

Efficacy of the Porcupine Inhibitor RXC004 in genetically-defined tumour types

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Introduction

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Signalling through the Wnt pathway is highly regulated at the level of ligand (Wnt), receptor (Fzd/LRP) and downstream components (e.g. destruction complex – APC/Axin/GSK3 β). Post-translational modification of Wnt via porcupine (PORCN), a membrane bound O-acyltransferase is essential for secretion of active Wnt¹. Activity of RNF43/ZNF3 (E3-ubiquitin ligases) results in ubiquitination and membrane clearance of Fzd, whilst RNF43/ZNF3 levels are kept in check via LGR and secreted RPSO ligands² (Fig. 1).

The potent and selective porcupine (PORCN) inhibitor RXC004 is being investigated in a Phase 1 clinical trial (NCT03447470)³, and has the potential to treat tumours dependent on Wnt-ligand. Upstream Wnt pathway aberrations, including RNF43/ZNF3 mutations and RPSO-fusions, result in high levels of surface Fzd receptors and increased Wnt-ligand dependent signalling⁴ (Fig. 1). These aberrations are implicated in pancreatic, gastric and colorectal cancer (CRC).

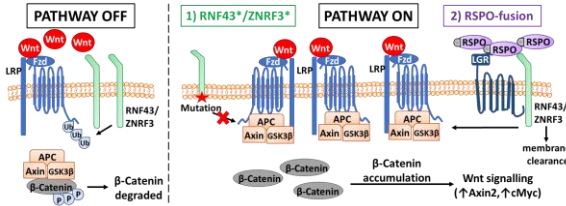


Figure 1: Upstream alterations trigger aberrant Wnt ligand-dependent signalling RNF43/ZNF3 keep surface Fzd in check, allowing the destruction complex to phosphorylate and degrade β -catenin - **Wnt pathway 'OFF'**. Loss-of-function (LOF) RNF43/ZNF3 mutations (1), or high RPSO expression through gene fusion (2), ultimately leads to accumulation of β -catenin - **Wnt pathway 'ON'**.

Results

In vitro pathway inhibition and anti-proliferative effects of RXC004 in genetically-defined tumour cell lines

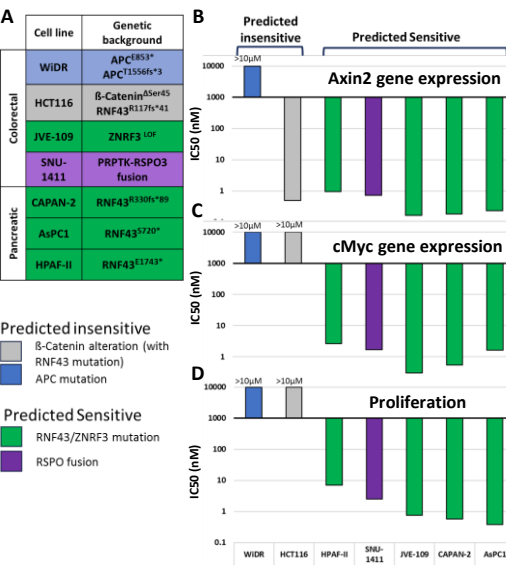


Figure 2: RXC004 potency in genetically-defined pancreatic and CRC cell lines (A) A dose response of RXC004 was evaluated across a panel of genetically-defined tumour cell lines. (B, C) Cells were treated with RXC004 for 72h. RNA was isolated and analysed using RT-qPCR to assess mRNA expression of the downstream markers of target engagement for the Wnt pathway, Axin2 (B) and cMyc (C). (D) Cells were treated with RXC004 for 5 days in 2D or 3D format, proliferation was measured using an ATP-lite assay. N \geq 3 throughout.

RXC004 efficacy and PD in genetically-defined xenografts

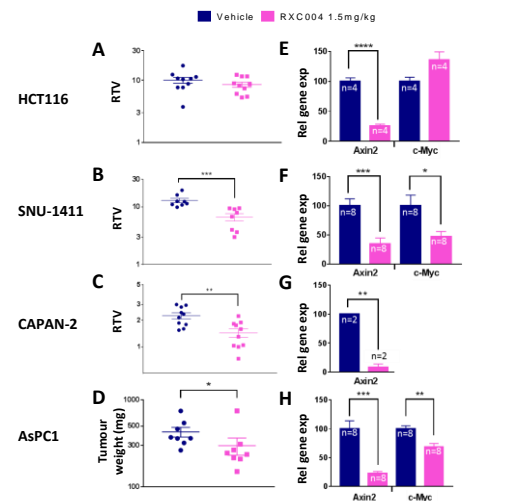


Figure 4: RXC004 efficacy and PD in human xenograft models Relative tumour volumes (RTV) (A-C), end of study tumour weights (D: mg) and end of study relative gene expression of Axin2 and cMyc (E-H). HCT116 (A; 3x10⁶ cells; athymic nude mice), SNU-1411 (B; 1x10⁷ cells; NOC-SCID mice), CAPAN-2 (C; 3x10⁶ cells; SCID-Beige mice) and AsPC1 (D; 3x10⁶ cells; NOD-SCID mice) were implanted subcutaneously. Treatment was initiated once tumour volumes reached \sim 100-150mm³. Tumours per group: 10 (A, C) and 8 (B, D). Dosing was 1.5mg/kg BID throughout (A, C) or BID for 7 or 13 days then QD for the remainder of the study (B, D). Tumour RNA was isolated for RT-qPCR analysis of the downstream target engagement biomarkers for the Wnt pathway, Axin2 and cMyc at 8h (E, G) or 12h (E, F, H) post final dose. Mann-Whitney U (A-D) or unpaired t-test (E-H) p values.

RXC004 arrests at G1/S and G2/M cell cycle checkpoints

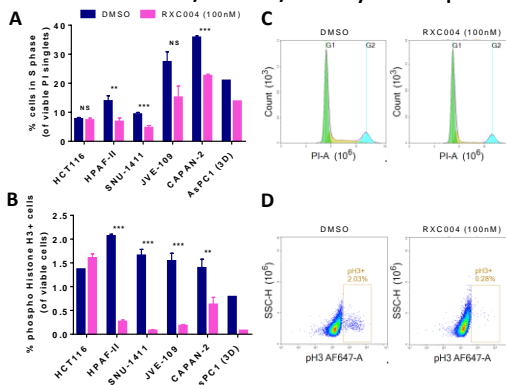
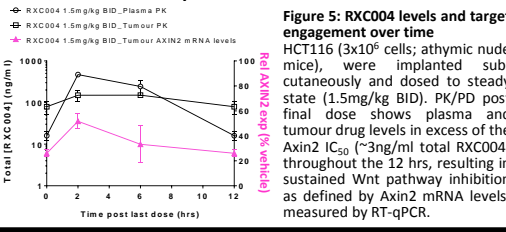


Figure 3: Effects of RXC004 treatment on the cell cycle Cells were treated with RXC004 (100nM) or vehicle (0.1% DMSO) for 72h. Cell cycle profiles were determined using propidium iodide (A, C) and phospho^{Ser10}-Histone H3 (mitotic marker; B, D) staining by flow cytometry. Data are N \geq 3 except for HCT116 and AsPC1 (N=1). Representative flow plots for cell cycle (C) and phospho-Histone H3 (D) in HPAF-II cells treated as indicated. T-test p values.

PK/PD relationship of RXC004 in the HCT116 model



Data represent Mean \pm SEM. *p<0.05; **p<0.01; ***p<0.001; ****p<0.0001

Conclusions

- Tumour cells carrying RNF43 mutations or RPSO fusions are sensitive to RXC004 treatment both *in vitro* and *in vivo*.
- RXC004 monotherapy could benefit patients with tumours bearing RNF43 mutations or RPSO fusions.
- Data supports a genetically-defined patient selection strategy for ongoing RXC004 clinical studies.

References

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