O TUMOUR METABOLISM

Translating the undruggable target

Clear cell renal cell carcinoma (ccRCC) is largely attributable to inactivating mutations in the von Hippel-Lindau tumour suppressor pVHL, which drive accumulation and activation of the transcription factor hypoxia-inducible factor 2a (HIF2a). HIF2a, like many transcription factors, was considered undruggable until the discovery of a large cavity within the HIF2a PAS-B domain capable of binding to small molecules. This breakthrough led to the identification of a series of selective HIF2a antagonists. Three studies have now evaluated the anti-tumour activity of two of these compounds, PT2399 and the closely related analogue PT2385, in preclinical models of pVHL-mutant ccRCC.

What is the mechanism of HIF2a antagonists? Using a VHL^{-/-} ccRCC cell line, 786-O, both Cho et al. and Wallace et al. showed that treatment with PT2399 or PT2385 could disrupt the heterodimerization between HIF2a and its transcriptional partner HIF1 β (also known as ARNT), thereby preventing HIF2a binding to DNA. Complementing these findings, Chen et al. demonstrated the same result with the treatment of patient-derived xenograft (PDX) ccRCC tumours. All three studies used quantitative PCR to show that both PT2399 and PT2385 suppressed the expression of a number of HIF2a-specific target genes, such as vascular endothelial growth factor A (VEGFA), cyclin D1 (CCND1) and erythropoietin (EPO), but importantly, not those of HIF1 α , such as phosphoglycerate kinase 1 (PGK1). What is the functional outcome

of specific HIF2a transcriptional

inhibition in vivo? Cho et al. and

mouse models, saw that extended

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Wallace et al., using 786-O xenograft

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extended treatment with PT2399 or PT2385 caused tumour stasis and regression

Do these HIF2a antagonists represent an improvement over current approved first-line angiogenesis inhibitors for advanced ccRCC? Chen et al. and Wallace et al. compared the activity of PT2399 or PT2385 against sunitinib. Both HIF2a antagonists were better tolerated, and showed greater efficacy

caused tumour stasis and regression coinciding with decreased cell proliferation and decreased angiogenesis. PT2399 and PT2385 also substantially reduced tumour growth and decreased tumour vascular area in *VHL*^{-/-} ccRCC PDXs. Together these studies validate HIF2a as a target driving pVHL-defective ccRCC.

Could HIF2a antagonists be used as therapy for all VHL^{-/-} ccRCCs? Surprisingly, Cho et al. and Chen et al. both observed that some VHL-mutant ccRCC cell lines and tumours were resistant to PT2399. These two groups came to the same conclusion that this resistance is probably the result of differential dependence on HIF2a, with both sensitive VHL-deficient ccRCC cell lines and PDX tumours having considerably higher HIF2a levels. However, the responsiveness of VHL^{-/-} ccRCCs may not be exclusively ascribed to HIF2a expression; Cho et al. identified two resistant ccRCC cell lines with high HIF2a harbouring a p53 R248W mutation, suggesting that p53 status could also be a factor. In addition, Chen et al. found that even sensitive PDX tumours could gain resistance to PT2399 after prolonged exposure, with one tumour harbouring a PT2399 binding site mutation in HIF2α and another a second site suppressor mutation in HIF1 β . This finding suggests that the design of complementary inhibitors may be required.

than sunitinib in reducing PDX tumour growth. Chen et al. even found that PT2399 inhibited tumour growth in some sunitinib-resistant PDX tumours.

The potential of PT2399 and PT2385 for clinical use is already evident, as Chen et al. demonstrated that a patient with extensively pretreated metastatic ccRCC, subsequently treated with PT2385 in a phase I trial, had a sensitive PDX tumour and exhibited progression-free survival for more than 11 months. However, these studies also underscore the need for predictive and pharmacodynamic biomarker-driven clinical trials, as patient responses are likely to depend, at least in part, on HIF2a levels.

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ORIGINAL ARTICLES Chen, W. et al. Targeting renal cell carcinoma with a HIF-2 antagonist. Nature http://dx.doi.org/10.1038/nature19796 (2016) | Cho, H. et al. On-target efficacy of a HIF2a antagonist in preclinical kidney cancer models. Nature http://dx.doi.org/10.1038/nature19795 (2016) | Wallace, E. M. et al. A small-molecule antagonist of HIF2α is efficacious in preclinical models of renal cell carcinoma. Cancer Res. 76, 5491-5500 (2016)