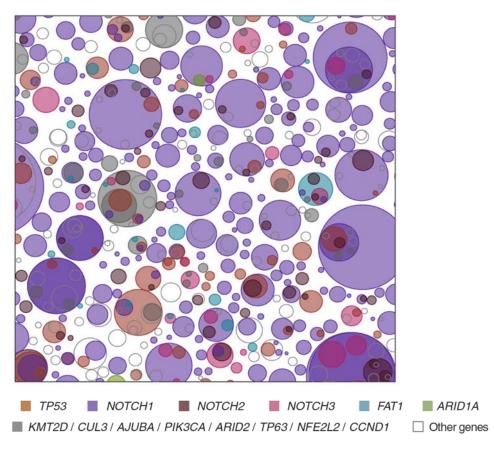
Maps from deep targeted sequencing

Top down views: mutant clones in 1cm² tissue

Coloured circles: positively selected mutants



esophagus

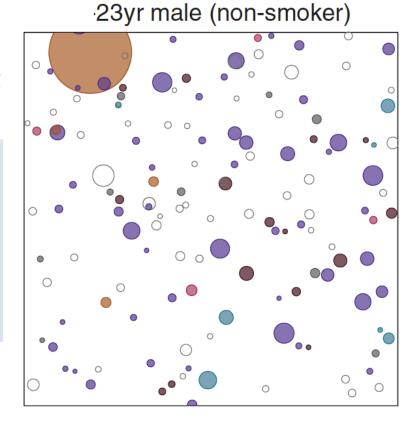


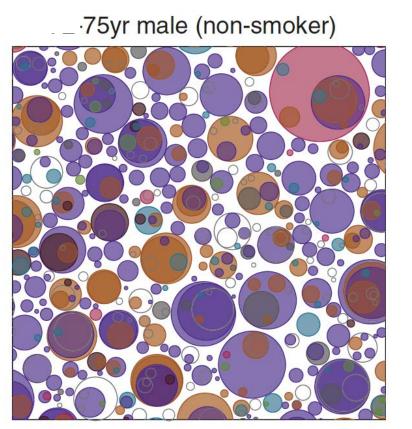
Martincorena Science 2015 Martincorena Science 2018 Fowler Cancer Discovery 2021

Esophagus: Mutants accumulate with age

Maps from deep targeted sequencing

Top down views of esophagus. Coloured circles: positively selected mutants





Martincorena, Fowler et al, Science 2018





NOTCH1





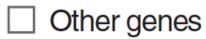
NOTCH3



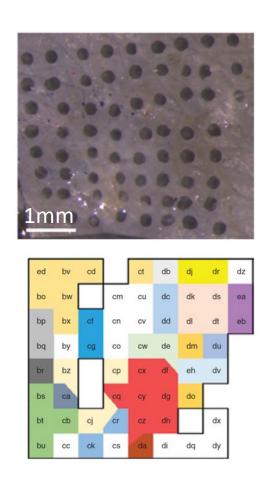
FAT1

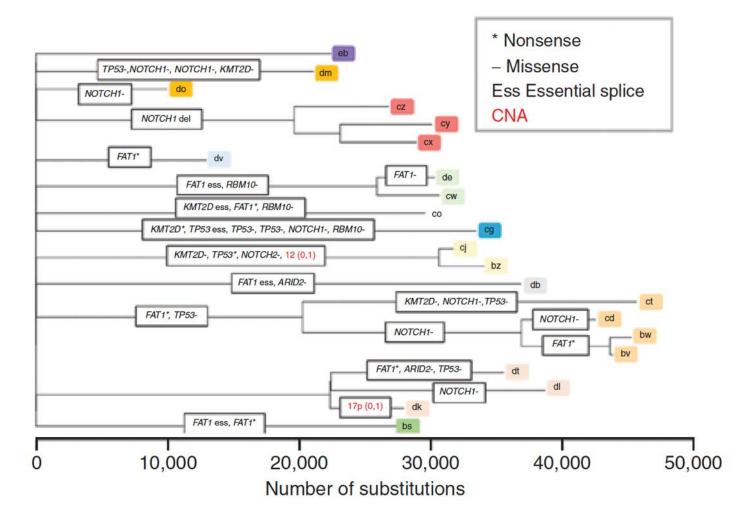


KMT2D / CUL3 / AJUBA / PIK3CA / ARID2 / TP63 / NFE2L2 / CCND1



Normal skin 75 year old: clonal genomes





Mapping mutants in normal tissues

No single simple assay for all tissues

Variation: with age

within tissues

across tissues

between individuals

What shapes the landscape?

Germline Variation

Mutagen exposure

Age

Cell lineage (selection varies markedly across tissues)

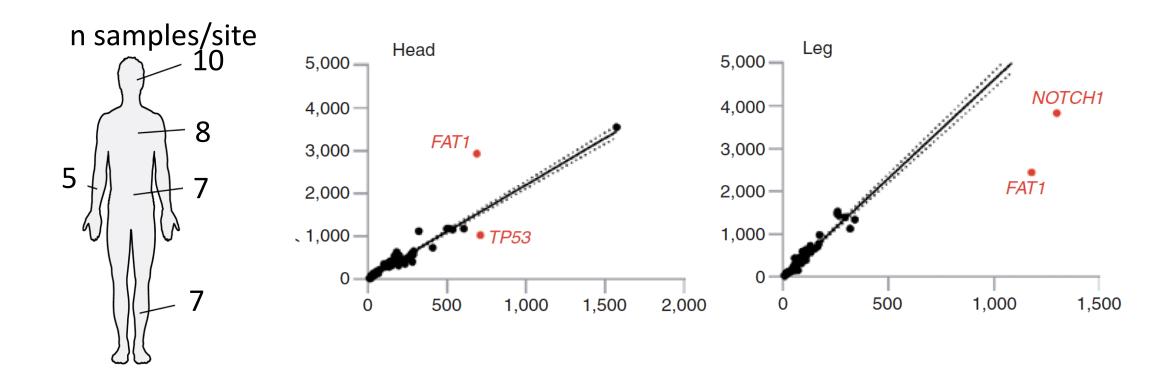
Tissue architecture (crypt, gland or epithelial sheet)

Disease (inflammatory bowel disease, cirrhosis)

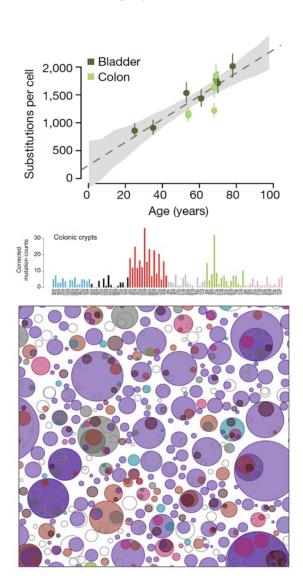
Density of competing clones (less competition=larger clones)

Environmental factors altering fitness UV, IR?

Detecting selection: UV exposed human skin



Epidemiology: Cancer risk prediction



Mutational burden?

Mutational signatures?

Proportion of oncogenic mutants?

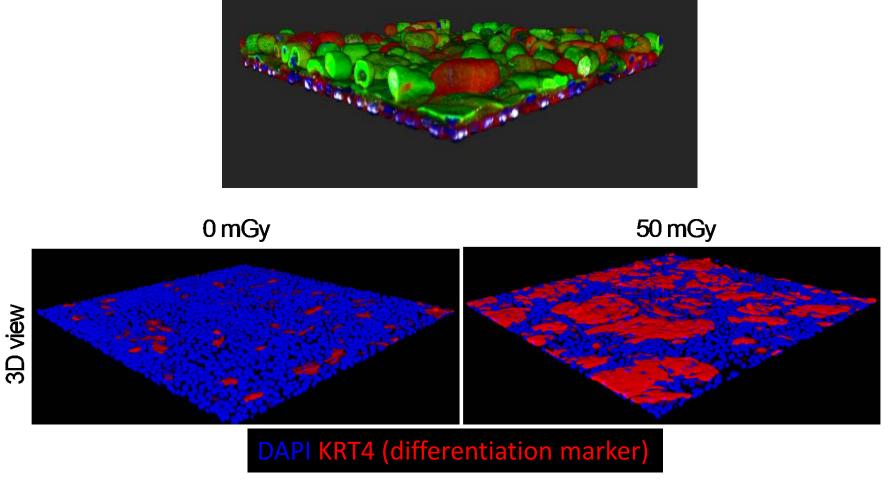
Genetic background?

Tissue type?

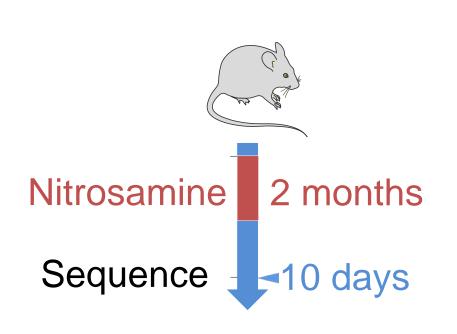
Somatic mutations in low dose research Challenging the selection hypothesis:

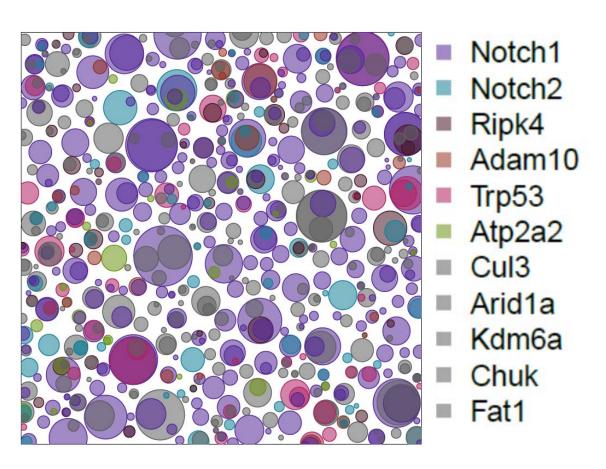
- Pre and post exposure sequencing of aged tissues
- Organotypic primary cultures with gene edited cells
 Live imaging, reporters, spatial omics.....
- Genetics of response: Crispr/Cas9 fitness screens +/- LDIR
- Mice: Lineage tracing, complex mutational landscapes

Organotypic Culture: esophagus



Replicating aged human esophagus in mice





Research Priorities

Basic biology of normal tissue responses is an urgent need

Engage a wider community in low dose research to diversify perspectives and technologies

Build collaborative groups linking radiation researchers with those new to the field

Diversity of models and tissues is an advantage

Ensure access to low dose irradiation facilities

Include cutting edge genomic and gene editing experts and data scientists









Emilie Abby
Irina Abnizova
Ujjwal Banerjee
Bartomeu Colom
Stefan Dentro
Jo Fowler

Albert Herms
Michael Hall
Charlotte King
Tom Metcalf
Swee Hoe Ong
Roshan Sood

Alumini Maria Alcolea Gabriel Piedrafita David Fernandez-Antoran Christopher Bryant



Christian Frezza



Bart Vanhaesebroeck Ben Hall



Moritz Gerstung