### HIGHLIGHTS

#### DEVELOPMENT

## Hedgehog proliferation

Mutations that activate the Hedgehog (Hh) signalling pathway have been linked to tumour formation, but it's not been clear how. The discovery of a direct link between Hh signalling and key regulators of the cell cycle might now provide the answer.

Wei Du and colleagues were studying eye development in *Drosophila melanogaster*. The expression pattern of Hh

during this

process, just posterior to cells entering S phase, indicated that reception of the Hh signal might be needed for entry to S phase. To test this, the authors looked at what would happen if Hh signalling was blocked during eye development. They found that second mitotic wave cells with mutated *smoothened* (*smo*), a gene that is required for Hh signalling, do not enter S phase. By contrast, overexpression of Cubitus interruptus (Ci) — the transcription factor that mediates Hh signalling — drove G1-arrested cells to enter S phase.

One protein that promotes S phase is Cyclin D. During eye

development, the highest expression of Cyclin D overlaps with that of

Ci — so could Ci promote the expression of Cyclin D? Support for this idea came from the observation that levels of Cyclin D are reduced in *smo*-mutant clones, and also that overexpression of Ci induces high levels of Cyclin D

As well as promoting entry into S phase, Cyclin D induces cell growth. Du and co-workers therefore wondered whether Hh might also regulate growth, so they studied the effects of overexpressing either Ci or Patched (Ptch; an inhibitor of Hh signalling) in clones of undifferentiated wing-disc cells. Whereas Ptch overexpression clones were considerably smaller than controls, Ci overexpression clones were much larger, which indicates that Hh signalling not only promotes S phase, but that it also regulates cell growth.

Cyclin E also promotes S phase, and reduced or increased levels of this protein could be detected with loss of Smo or overexpression of Ci, respectively. The authors then looked at how Hh signalling might induce the transcription of Cyclin E. They identified several sequences in the Cyclin E promoter with homology to the consensus Cibinding site, and used chromatin immunoprecipitation to show that

#### EPITHELIAL-MESENCHYMAL TRANSITION

# A deadly combination

In some cell lines, an epithelial– mesenchymal transition (EMT) arises as the result of a joint effort between Hras and transforming growth factor- $\beta$  (Tgf- $\beta$ ). How relevant this is to the multistage nature of *in vivo* tumour progression, though, is a burning question.

So, Allan Balmain's group studied whether changes in the levels of Hras and Tgf- $\beta$  affect tumour progression, using a series of wellcharacterized tumour cell lines that arise from initiated cells that carry activating mutations in the *Hras1* gene. And, as they now report in *Nature Cell Biology*, Smad2 (a downstream target of Tgf- $\beta$  signalling) and Hras surpass discrete thresholds during progression from early-stage papillomas, through squamous carcinomas, to late-stage undifferentiated spindle-cell tumours.

First, the authors studied the molecular changes that occur when squamous carcinomas are converted into spindle-cell tumours. Tgf-β-mediated transcriptional activity was very high in the spindle cells, and phosphorylated Smad2 accumulated in the nucleus, which indicated that the Tgf-β pathway was activated in these cells. Furthermore, in primary material from spindle-cell tumours, but not from differentiated tumours or squamous carcinomas, Smad2 was phosphorylated and predominantly localized in the cytoplasm.

mRNA and protein.

Although Smad2 alone induced changes in the migration of squamous carcinoma cells, only in the presence of increased levels of mutated Hras did changes in cell shape and the expression of genes such as  $\alpha$ -smoothmuscle actin (a mesenchymal marker) occur, resulting in EMT.

The authors then investigated whether, once this stage has been reached, Tgf- $\beta$  signalling by Smad2 is still necessary for tumour progression. Expression of a dominant-negative form of Smad2 showed that this is indeed the case; spindle cells that expressed this construct reverted to a more epithelial phenotype and took on many features of epithelial gene expression. Notably, surface expression of  $\alpha\nu\beta3$  integrin was lost, and this was coincident with the loss of collagen-matrix invasion. *In vivo*, this correlated with an

inability to form tumours. By contrast, parental spindle cells or spindle cells that express a dominant-active form of Smad2 formed tumours, and those formed by dominant-active Smad2 were particularly invasive. Expression of dominant-active Smad2 also promoted extravasation into the target tissue, and a subsequent increase in lung metastases.

As the ability of a tumour to metastasize is the main determinant of whether or not patients with cancer die of their disease, these findings that different thresholds of Hras and Tgf- $\beta$  activity — intermediate levels of Smad2 co-operating with Hras to induce EMT and invasiveness, and even higher levels of Smad2 being required for metastasis — are crucial for metastasis offer the opportunity for the design of small-molecule inhibitors to prevent the spread of tumours.

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#### References and links

**ORIGINAL RESEARCH PAPER** Oft, M., Akhurst, R. J. & Balmain, A. Elevated levels of activated Smad2 and Hras control epithelial-mesenchymal transformation, turnour cell extravasation and metastasis. *Nature Cell Biol.* **4**, 487–494 (2002) **FURTHER READING** Frame, S. & Balmain, A. Integration of positive and negative growth signals during ras pathway activation *in vivo. Curr. Opin. Genet. Dev.* **10**, 106–113 (2000) | Akhurst, R. J. & Derynck, R. TGF- $\beta$  signaling in cancer – a double-edged sword. *Trends Cell Biol.* **11**, S44–S51 (2001) **WEB SITE** 

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