

Putting the Wnt up colon cancer

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If there is one cancer signalling pathway that *Gut* readers should be comfortable with, it is the Wnt pathway. Central to this pathway is the adenomatous polyposis coli (APC) gene responsible for familial adenomatous polyposis. Identification of the APC gene in 1991 heralded the start of an ongoing period of exponential growth in our understanding of the molecular basis of cancer, particularly of the large bowel. Clinicians can perhaps be forgiven for finding the ever-expanding list of genes and interacting pathways with their inaccessible names and acronyms daunting. For many, the paper in this issue by Elvira Bakker and coworkers at the Netherlands Cancer Institute revealing the detailed role of *RSPO3* in colorectal carcinogenesis may only serve to widen the expanding clinician–researcher divide. However, those brave enough to dive deeper into this paper will be rewarded with an insight into the current state of the art of molecular research into colorectal cancer with an intriguing mix of confirmation of what we thought we knew and findings that challenge previous assumptions.

R-spondin 3 (*RSPO3*) is a secreted protein that acts to potentiate Wnt signalling. *RSPOs* first attracted attention as being specific ligands of the LGR5 (leucine-rich repeat-containing G-protein-coupled receptor 5) receptor. LGR5 is the marker for intestinal stem cells first described by Hans Clevers. LGR5 was itself identified in a screen for Wnt pathway targets in the intestine. So in a sense we have come full circle investigating a ligand, *RSPO3*, which impacts on the quintessential colon cancer pathway, the Wnt pathway, and itself discovered through a detailed investigation of Wnt signalling. But why investigate a ligand that is seemingly once removed from the action? For biologists, the mountaineers' adage 'because it's there' might suffice, but *Gut* readers will ask how is this likely to impact clinical practice?

RSPO-activating translocation mutations have been recently found in around 10%

of human colorectal cancers¹ and these tumours respond to *RSPO*-neutralising antibodies.² *RSPO3* mutations occur mutually exclusive with other classical Wnt-activating mutations (eg, in *APC* or *CTNNB1* (the gene for β -catenin)), suggesting that *RSPO3* mutations can substitute for them with a similar net effect on Wnt signalling and oncogenesis. Furthermore, *RSPO* may potentially represent a handle by which to pharmacologically target the notoriously 'undruggable' Wnt pathway.³ The problem with the Wnt pathway itself is that it is corrupted by pathway-activating mutations downstream of the Wnt ligands and receptors. Most successful molecularly targeted therapies to date have been aimed at ligands or their receptors. The mutated intracellular signal transduction elements of the pathway such as APC and β -catenin are tricky therapeutic targets. Recent work in a broader array of tumour types suggests that an *RSPO*-based therapy may have a bright future.⁴

Further tantalising insights come from the analysis of serrated polyps. The most aggressive colorectal cancers develop from serrated polyps via the serrated pathway⁵ and follow a different molecular chain of events than the classical Vogelstein adenoma-to-carcinoma molecular model, with no classical Wnt pathway mutations. Late recognition of serrated polyps and familial forms of serrated polyposis has meant that molecular understanding of the serrated pathway lags behind that of classical adenomas. *RSPO3*-activating mutations have recently been found to occur very frequently (30%) in serrated polyps.⁶ Could this explain their aggressive nature?

What have the Dutch researchers done? Using advanced genetically modified mouse models, they have ensured that mice, after receiving tamoxifen, express large amounts of *RSPO3* specifically in stem cells thereby mimicking the effects of the human *RSPO3* mutations. Within only 2 months, these mice develop dramatic changes in their intestine with widespread hyperplasia and grossly disrupted crypt-villus architecture. They do not use the term 'serrated'. However, the morphological characteristics of serrated architecture have not been clearly defined in animal models⁷ and agreement among pathologists is poor even in human lesions.⁸ The mice also develop multiple adenomas and even invasive carcinomas, something that

is rarely seen in mice where only a single gene is altered. Intriguingly, the expected high levels of Wnt pathway activity were absent as judged by nuclear accumulation of β -catenin. A further analysis using RNA sequencing confirmed only modest increases in Wnt signalling-related genes. While perhaps unexpected, this would fit well with the previous evidence published in *Gut* that there is a subtle relationship between levels of Wnt signalling and polyp development.⁹ More Wnt signalling does not translate directly into more polyps and cancer. The lower levels of Wnt signalling induced by *RSPO3*-activating translocations may be 'just right' in this respect. Does this mean that therapeutic efforts to lower Wnt signalling could have unexpected adverse effects?

The paper goes on to use lineage-tracing methods to assess which cells turn into the polyps and cancers they find. By expressing *RSPO3* specifically in stem cells, one might expect expansion of the stem cell pool and that polyps and cancers also arise from these cells. Surprisingly, there was little expansion of the stem cell compartment in contrast to the traditional *APC*-mutant models.⁹ *RSPO3* seems to act predominantly on non-stem cells in the intestine via an alternative more widely expressed receptor, LGR4. The final surprise is the effect of adding in mutant *KRAS* on top of increased *RSPO3*. In human tumours, mutations in *RSPO3* are always found in conjunction with *KRAS* or *BRAF* mutations. The *KRAS* mutations have been added on top of multiple other single colorectal cancer gene manipulations in mice (*APC*, *TGFBRII*, *PTEN* and *CDKN2A*) and consistently lead to a more aggressive phenotype.¹⁰ It would therefore be expected that the addition of a *KRAS* mutation would increase the number of invasive tumours. However, this was not seen. What is apparent is that *RSPO3* and *KRAS* show additive effects as far as the hyperplastic phenotype is concerned.

In summary, this paper confirms *RSPO* as an important driver mutation in colorectal cancer and as such it represents an attractive target for new therapies.

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Commentary

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