

The background of the slide is a dark-field microscopy image showing several clusters of cells. The cells are stained with a blue dye, likely DAPI, which highlights their nuclei. The image is overlaid with a light gray grid. Numerous horizontal bars of various colors (red, green, blue, yellow, magenta, cyan) are superimposed across the image, some appearing as thin lines and others as thicker bands, creating a layered, digital effect.

Babraham Institute
Annual Research Report
Signalling



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24-37

Signalling

The process of cell signalling consists of several interconnected mechanisms that allow cells to communicate, co-ordinate and respond rapidly to change. By examining these signalling mechanisms and their interactions we seek to understand the effects of signalling on cell growth, survival and behaviour.

Our current focus is to discover the role that signalling has in helping cells to respond and adapt to damage, illness, dietary changes and ageing by investigating:

- n How cells called neutrophils detect and respond to infections
- n How changes in diet affect metabolism and growth
- n The effect of signalling mechanisms on the rate of ageing
- n The role of autophagy in recycling cell components following damage or starvation

Group Leaders



Simon
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Oliver
Florey



Phill
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Nicholas
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Rahul
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Hayley
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Len
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Group members

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Sabine Suire

Senior research scientist:
Simon Rudge

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Research fellow

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Nicholas Ktistakis

Group members

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Current Aims

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selective autophagy
(mitochondrial autophagy)
aggrephagy (autophagy of
aggregates). In addition
tissue culture cells, we
on iPSC-derived neurons
understand how autophagy
neurodegeneration.

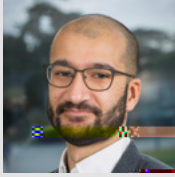
Progress in 2019 and 2020

We have modelled the process of
autophagy and mitophagy using
extensive collection of live imaging
and discovered a possible explanation
for why the process of mitophagy
involves the sequential translocation
of autophagy components to the

Publications

www.babraham.ac.uk/our-research





Rahul Samant

Group members

Postdoc research scientist:
Harvey Johnston

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Yasmeen Al-Mufti

Research assistant:
Estelle Wu

Why misfolded proteins accumulate with age

Cellular accumulation of misfolded proteins is a hallmark of ageing. In young cells, the proteostasis network limits toxicity by activating one or more systems for misfolded protein clearance. We focus on how these clearance systems are integrated within the network to maintain proteome health during youth, and how loss of this integration contributes to cellular senescence, another ageing hallmark with strong links to chronic inflammation and organismal frailty.

Current Aims

We use two evolutionarily distant cell types, budding yeast and primary human fibroblasts, to identify common, conserved lines of communication between different clearance systems of the proteostasis network, and investigate how these are re-wired during replicative ageing (yeast) and senescence (mammals). Our lab employs multi-disciplinary approaches such as super-resolution imaging, flow cytometry, and mass spectrometry-based proteomics to measure proteostasis capacity and



Hayley Sharpe

Group members

Postdoc research scientists:
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Cell signalling through tyrosine phosphatases

The reversible phosphorylation of protein tyrosine residues enables cells to dynamically respond to changes in their environment and is regulated by the antagonistic actions of kinases and phosphatases. We focus on the understudied phosphatases to understand how they signal, their roles in health and disease and how they are regulated, particularly by reactive oxygen species, which are implicated in the ageing process.

Current Aims

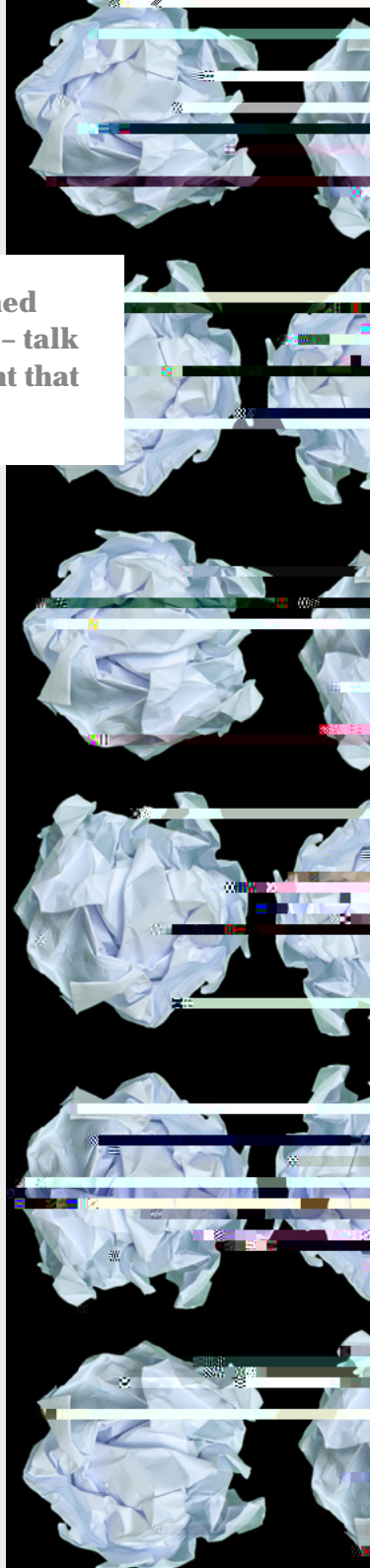
Our overarching aim is to understand mechanisms of tyrosine phosphatase signalling in order to understand their

fundamental functions but also to reveal new approaches to targeting them in disease, to overcome their undruggable reputation. Our current work is focused on a family of receptor tyrosine phosphatases that are present on the cell surface and form homophilic interactions at points of cell–cell contact. The receptor PTPRK is a tumour suppressor and had been suggested to regulate cell adhesion. To as dec diseasso tplaTPRI tritheince proachesecherec

The adhesive receptor protein tyrosine phosphatase PTPRK is expressed in epithelial cells at sites of cell–cell contact. We have identified key substrates linked to cell adhesion (left). Transmission electron microscopy images reveal that deleting PTPRK from mammary epithelial cells leads to disrupted cell junctions and adhesions, as well as decreased cell height, reminiscent of an epithelial to mesenchymal transition (right).



Setting up a new group is exciting and daunting. Two group leaders who joined the Signalling programme in 2019 – Dr Hayley Sharpe and Dr Rahul Samant – talk about their research and the supportive, collaborative and open environment that they say marks out the Institute.



This quality control machinery declines as we age. As a result, most age-related diseases, including Alzheimer's disease, Parkinson's



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