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1	Frizzled-7 is required for Wnt signaling in gastric tumors with and without Apc mutations.
2	
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 45

46 Abstract

A subset of gastric cancer (GC) patients have mutations in genes that participate in or regulate 47 48 What signaling at the level of ligand (What) receptor (Fzd) binding. Moreover, increased Fzd expression is associated with poor clinical outcome. Despite these findings, there are no in vivo 49 studies investigating the potential of targeting Wnt receptors for treating GC, and the specific Wnt 50 51 receptor transmitting oncogenic Wnt signaling in GC is unknown. Here we use inhibitors of 52 Wnt/Fzd (OMP-18R5/Vantictumab) and conditional gene deletion to test the therapeutic potential 53 of targeting Wnt signaling in preclinical models of intestinal-type gastric cancer and ex vivo 54 organoid cultures. Pharmacological targeting of Fzd inhibited the growth of gastric adenomas in 55 vivo. We identified Fzd7 to be the predominant Wnt receptor responsible for transmitting Wnt signaling in human gastric cancer cells and mouse models of GC, whereby Fzd7-deficient cells 56 57 were retained in gastric adenomas but were unable to respond to Wnt signals and consequently failed to proliferate. Genetic deletion of Fzd7 or treatment with Vantictumab was sufficient to inhibit 58 59 the growth of gastric adenomas with or without mutations to Apc. Vantictumab is currently in phase Ib clinical trials for advanced pancreatic, lung, and breast cancer. Our data extend the 60 scope of patients that may benefit from this therapeutic approach as we demonstrate that this 61 62 drug will be effective in treating gastric cancer patients regardless of Apc mutation status.

63

64 Statement of significance

The Wnt receptor Fzd7 plays an essential role in gastric tumorigenesis irrespective of Apc mutation status therefore targeting Wnt/Fzd7 may be of therapeutic benefit to gastric cancer patients.

68

69 Introduction

70 Gastric cancer (GC) is a common malignancy, ranking in the top 4 of global cancer incidence [1]. 71 Often due to advanced stage diagnosis, gastric cancer patients have a very poor 5-year survival 72 rate [1]. This highlights a desperate need for novel clinical treatments as there are very few 73 approved targeted therapies for GC [2, 3]. Gastric cancer is divided histologically into two groups; intestinal-type and diffuse-type, with intestinal-type being more prevalent. Members of the cell-74 75 surface Frizzled (Fzd) receptor family are deregulated or overexpressed in several cancer types, 76 including GC [4]. Whits are lipid-modified glycoproteins that initiate signal transduction by binding 77 to Fzd via a palmitate group, which is appended by the palmitovltransferase Porcupine (PORCN) [5, 6]. White also bind cell surface co-receptors, such as Lrp5/6, forming a ternary complex [7]. 78 Formation of the Wnt-receptor complex leads to inhibition of a multiprotein 'destruction complex' 79 comprised of Axin, glycogen synthase kinase-3 (GSK3), calcium kinase-1 (CK1) and 80 81 adenomatous polyposis coli (APC), which targets β -catenin for proteosomal degradation. Newly synthesised cytoplasmic β -catenin can now escape degradation, accumulate and translocate to 82 83 the nucleus, where it forms a transcriptionally active complex with T-cell factor (TCF)/lymphoid enhancing factor (LEF) family of transcription factors to induce Wnt target gene transcription [8]. 84 85 However, deregulated Wnt signaling can initiate cell transformation and subsequent 86 carcinogenesis [8].

87

88 Furthermore, several Wht/Fzd antagonists [9] are epigenetically silenced through promoter hypermethylation, including DKK3 (67.6% of gastric tumors [10]), sFRP1 (91%), sFRP2 (96%), sFRP5 89 (65%) [11], whilst others such as the E3 ligase RNF43, which regulates Fzd turnover on the cell 90 91 surface [12], are mutated in 54% and 4.8% of microsatellite instable (MSI) and microsatellite 92 stable (MSS) gastric tumors, respectively [13]. Exogenous re-introduction of sFRP or DKK can significantly reduce gastric tumor burden in APC or β -catenin-mutant gastric cancer cells by 93 attenuating Wnt signaling [11, 14]. Critically, this provides proof-of-principle that modulation of 94 ligand/receptor signaling components can further regulate Wnt signaling irrespective of 95 96 downstream mutations that constitutively activate the pathway, which has been reported in colorectal cancer cells [15-18]. Together, these data strongly implicate a role for Wnt/Fzd in GC 97 98 which could be exploited for targeted therapy.

99

We recently demonstrated that Frizzled-7 (Fzd7) regulates stem cell function in the gastric and intestinal epithelium [19, 20]. In addition, *FZD7* is abundantly expressed in human gastric cancer tissue [21-23], which is also associated with poor patient outcome [24]. Despite compelling

evidence implicating Fzd receptors in GC, there has been no formal investigation of the 103 104 therapeutic benefit of targeting Fzd receptors in GC in vivo. These types of in vivo studies are 105 crucial to fully understand the potential of novel therapeutic strategies due to the complex cellular and molecular interactions of a tumor, which can directly inform clinical trials and cannot be 106 107 replicated in vitro. Our results demonstrate that Fzd receptors, specifically Fzd7, are rate-limiting for the growth of gastric adenomas with or without Apc mutations in vivo. These findings have 108 significant clinical utility as targeted Fzd therapeutics (OMP-18R5/Vantictumab), currently being 109 tested in other solid cancer types (http://www.oncomed.com/Pipeline), can now be extended to 110 111 GC patients with and without APC mutations.

112

113 Materials and Methods

114 **Mice**

The *Tff1Cre^{ERT2}*[25], *Fzd7^{tl/fl}*[20], *Apc⁵⁸⁰ (Apc^{fl/fl})*[26], *c-Myc^{fl/fl}*[27], *Rosa26LacZ*[28] and *gp130^{F/F}* [29] are previously described. Mice were interbred to generate compound mice with appropriate alleles on an inbred C57Bl/6 genetic background. Mice were co-housed using appropriate littermates as controls. All animal experiments were approved by the Animal Ethics Committee, Office for Research Ethics and Integrity, University of Melbourne.

120

121 **Treatments**

In vivo Cre induction was performed in 8-10 week old mice with a single daily intraperitoneal (ip) injection of 2mg of tamoxifen/mouse/day over four consecutive days. *gp130^{F/F}* mice aged 8-9 weeks were injected ip with 20mg/kg of OMP-18R5 (OncoMed) or vehicle control (2.5%DMSO+IgG) twice weekly over the course of 30 days at which point animals were sacrificed and tissues harvested.

127

128 **Tumor xenografts**

A total of 4x10⁶ cells in 100µl of PBS were injected subcutaneously into the hind flank of 6-8 week
old nude mice (nu(ncr)-foxn1 nu/nu). 7 mice were used for each cohort which were treated with
20mg/kg OMP-18R5 or vehicle control (2.5%DMSO+lgG) once tumors were palpable, five days
following injection of cells. Xenografts were measured with calipers twice a week to monitor tumor
growth.

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- 136

137 Tissue collection and histological analysis

Mouse stomachs were isolated, flushed with PBS, fixed overnight at 4°C in 10% neutral buffered formalin (NBF) and processed for immunohistochemistry and immunofluorescence as we previously described [20, 30, 31], with antibodies used on Table S2.

141

142 Isolation and culture of normal and tumor organoids

Organoids were cultured from mouse stomachs as previously described [31]. Adenomas from 143 gp130^{F/F} mice were isolated from the stomach, washed in PBS, roughly minced and incubated in 144 digestion buffer (Dispase I (125µg/ml), Collagenase IV (75U/ml) and DMEM+2.5% FCS) at 37°C 145 until epithelial fragments dissociate from tumor bulk. Dissociated cells were passed through a 146 70µM cell strainer, counted, centrifuged and resuspended in Matrigel. In vitro Cre recombinase 147 was activated by treating gastric organoid cultures with 100nM 4-hydroxytamoxifen (4-OHT) as 148 previously described [31]. R-Spondin and Wnt conditioned medium were withdrawn from 149 *Tff1Cre⁺;Apc^{fl/fl}* organoid cultures following 4-OHT treatment. Differential interference contrast 150 (DIC) images were captured as Z-sections and final image generated as previously described 151 152 [20, 32].

153

154 **RNA extraction and analysis**

Gastric glands were homogenized in TRizol and total RNA purified, DNAse treated, quantified and subjected to quantitative reverse transcriptase PCR (qRT-PCR). qRT-PCR and calculating gene expression levels relative to the house-keeping gene 18S (2-ΔΔCT) were performed as previously described [16].

159

160 MTT assay

Following treatment, gastric organoids were mechanically dissociated, washed with ADF, resuspended in fresh Matrigel and seeded in a flat bottom 96 well tissue culture plate for enumeration using the MTT assay performed exactly as we previously described [19, 20].

164

165 Cell culture and transfection

Human gastric cancer cell lines (MKN28, MKN74, MKN7, MKN1, AGS and MKN45) were
maintained in RPMI 1640 medium (Invitrogen) supplemented with 10% fetal calf serum (FCS)
(Invitrogen) and 1% penicillin/streptomycin (Invitrogen) and L-Glutamine (Invitrogen) and were
not taken past passage 15 for experimental use. All cells were tested for Mycoplasma,
authenticated and cultured at 37°C in 5% CO₂. Gastric cancer cells were transfected with Short-

hairpin RNA (shRNA) and expression constructs designed to knockdown and stably express
 FZD7 respectively, as previously described [16, 33] or MSCV-MYC from Addgene (18119).

173

174 Soft agar colony assay

175 Cells were cultured in 60mm tissue culture dishes until 50% confluency and transfected with 5µg of plasmid DNA using Lipofectamine LTX (Invitrogen) following manufacturer's instructions. After 176 48hrs incubation, cells were washed in PBS, detached using trypsin, resuspended 177 RPMI+10%FCS, counted and mixed with pre-warmed 1% agar/RPMI culture medium to a final 178 179 concentration of 500 cells/well of a 6-well plate. Once agar/cell suspensions solidified, cultures were overlaid with RPMI+10% FCS culture medium and incubated at 37°C in 5% CO₂ for 14 days. 180 181 For Wnt inhibition experiments, cells were treated with OMP-18R5 (10µg/ml), IWP-2 (10µM) [34] 182 or vehicle control (2.5%DMSO+IgG) 3 days after plating. Treatments were removed and replaced 183 every 4 days over the 2 weeks. Cells were fixed in 4%PFA and stained with crystal violet and 184 colonies consisting of \geq 50 cells scored and imaged.

185

186 Genomic recombination PCR

187 Conventional PCR to detect the *Fzd7* and *Apc* mutant alleles following recombination in genomic

- 188 DNA extracted from compound transgenic mice was performed as previously described [20, 35].
- 189 See also supplementary experimental procedures.
- 190

191 Luciferase assay

192 Cells were cultured in 24-well tissue culture plates until 50% confluency and transfected with a total of 1µg plasmid DNA/well (500ng of SuperTOPflash or SuperFOPflash TCF reporter plasmids 193 194 expressing firefly luciferase [36], plus 500ng of either "control" or "treatment" DNA, plus 2ng of 195 renilla luciferase plasmid to normalize transfection efficiency). Cells were transfected using Lipofectamine LTX with Plus reagent (Invitrogen) according to manufacturer's instructions. Cells 196 were harvested 48hr later and analysed using the dual luciferase reporter assay system 197 198 (Promega). Ratio of luciferase/renilla reporter activity was calculated and results expressed relative to control cultures. 199

200

201 Analysis of gastric adenocarcinoma genomic dataset

Analysis of somatic mutations and copy number alterations (CNA) for a panel of 21 Wnt pathway

203 genes was performed on the TCGA stomach adenocarcinoma dataset [37] using the cBioPortal

platform [38]. Only samples with sequencing and CNA data were assessed across all molecular
subtypes, n = 287.

206

207 Statistical analysis

Data are expressed as mean \pm SEM, where mean represents number of mice (\geq 3 per genotype) or number of independent experiments (\geq 3). Statistical tests used are Mann-Whitney with Prism7 (GraphPad software) where *P* values of \leq 0.05 were considered significant. Heatmap generated in R version 3.0.2 using the *heatmap* function in the stats base package. Raw Ct values were transformed to delta Ct values using β 2M as housekeeping gene.

213

214 Results

Gastric cancer cells require cell intrinsic Wnt signaling for growth

Gastric cancer, like many malignancies, is genetically heterogeneous, which complicates 216 217 identifying non-redundant signaling pathways suitable for targeted therapy. To investigate the expression of Fzd receptors, which transmit oncogenic Wnt signals, we performed gRT-PCR for 218 219 all 10 mammalian Fzd genes on a panel of human GC cell lines. Several Fzd receptors were 220 abundantly expressed, including FZD7 (Figs. 1A, B and Supplementary Fig. S1A-D), suggesting 221 these might be attractive therapeutic targets. Although the pan-Fzd antibody, OMP-18R5 222 (Vanticumab) has shown efficacy in several solid cancer types [39], its therapeutic potential for GC has not been explored. MKN28 (APC mutant), MKN74 (APC mutant) and MKN45 (APC wild-223 224 type) GC cells treated with OMP-18R5 formed significantly fewer anchorage-independent colonies compared to vehicle control treated cells (Figs. 1C, D and Supplementary Fig. S1E). Of 225 note, MKN28 and MKN45 cells grown as conventional 2D monolayers do not show growth 226 inhibition following OMP-18R5 treatment (Supplementary Fig.1F and G), which highlights the 227 228 importance of testing drug efficacy in conditions that better mimic tumor biology. This suggests that cell intrinsic Wnt ligands are required for the 3D-growth of GC cells, which we confirmed by 229 230 treatment with IWP-2, which prevents Wnt secretion [40] (Figs. 1C and D). TOPflash assays and gRT-PCR demonstrate that either IWP-2 or OMP-18R5 treatment inhibit Wnt signaling in GC cells 231 232 (Figs. 1E-H). These data demonstrate cell intrinsic secretion of Wnt ligands and Fzd receptor availability are required for the sustained growth of GC cells. To determine whether Fzd regulates 233 the growth of established gastric tumors, MKN28 and MKN45 cells were subcutaneously injected 234 235 into the hind flanks of nude mice and allowed to develop into palpable gastric tumors. Compared to vehicle control treated gastric tumor xenografts, OMP-18R5 treated mice had significantly 236 237 smaller gastric tumors (Supplementary Figs. 2A-D), which demonstrates Fzd inhibition is sufficient to block the initiation (Figs 1C and D) and progression (Supplementary Figs. 2A-D) of human
gastric cancer cells.

240

241 Inhibiting Fzd receptors limits gastric tumorigenesis in vivo

242 We next utilised the well-characterised gp130^{F/F} mouse-model of intestinal-type gastric tumorigenesis [29, 41], which develop prominent antral lesions with adenomatous hyperplasia to 243 explore the relative expression of Fzd receptors. Compared to normal gastric epithelium, 244 upregulation of several Fzds was observed in *gp130^{F/F}* gastric adenomas (Figs. 2A-C), supporting 245 246 expression levels observed in human GC cells (Figs. 1A, B, and Supplementary Figs. S1A-D). Expression of Wnt ligands and target genes are also increased in *qp130^{F/F}* gastric adenomas 247 compared to non-adenoma gastric epithelium (Figs. 2A-C, Table S1). To determine if Fzd 248 inhibition could also reduce the growth of antral gastric adenomas in vivo, we treated 8-week-old 249 gp130^{F/F} mice, which at this age have small antral gastric adenomas (Supplementary Fig. S2E), 250 with OMP-18R5 twice a week for 30 days, following published protocols (Supplementary Fig. S3A) 251 [39]. Gastric adenomas were significantly smaller and fewer in OMP-18R5-treated *gp130^{F/F}* mice 252 253 compared to vehicle control treated mice (Figs. 2D-F), which was associated with a significant 254 reduction in the expression of Wnt target genes and cell proliferation (PCNA IHC) (Figs. 2G-J). 255 As previously reported [39], no toxicity was observed in OMP-18R5-treated mice, which displayed 256 consistent bodyweight, no signs of morbidity and no reduction in proliferation of normal nonadenoma gastric epithelial cells for the duration of treatment (Supplementary Figs. S3B-D). These 257 258 data strongly suggest Fzd receptors are rate-limiting for the growth of gastric adenomas in vivo, and in human GC cells in vitro. Given that Whts and Fzds can be expressed by non-epithelial 259 cells, we established gastric organoids from gp130^{F/F} antral adenomas using defined culture 260 conditions to determine if the anti-growth effects observed in *ap130^{F/F}* mice following OMP-18R5 261 treatment was systemic or cell intrinsic. gp130^{F/F} gastric adenoma organoids treated with OMP-262 18R5 or IWP-2 displayed reduced viability (MTT assay) and growth compared to vehicle control 263 treated organoids (Figs. 2K-M). This data confirms that Wnt ligands and Fzd receptors are 264 265 required cell intrinsically for the growth of gastric adenoma cells ex vivo.

266

267 Targeted FZD7 knockdown reduces gastric cancer colony formation

Inhibition of cell growth following OMP-18R5 treatment suggest that one of several Fzds targeted
by OMP-18R5 (FZD1, 2, 5, 7 and 8) is responsible for transmitting Wnt signals to GC cells. Gene
expression analysis narrows this down to *FZD2* and/or *FZD7*, as *FZD1*, *FZD5* and *FZD8* are
undetectable in these cell lines (Figs. 1A and B). We have previously shown that Fzd2 is unable

272 to compensate for the loss of Fzd7 in the intestinal epithelium [20], which may indicate Fzd7 plays 273 a predominant role in Wht signal transmission in gastric tissue. Indeed, FZD7 is commonly 274 upregulated in a variety of different cancer types, including gastric cancer, which is associated with poor clinical outcome [24, 42]. To determine the specific requirement of FZD7 for the growth 275 276 of human GC cells we performed colony formation assays. Cells transfected with FZD7-targeted shRNA (shFZD7) [16] had a marked decrease in colony growth, compared to scrambled shRNA 277 (shSCRAM) or empty vector (EV) controls (Figs. 3A and B), associated with decreased Wnt 278 279 signaling (Figs. 3C and D). These data suggest that Fzd7 is the predominant Wnt receptor 280 transmitting oncogenic Wnt signaling in GC cells. Importantly, growth inhibition following FZD7knockdown was rescued by co-transfection with a full-length FZD7 expression construct [33], 281 demonstrating the specificity of the shRNA and FZD7-regulated growth in human GC cells 282 283 (Supplementary Figs. S4A and B).

284

285 Conditional deletion of Fzd7 from gp130^{F/F} gastric tumors reduces cell proliferation

To determine the functional requirement of Fzd7 for gastric adenoma growth in vivo, we 286 conditionally deleted Fzd7 in the gastric adenomas of 8-week old Tff1CreERT2/+;gp130F/F:Fzd7^{1//1} 287 mice (*Cre⁺:ap130^{F/F}:Fzd7^{fl/fl}*) (Supplementary Fig. S4C), which allows robust recombination in 288 these adenomas [25]. Tamoxifen injected Cre^+ ; $gp130^{F/F}$; $Fzd7^{fl/fl}$ mice developed significantly 289 smaller and fewer antral gastric adenomas than their Cre-negative (Cre-; ap130^{F/F}:Fzd7^{fl/fl}) 290 tamoxifen-treated littermates (Figs. 3E-G and Supplementary Fig. S4D), supporting our previous 291 292 in vitro experiments demonstrating FZD7 inhibition is sufficient to block gastric adenoma growth 293 (Figs. 3A-D).

294

Fzd7 deficient cells are retained in gastric tumors and fail to proliferate

The growth of *qp130^{F/F}* gastric adenomas requires Stat3 [43]. Therefore we performed p-Stat3 296 IHC and Socs3 gRT-PCR which identified no alterations in Stat3 activity, and did not cause the 297 reduced growth of gastric adenomas in Cre^+ ; $gp130^{F/F}$; $Fzd7^{II/II}$ mice (Figs. 4A and B). This identifies 298 that Fzd7-mediated Wnt signaling is rate-limiting for Stat3-driven gastric adenomas, which have 299 300 no Wnt-activating mutations. Deletion of Fzd7 in normal, non-transformed gastric epithelium causes repopulation with Fzd7-proficient cells [19]. To monitor if repopulation occurs in 301 Cre⁺;gp130^{F/F};Fzd7^{fl/fl} adenomas, we performed PCR for the recombined Fzd7 floxed allele 302 (Fzd7⁴), which we have previously shown is lost during repopulation in the normal gastric 303 epithelium following Fzd7 deletion [19]. However, in gastric adenomas of Cre+;gp130^{F/F};Fzd7^{fl/fl} 304 mice 30 days post tamoxifen, we detect robust recombination of the $Fzd7^{\Delta}$ allele, demonstrating 305

306 that Fzd7 deleted cells are retained in these adenomas (Fig. 4C). In support, the expression of 307 *Fzd7* and many Wnt pathway components and target genes remain low in these adenomas (Fig. 308 4D, Supplementary Fig. S4E and Table S1). This suggests that the mechanism underlying smaller gastric adenomas following Fzd7 deletion is due to retention of Fzd7-deficient cells in the 309 310 adenoma that are unable to respond to proliferative Wnt signals, and thus fail to proliferate (Fig. 4E). To investigate this further, we performed IHC on serial sections to detect recombined (β -gal⁺, 311 Fzd7 deleted) cells and proliferating cells (PCNA⁺) in Cre⁺;gp130^{F/F};Fzd7^{fl/fl};LacZ mice and 312 observed a marked co-localisation of non-proliferative (PCNA⁻) cells with recombined cells (β-313 314 gal⁺) (Fig. 4F).

315

To monitor cellular changes following Fzd7 deletion in Cre+;gp130^{F/F};Fzd7^{#/#} mice, IHC for 316 apoptosis (Caspase-3) and differentiation (Muc5a and Gastrin) was performed (Supplementary 317 Fig. S4F). Muc5a⁺ and Gastrin⁺ cells were increased following Fzd7 deletion in 318 $Cre^+;gp130^{F/F};Fzd7^{fl/fl}$ mice compared to Fzd7-proficient gastric adenomas ($Cre^-;gp130^{F/F};Fzd7^{fl/fl}$). 319 This also suggests that gastric adenomas do not repopulate following Fzd7 deletion, as 320 321 repopulation in the normal gastric epithelium following Fzd7 deletion is associated with reduced 322 cell differentiation [19]. No change in the frequency of Caspase-3⁺ cells was observed (Supplementary Fig. S4D), indicating that deletion of *Fzd7* from adenoma cells does not trigger 323 324 apoptosis.

325

326 Cell intrinsic Wnt signaling via Fzd7 is required for Wnt-driven gastric adenomas

The *gp130^{F/F}* mice and MKN45 GC cells are wild-type for *APC*, and have no known Wnt-activating 327 mutations, suggesting that targeting Fzd7 may be effective in gastric adenomas and GC cells 328 without mutations to the Wnt pathway. However, some of the GC cell lines that responded to Fzd 329 330 therapy (MKN28 and MKN74) have mutant APC (https://portals.broadinstitute.org/ccle), suggesting that Fzd therapies can be effective in gastric adenomas with and without mutant APC. 331 In silico analysis of GC patient datasets identify mutations in several genes that regulate Wnt 332 signaling, demonstrating that this pathway is aberrantly activated in GC (Supplementary Fig. 333 S5A). To functionally investigate this, gastric organoids established from Tff1Cre^{ERT2/+};Apc^{fl/fl} 334 (Cre+:Apc^{#/#}) mice were treated with tamoxifen, to truncate Apc, and showed significant increase 335 in growth and proliferation (Fig. 5A), which was confirmed by Ki-67 staining and increased cell 336 337 viability (MTT assay) (Figs. 5A-C). A concordant increase in Wnt target gene expression was observed in hyperproliferative Apc mutant organoids (Fig. 5D). Treatment of Apc mutant 338 organoids with IWP-2 or OMP-18R5 prevented upregulation of the Wnt pathway and blocked 339

organoid proliferation (Figs. 5A-D), demonstrating that cell intrinsic Wnt secretion and Fzd
 receptors are required for gastric cells to activate Wnt signaling and regulate growth, even after
 mutation of *Apc* (Figs. 5A-D).

343

344 Fzd7 expression was increased in Apc mutant gastric organoids and subsequently downregulated in IWP-2 or OMP-18R5 treated organoids (Fig. 5E), therefore we examined whether Fzd7 is 345 responsible for transmitting Wnt signaling in Apc mutant gastric adenoma cells in vivo. 30 days 346 following tamoxifen, *Cre⁺;Apc^{fl/fl}* mice developed multiple, large intestinal-type gastric adenomas 347 348 with extensive hyperplasia in the antral stomach (Figs. 6A and B), which were not observed in tamoxifen-treated Cre⁻:Apc^{#/#} mice (Fig. 6A). Remarkably, co-recombination of Apc and Fzd7 349 alleles in Cre+; Apc^{1//I}; Fzd7^{1//I} mice inhibited the ability of Apc mutant cells to develop antral 350 adenomas (Figs. 6A and B). Gastric adenomas of Cre+;Apc^{11/11};Fzd7^{11/11} mice had significantly less 351 PCNA⁺ cells compared to Cre⁺;Apc^{fl/fl} mice (Figs. 6A and C). In common with gp130^{F/F} tumors, 352 353 deletion of Fzd7 in Apc deficient gastric adenomas also results in retention of Fzd7-deficient cells as monitored by expression of the $Fzd7^{\Delta}$ allele (Fig. 6D). 354

355

As expected, Wnt signaling is increased in gastric adenomas of *Cre⁺:Apc^{1//1}* mice, however, Wnt 356 signaling is not elevated in the non-adenoma antral epithelium of Cre^+ ; $Apc^{t/t}$; $Fzd7^{t/t}$ mice (Fig. 357 358 6E). This is supported by IHC for the surrogate markers of active Wnt signaling, β -catenin and 359 Myc (Supplementary Fig. S6A). IHC revealed a decrease in Muc5a⁺ and Gastrin⁺ cells following 360 Apc mutation (Supplementary Fig. S6B), while tamoxifen-treated Cre⁺; Apc^{1//1}; Fzd7^{1//1} mice display a modest restoration of mucus-secreting and gastrin-producing cells, similar to that observed in 361 *Cre⁺:ap130^{F/F}:Fzd7^{fl/fl}* mice (Supplementary Fig. S4D). Collectively, these data demonstrate that 362 Apc-mutant gastric phenotypes require functional Fzd7. 363

364

Fzd7-dependant Myc expression is required for the growth of gastric adenomas.

The transcription factor c-Myc is a well-characterised β-catenin/TCF target gene in the 366 gastrointestinal tract as c-Myc is required for all intestinal tumor phenotypes following Apc-367 mediated activation of Wnt signaling [35]. Myc is upregulated in our gastric adenoma mouse 368 models and human GC cell lines, and inhibition of Fzd7 prevents this upregulation (Figs. 2H, 3C, 369 4D, 5D and 6E). Conditional deletion of *c-Myc* in *Tff1Cre*^{ERT2/+};*Apc*^{fl/fl};*c-Myc*^{fl/fl}(*Cre*⁺;*Apc*^{fl/fl};*Myc*^{fl/fl}) 370 mice showed complete absence of antral adenoma formation and Wnt activation compared to 371 Cre⁺;Apc^{il/il} mice (Supplementary Fig. S7), indicating Fzd7-dependant expression of Myc is 372 373 required for the growth of Apc mutant gastric adenomas.

To determine whether elevated levels of MYC can rescue GC cell growth suppression following *FZD7* knockdown, GC cells were co-transfected with FZD7shRNA and MSCV-MYC expression plasmids and grown as colonies in soft agar for 2 weeks. Compared to control (EV) transfected cells, co-transfected cells (FZD7shRNA and MSCV-MYC) showed no difference in the number of colonies formed (Supplementary Fig. S7G-I), which suggests that overexpression of MYC is able to rescue the growth suppressive effects of FZD7 knockdown in GC cells.

380

381 Discussion

Expression of Fzd receptors is deregulated in several cancers, including gastric cancer [4, 21, 42]. Here we show for the first time that Fzd receptors are rate-limiting for the growth of gastric adenomas *in vivo*. We further elucidate that Fzd7 is the predominant Wnt receptor transmitting cell-intrinsic Wnt signals in human GC cells.

386

387 In vitro studies have shown that targeted inhibition of Fzd is sufficient to block growth of GC cells [24, 44]. However, it is well documented that in vitro studies do not fully recapitulate the complex 388 389 cellular and molecular interactions present in tumors [45]. Here, we demonstrate that gastric 390 adenomas require Fzd7 for optimal growth using genetic and pharmacological strategies in two 391 independent mouse models. Our findings support our previous work [39] demonstrating that 392 targeting multiple Fzd receptors blocks the growth of several different cancers, which we now 393 extend to GC. Using ex-vivo adenoma-derived organoids we demonstrate these anti-growth 394 effects are cell intrinsic as OMP-18R5 blocks the growth of gastric adenoma-derived organoids in 395 the absence of immune or stromal cells.

396

As previously observed in the normal gastric epithelium [19], genetic inhibition of *Fzd7* in gastric adenomas induces upregulation of other *Fzd* genes (Table S1), however, these are insufficient to compensate and promote gastric adenoma growth. This suggests that specific targeting of Fzd7 is an attractive therapeutic strategy for the treatment of gastric cancer.

401

Deletion of *Fzd7* in the normal gastric epithelium triggers repopulation [19] which could be a possible explanation for why *Fzd7*-deficient gastric adenomas are smaller. Epithelial repopulation is an effective tissue mechanism that helps the gastric epithelium to survive the harsh conditions of the stomach. Here we show that repopulation is not preserved in gastric adenomas, which contain aberrant cell signaling and tissue architecture, and therefore *Fzd7*-deficient cells remain in the adenoma but are unable to respond to Wnt signals and thus do not proliferate. 408 One feature of inflammation-associated tumors in the gastrointestinal tract is phosphorylated Stat3 (p-Stat3), which regulates many cancer hallmarks [43]. Gastric adenomas in *gp130^{FF}* mice 409 410 do not harbor any Wnt-activating mutations [41], however, they display high levels of Wnt signaling. Stat-3 has been shown to activate Wnt signaling, which would allow pathway activation 411 412 in the absence of Wnt mutations in gp130^{FF} adenomas [46, 47]. Indeed, Wnt and gp130/Stat3 signaling operate in parallel during gastric tumorigenesis as active p-Stat3 levels remain high in 413 Fzd7 deleted adenomas, demonstrating that Wnt/Fzd7 signaling is rate-limiting for Stat3-driven 414 gastric adenomas. Similarly, mTORC1 signaling is also rate-limiting for *ap130^{FF}* adenoma growth 415 416 independent of Stat3 [41].

417

Recent large-scale sequencing of human gastric tumors has identified environmental and genetic 418 factors associated with increased pathology, which include aberrant Wnt signaling [48-50]. 419 420 Importantly, these genomic studies are yet to be validated with functional interrogation in vivo, 421 which are essential to understand the therapeutic potential of targeting Wnt signaling in gastric cancer [21]. We and others have demonstrated that Fzd7 inhibition is sufficient to block Wnt 422 423 signaling in cells with mutant APC [17, 51]. Interestingly, ~37% of APC mutant gastric tumors are 424 mutant for RNF43 (regulates Fzd on the cell surface [12]), demonstrating that Fzd is deregulated 425 in a subset of APC mutant gastric tumors (http://www.cbioportal.org/). Interestingly RNF43 and 426 APC mutations are mutually exclusive in colon tumors suggesting that CRC and GC cells 427 preferentially select different Wnt mutations that confer optimal or 'just-right' levels of Wnt 428 signaling required for tumor growth [52, 53].

429

Furthermore, we have shown that Myc is required for the gastric adenoma phenotypes associated with *Apc* mutation. These findings are reminiscent of the role played by Myc in the intestinal epithelium following *Apc* mutation [35], and thus place the Wnt/Fzd7/Myc signaling axis as an attractive therapeutic target for gastric cancer. Encouragingly, next generation bromodomain (BET) inhibitors are effective in killing patient-derived GC cells [54]. Importantly, this provides justification for testing a combination of BET and Wnt inhibitors in GC, which we have previously shown is effective at blocking the growth of human colon cancer cells [55].

437

New generation PORCN inhibitors are in clinical trials for solid tumors, which our results show
may be effective in gastric cancer, however these target the secretion of all Wnt ligands.
Collectively, we demonstrate that targeted inhibition of Wnt receptors, specifically Fzd7, is ratelimiting for the growth of gastric adenomas with and without *Apc* mutations. This provides a broad

scope for the application of this therapeutic strategy for the treatment of GC, with potentially less

side effects than targeting all Wnt secretion with PORCN inhibitors, and will directly inform clinical

444 trials to treat GC patients with OMP-18R5 (Vantictumab), which only targets 5 out of the 10 Fzd

445 family.

446

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579 Figure Legends

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581 Figure 1. Inhibition of Wnt or Fzd blocks gastric cancer cell growth.

- 582 A. qRT-PCR for *FZD* gene expression in MKN28 gastric cancer cells. Expression shown 583 relative to housekeeper (β 2M), n=4 biological replicates.
- 584 B. qRT-PCR for *FZD* gene expression in MKN74 gastric cancer cells. Expression shown 585 relative to housekeeper (β 2M), n=4 biological replicates.
- 586 C. Quantification of cell colonies (>50 cells) from MKN28 gastric cancer cells grown in agar 587 for 2 weeks following treatment with vehicle control (2.5%DMSO+IgG), IWP-2 (10μ M) or 588 OMP-18R5 (10μ g/mI). Treatments were replaced every 4 days for the duration of 2 weeks. 589 Individual experiments were repeated three times. Colonies were counted with ImageJ (*= 590 p<0.05, mean ±SEM, Mann-Whitney).
- 591 D. Quantification of cell colonies (>50 cells) from MKN74 gastric cancer cells grown in agar 592 for 2 weeks following treatment with vehicle control (2.5%DMSO+IgG), IWP-2 (10μ M) or 593 OMP-18R5 (10μ g/mI). Treatments were replaced every 4 days for the duration of 2 weeks. 594 Individual experiments were repeated three times. Colonies were counted with ImageJ (*= 595 p<0.05, mean ±SEM, Mann-Whitney).
- E. TOPflash assay on MKN28 cells treated 24hrs with DMSO, IWP-2 (10μM) or OMP-18R5 (10μg/ml) (**= p<0.005, mean ±SEM, n=9 biological replicates, Mann-Whitney). Individual experiments were repeated three times.
- 599 F. TOPflash assay on MKN74 cells treated 24hrs with DMSO, IWP-2 (10 μ M) or OMP-18R5 600 (10 μ g/ml) (**= p<0.005, mean ±SEM, n=9 biological replicates, Mann-Whitney). Individual 601 experiments were repeated three times.
 - G. qRT-PCR for *CD44* in MKN28 and MKN74 cells described in E and F (mean ±SEM, n=6 biological replicates, Mann-Whitney). Individual experiments were repeated twice.
 - H. qRT-PCR for *AXIN2* in MKN28 and MKN74 cells described in E and F (mean ±SEM, n=6 biological replicates, Mann-Whitney). Individual experiments were repeated twice.

Figure 2. Inhibition of Fzd receptors reduces cell intrinsic Wnt signaling and gastric adenoma burden.

- 609A. qRT-PCR for Wnt ligands in $gp130^{F/F}$ adenomas compared to normal gastric epithelium610(*= p<0.05, mean ±SEM, n=4 mice, Mann-Whitney).</td>
- 611 B. qRT-PCR for Fzd receptors in $gp130^{F/F}$ adenomas compared to normal gastric epithelium 612 (*= p<0.05, mean ±SEM, n=4 mice, Mann-Whitney).
- 613 C. qRT-PCR for Wnt target genes in $gp130^{F/F}$ adenomas compared to normal gastric 614 epithelium (*= p<0.05, mean ±SEM, n=4 mice, Mann-Whitney).
- D. Whole mount images of 8-9 week old *gp130^{F/F}*mice treated with control IgG or OMP-18R5
 over the course of 30 days and harvested. Black and white arrows show gastric tumors.
- E. Weights of gastric adenomas from mice described in D (***= p<0.001, mean ±SEM, n=9 mice, Mann-Whitney).
- F. Quantification of gastric adenomas in mice described in D (***= p<0.001, mean ±SEM, n=9 mice, Mann-Whitney).
- 621 G. qRT-PCR for Fzd receptors in mice described in D (**= p<0.005, mean ±SEM, n=9 mice, 622 Mann-Whitney).

H. qRT-PCR for Wnt target genes in mice described in D (**= p<0.005, mean ±SEM, n=9 mice, Mann-Whitney).

 bars = 100µm. Quantification of PCNA* cells from adenomas sections described in 1 (*= p<0.05, mean ±SEM, n=4 mice, Mann-Whitney). K. Representative DIC images of <i>gp130^{-pE}</i> adenoma-derived organoids treated with vehicle control (25%DMSO-1(g), IWP-2 (10µM) or OMP-18RS (10µg/ml) and cultured for 5 days. Green arrows indicate viable organoids. Red arrows indicate dying/atrophic organoids. Scale bar = 200 µm L. MTT viability assay performed on organoids described in K (*= p<0.05, mean ±SEM, n=3 biological replicates, Mann-Whitney). M. Measurement (diameter) of organoids described in K. Measurements were quantified in ImageJ (*= p<0.05, mean ±SEM, n=3 biological replicates, Mann-Whitney). Figure 3. Targeted inhibition of Fzd7 reduces gastric cancer clonogenicity and adenoma burden. A. Representative DIC images of MKN28 and MKN74 cells transfected with empty vector (EV), scrambled (shSCRAM) or FZD7-specific shRNA (FZD7shRNA) and grown in agar. Scale bars = 200µm Quantification of cell colonies from experiment described in A (*= p<0.05, mean ±SEM, n=3 biological replicates, Mann-Whitney). Individual experiments were repeated twice. q RT-PCR for Wnt taget genes on MKN28 and MKN74 cells transfected with empty vector (EV), scrambled (shSCRAM) or FZD7-specific ShRNA (Fzd7shRNA) (*= p<0.05, mean ±SEM, n=3 biological replicates, Mann-Whitney). D TOPHash assay on MKN28 and MKN74 cells described in C (**= p<0.05, mean ±SEM, n=3 biological replicates, Mann-Whitney). D TOPHash assay on MKN28 and MKN74 cells described in C (**= p<0.05, mean ±SEM, n=3 biological replicates, Mann-Whitney). C Quantification of gastric adenomas per mouse described in E (**= p<0.005, mean ±SEM, n=3 biological replicates, Mann-Whitney). G Quantification of gastric adenomas per mouse described in E (**= p<0.005, mean ±SEM, n=7 mice, Mann-Whitney). G Quantification of gast	625	١.	Immunohistochemistry for PCNA on adenomas sections from mice described in D. Scale
 J. Quantification of PCNA⁺ cells from adenomas sections described in 1 (*= p<0.05, mean ±SEM, n=4 mice, Mann-Whitney). K. Representative DIC images of <i>gp</i>/30^{FF} adenoma-derived organoids treated with vehicle control (2.5%DMSO+IgG), IMP-2 (10µM) or OMP-18RS (10µg/ml) and cultured for 5 days. Green arrows indicate viable organoids. Red arrows indicate dying/atrophic organoids. Scale bar = 200 µm L. MTT viability assay performed on organoids described in K (*= p<0.05, mean ±SEM, n=3 biological replicates, Mann-Whitney). M. Measurement (diameter) of organoids described in K. Measurements were quantified in Image) (*= p<0.05, mean ±SEM, n=3 biological replicates, Mann-Whitney). Flgure 3. Targeted inhibition of Fzd7 reduces gastric cancer clonogenicity and adenoma burden. A. Representative DIC images of MKN28 and MKN74 cells transfected with empty vector (EV), scrambled (shSCRAM) or FZD7-specific shRNA (FZD7shRNA) and grown in agar. Scale bars = 200µm Quantification of cell colonies from experiment described in A (*= p<0.05, mean ±SEM, n=3 biological replicates, Mann-Whitney). Individual experiments were repeated twice. C. qRT-PCR for Wnt taget genes on MKN28 and MKN74 cells transfected with empty vector (EV), scrambled (shSCRAM) or FZD7-specific shRNA (Fzd7shRNA) (*= p<0.05, mean ±SEM, n=3 biological replicates, Mann-Whitney). Individual experiments were repeated twice. Representative images of tamoxifen-treated <i>Tff1Cre^{ERT2+}</i> (Cre) or <i>Tff1Cre^{ERT2+}</i> (Cre) stomachs following <i>Fzd7</i> deletion. Black arrows indicate gastric tumors. F. Weights of gastric adenomas per mouse described in E (**= p<0.005, mean ±SEM, n=7 mice, Mann-Whitney). G. Quantification of gastric adenomas per mouse described in E (**= p<0.005, mean ±SEM, n=7 mice, Mann-Whitney). G. Quantification of PZd7 from gastric tumors decreases cell proliferation. K. Weights	626		bars = 100μm.
 45EM, n=4 mice, Mann-Whitney). K. Representative DIC images of <i>gp130^{erf}</i> adenoma-derived organoids treated with vehicle control (2.5%DMSO+IgG). IMP-2 (10µM) or OMP-18RS (10µg/ml) and cultured for 5 days. Scale bar = 200 µm L. MTT viability assay performed on organoids described in K (*= p<0.05, mean ±SEM, n=3 biological replicates, Mann-Whitney). M. Measurement (diameter) of organoids described in K. Measurements were quantified in ImageJ (*= p<0.05, mean ±SEM, n=3 biological replicates, Mann-Whitney). Figure 3. Targeted inhibition of Fzd7 reduces gastric cancer clonogenicity and adenoma burden. A. Representative DIC images of MKN28 and MKN74 cells transfected with empty vector (EV), scrambled (shSCRAM) or FZD7-specific shRNA (FZD7shRNA) and grown in agar. Scale bars = 200µm Quantification of cell colonies from experiment described in A (*= p<0.05, mean ±SEM, n=3 biological replicates, Mann-Whitney). Individual experiments were repeated twice. C. qRT-PCR for Wnt taget genes on MKN28 and MKN74 cells transfected with empty vector (EV), scrambled (shSCRAM) or FZD7-specific shRNA (FZd7shRNA) (*= p<0.05, mean ±SEM, n=3 biological replicates, Mann-Whitney). D. TOPflash assay on MKN28 and MKN74 cells described in C (***= p<0.01, mean ±SEM, n=9 biological replicates, Mann-Whitney). E. Representative images of tamoxifen-treated <i>Tff1Cre^{ERT2+}</i> (Cre) or <i>Tff1Cre^{ERT2+}</i> (Cre) stomachs following <i>Fzd7</i> deficien. Black arrows indicate gastric tumors. Figure 4. Deletion of <i>Fzd7</i> from gastric tumors decreases cell proliferation. A. Immunohistochemistry (IHC) for p-Stat3 on adenoma sections from <i>Fzd7^{thf1}gp130^{erf}</i> mice (C<i>re</i> or <i>Cre</i>) 30 days after tamoxifen treatment. Scale bars = 100µm. G. Quantification of gastric adenomas from mice described in A (*= p<0.05, mean ±SEM, n=7 mice, Mann-Whitney). G. Quantification of PCA7 from gastric	627	J.	Quantification of PCNA ⁺ cells from adenomas sections described in I (*= p<0.05, mean
 K. Representative DIC images of <i>gp130^{cFE}</i> adenoma-derived organoids treated with vehicle control (2.5%DMSO+IgG), IWP-2 (10µM) or OMP-18R5 (10µg/ml) and cultured for 5 days. Green arrows indicate viable organoids. Red arrows indicate dying/atrophic organoids. Scale bar = 200 µm L. MTT viability assay performed on organoids described in K. (*= p<0.05, mean ±SEM, n=3 biological replicates, Mann-Whitney). Measurement (diameter) of organoids described in K. Measurements were quantified in ImageJ (*= p<0.05, mean ±SEM, n=3 biological replicates, Mann-Whitney). Figure 3. Targeted inhibition of Fzd7 reduces gastric cancer clonogenicity and adenoma burden. A. Representative DIC images of MKN28 and MKN74 cells transfected with empty vector (EV), scrambled (shSCRAM) or FZD7-specific shRNA (FZD7shRNA) and grown in agar. Scale bars = 200µm Guantification of cell colonies from experiment described in A (*= p<0.05, mean ±SEM, n=3 biological replicates, Mann-Whitney). Individual experiments were repeated twice. C. qRT-PCR for Wnt taget genes on MKN28 and MKN74 cells transfected with empty vector (EV), scrambled (shSCRAM) or FZD7-specific shRNA (Fzd7shRNA) (*= p<0.05, mean ±SEM, n=3 biological replicates, Mann-Whitney). TOPItash assay on MKN28 and MKN74 cells transfected with empty vector (EV), scrambled (shSCRAM) or FZD7-specific shRNA (Fzd7shRNA) (*= p<0.05, mean ±SEM, n=9 biological replicates, Mann-Whitney). TOPItash assay on MKN28 and MKN74 cells transfected with empty vector (EV), scrambled (shSCRAM) or FZD7-specific shRNA (Fzd7shRNA) (*= p<0.05, mean ±SEM, n=9 biological replicates, Mann-Whitney). TOPItash assay on MKN28 and MKN74 cells transfected with empty vector (EV), scrambled (shSCRAM) or FZD7-specific shRNA (Fzd7shRNA) (*= p<0.05, mean ±SEM, n=7 mice, Mann-Whitney). Guantification of gastric adenomas per mouse described in C (**= p<0.005, mean ±SEM, n=7 mice, Mann-Whitn	628		±SEM, n=4 mice, Mann-Whitney).
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 E. Representative images of tamoxifen-treated <i>Tff1Cre^{ERT2/-}</i> (<i>Cre⁺</i>) or <i>Tff1Cre^{ERT2/+}</i> (<i>Cre⁺</i>) stomachs following <i>Fzd7</i> deletion. Black arrows indicate gastric tumors. F. Weights of gastric adenomas per mouse described in E (**= p<0.005, mean ±SEM, n=7 mice, Mann-Whitney). G. Quantification of gastric adenomas per mouse described in E (**= p<0.005, mean ±SEM, n=7 mice, Mann-Whitney). Figure 4. Deletion of <i>Fzd7</i> from gastric tumors decreases cell proliferation. A. Immunohistochemistry (IHC) for p-Stat3 on adenoma sections from <i>Fzd7^{Mfl};gp130^{F/F}</i> mice (<i>Cre⁻</i> or <i>Cre⁺</i>) 30 days after tamoxifen treatment. Scale bars = 100µm. B. qRT-PCR for <i>Socs3</i> on gastric adenomas from mice described in A (*= p<0.05, mean ±SEM, n=4 mice, Mann-Whitney). C. Conventional PCR to detect recombination of <i>Fzd7^{Mfl}</i> allele (Fzd7^Δ) in gastric adenomas from mice described in A (*= p<0.005, mean ±SEM, n=7 mice, Mann-Whitney). E. Quantification of PCNA⁺ cells from adenoma sections described in A (*= p<0.05, mean ±SEM, n=3 mice, Mann-Whitney). F. Representative IHC images for β-galactosidase (detecting allelic recombination) and PCNA (proliferation) on serial sections from <i>Tff1Cre⁺;Fzd7^{Mfl};gp130^{F/F};LacZ</i> or <i>Tff1Cre⁺;Fzd7^{Mfl};gp130^{F/F};LacZ</i> mice 30 days following tamoxifen. Note, yellow dashed 	650		times.
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 B. qRT-PCR for <i>Socs3</i> on gastric adenomas from mice described in A (*= p<0.05, mean ±SEM, n=4 mice, Mann-Whitney). C. Conventional PCR to detect recombination of <i>Fzd7</i>^{#/#} allele (Fzd7^Δ) in gastric adenomas from mice described in A. D. qRT-PCR for Wnt target genes in gastric adenomas from mice described in A (**= p<0.005, mean ±SEM, n=7 mice, Mann-Whitney). E. Quantification of PCNA⁺ cells from adenoma sections described in A (*= p<0.05, mean ±SEM, n=3 mice, Mann-Whitney). F. Representative IHC images for β-galactosidase (detecting allelic recombination) and PCNA (proliferation) on serial sections from <i>Tff1Cre⁻;Fzd7</i>^{#/#};gp130^{F/F};LacZ or <i>Tff1Cre⁺;Fzd7</i>^{#/#};gp130^{F/F};LacZ mice 30 days following tamoxifen. Note, yellow dashed 	660		(<i>Cre</i> ⁻ or <i>Cre</i> ⁺) 30 days after tamoxifen treatment. Scale bars = $100\mu m$.
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 D. qRT-PCR for Wnt target genes in gastric adenomas from mice described in A (**= p<0.005, mean ±SEM, n=7 mice, Mann-Whitney). E. Quantification of PCNA⁺ cells from adenoma sections described in A (*= p<0.05, mean ±SEM, n=3 mice, Mann-Whitney). F. Representative IHC images for β-galactosidase (detecting allelic recombination) and PCNA (proliferation) on serial sections from <i>Tff1Cre⁻;Fzd7^{fl/fl};gp130^{F/F};LacZ</i> or <i>Tff1Cre⁺;Fzd7^{fl/fl};gp130^{F/F};LacZ</i> mice 30 days following tamoxifen. Note, yellow dashed 	664		from mice described in A.
666p<0.005, mean ±SEM, n=7 mice, Mann-Whitney).667E. Quantification of PCNA+ cells from adenoma sections described in A (*= p<0.05, mean ±SEM, n=3 mice, Mann-Whitney).668±SEM, n=3 mice, Mann-Whitney).669F. Representative IHC images for β-galactosidase (detecting allelic recombination) and PCNA (proliferation) on serial sections from Tff1Cre ⁻ ;Fzd7 ^{fl/fl} ;gp130 ^{F/F} ;LacZ or Tff1Cre ⁺ ;Fzd7 ^{fl/fl} ;gp130 ^{F/F} ;LacZ mice 30 days following tamoxifen. Note, yellow dashed	665	D.	qRT-PCR for Wnt target genes in gastric adenomas from mice described in A (**=
667E. Quantification of PCNA+ cells from adenoma sections described in A (*= p<0.05, mean668 \pm SEM, n=3 mice, Mann-Whitney).669F. Representative IHC images for β-galactosidase (detecting allelic recombination) and670PCNA (proliferation) on serial sections from <i>Tff1Cre⁻;Fzd7^{fl/fl};gp130^{F/F};LacZ</i> or671 <i>Tff1Cre⁺;Fzd7^{fl/fl};gp130^{F/F};LacZ</i> mice 30 days following tamoxifen. Note, yellow dashed	666		p<0.005, mean ±SEM, n=7 mice, Mann-Whitney).
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669F. Representative IHC images for β-galactosidase (detecting allelic recombination) and670PCNA (proliferation) on serial sections from $Tff1Cre^{-};Fzd7^{fl/fl};gp130^{F/F};LacZ$ or671 $Tff1Cre^{+};Fzd7^{fl/fl};gp130^{F/F};LacZ$ mice 30 days following tamoxifen. Note, yellow dashed	668		±SEM, n=3 mice, Mann-Whitney).
670 PCNA (proliferation) on serial sections from $Tff1Cre^{-};Fzd7^{fl/fl};gp130^{F/F};LacZ$ or 671 $Tff1Cre^{+};Fzd7^{fl/fl};gp130^{F/F};LacZ$ mice 30 days following tamoxifen. Note, yellow dashed	669	F.	Representative IHC images for β -galactosidase (detecting allelic recombination) and
671 <i>Tff1Cre⁺;Fzd7^{#//#};gp130^{F/F};LacZ</i> mice 30 days following tamoxifen. Note, yellow dashed	670		PCNA (proliferation) on serial sections from Tff1Cre ⁻ ;Fzd7 ^{#/#} ;gp130 ^{F/F} ;LacZ or
	671		Tff1Cre+;Fzd7 ^{fl/fl} ;gp130 ^{F/F} ;LacZ mice 30 days following tamoxifen. Note, yellow dashed

672		lines demarcate areas of allelic recombination, which correspond to reduced proliferation
673		and black dashed lines represent areas of non-recombined cells. Scale bars = $100\mu m$.
674	Figure	5. Wat/End inhibition reduces Anomytent restric erronsid preliferation
675	rigure	Depresentative DIC and immunofluereseance images of <i>Tff1CrevAnd^{fff}</i> ergeneide treated
070 677	А.	for 24brs with tamovifon (tmx 100nM) IWP-2 (10uM) or OMP-18B5 (10ug/ml) Green
678		arrows indicate hyperproliferative organoids Red arrows indicate growth-constrained
679		organoids. Scale bars = $200 \mu m$.
680	В.	MTT viability assay performed on organoid cultures described in A (***= p<0.001, mean
681		±SEM, n=3 biological replicates, Mann-Whitney). Individual experiments were repeated
682		twice.
683	C.	Measurement of organoid size (μ m) from cultures described in A (***= p<0.001, mean
684		±SEM, n=3 biological replicates, Mann-Whitney).
685	D.	qRT-PCR for Wnt target genes on organoid cultures described in A (*= p<0.05, mean
686	_	±SEM, n=3 biological replicates, Mann-Whitney).
687	E.	qRI-PCR for Fzd receptors on organoid cultures described in A. Expression of Fzd snown
600		and any water
688 680		as Log₂ ratio.
688 689 690	Figure	as Log_2 ratio. 6 Deletion of <i>EzdZ</i> rescues gastric adenoma formation following <i>Apc</i> truncation
688 689 690 691	Figure A	as Log ₂ ratio. 6. Deletion of <i>Fzd7</i> rescues gastric adenoma formation following <i>Apc</i> truncation. Bepresentative whole mount and IHC (PCNA) on wild-type (<i>Tff1Cre⁻:Apc</i>^{fi/fi}). <i>Apc</i> mutant
688 689 690 691 692	Figure A.	as Log ₂ ratio. 6. Deletion of <i>Fzd7</i> rescues gastric adenoma formation following <i>Apc</i> truncation. Representative whole mount and IHC (PCNA) on wild-type (<i>Tff1Cre⁻;Apc^{fl/fl}</i>), <i>Apc</i> mutant (<i>Tff1Cre⁺:Apc^{fl/fl}</i>) and <i>Apc/Fzd7</i> mutant mice (<i>Tff1Cre⁺:Apc^{fl/fl};Ezd7^{fl/fl}</i>) 30 days following
688 689 690 691 692 693	Figure A.	as Log ₂ ratio. 6. Deletion of <i>Fzd7</i> rescues gastric adenoma formation following <i>Apc</i> truncation. Representative whole mount and IHC (PCNA) on wild-type ($Tff1Cre^{-};Apc^{fl/fl}$), <i>Apc</i> mutant ($Tff1Cre^{+};Apc^{fl/fl}$) and <i>Apc/Fzd7</i> mutant mice ($Tff1Cre^{+};Apc^{fl/fl};Fzd7^{fl/fl}$) 30 days following tamoxifen. Black arrows indicate gastric adenomas in top panels. Scale bars = 100µm.
688 689 690 691 692 693 694	Figure A. B.	as Log ₂ ratio. 6. Deletion of <i>Fzd7</i> rescues gastric adenoma formation following <i>Apc</i> truncation. Representative whole mount and IHC (PCNA) on wild-type (<i>Tff1Cre⁻;Apc^{fl/fl}</i>), <i>Apc</i> mutant (<i>Tff1Cre⁺;Apc^{fl/fl}</i>) and <i>Apc/Fzd7</i> mutant mice (<i>Tff1Cre⁺;Apc^{fl/fl};Fzd7^{fl/fl}</i>) 30 days following tamoxifen. Black arrows indicate gastric adenomas in top panels. Scale bars = 100µm. Weights of gastric adenomas from harvested mice described in A (**= p<0.005, mean
688 689 690 691 692 693 694 695	Figure A. B.	as Log ₂ ratio. 6. Deletion of <i>Fzd7</i> rescues gastric adenoma formation following <i>Apc</i> truncation. Representative whole mount and IHC (PCNA) on wild-type (<i>Tff1Cre⁻;Apc^{fl/fl}</i>), <i>Apc</i> mutant (<i>Tff1Cre⁺;Apc^{fl/fl}</i>) and <i>Apc/Fzd7</i> mutant mice (<i>Tff1Cre⁺;Apc^{fl/fl};Fzd7^{fl/fl}</i>) 30 days following tamoxifen. Black arrows indicate gastric adenomas in top panels. Scale bars = 100µm. Weights of gastric adenomas from harvested mice described in A (**= p<0.005, mean ±SEM, n=7 mice, Mann-Whitney).
688 690 691 692 693 694 695 696	Figure A. B. C.	as Log ₂ ratio. 6. Deletion of <i>Fzd7</i> rescues gastric adenoma formation following <i>Apc</i> truncation. Representative whole mount and IHC (PCNA) on wild-type (<i>Tff1Cre⁻;Apc^{fl/fl}</i>), <i>Apc</i> mutant (<i>Tff1Cre⁺;Apc^{fl/fl}</i>) and <i>Apc/Fzd7</i> mutant mice (<i>Tff1Cre⁺;Apc^{fl/fl};Fzd7^{fl/fl}</i>) 30 days following tamoxifen. Black arrows indicate gastric adenomas in top panels. Scale bars = 100µm. Weights of gastric adenomas from harvested mice described in A (**= p<0.005, mean ±SEM, n=7 mice, Mann-Whitney). Quantification of PCNA ⁺ cells in adenoma sections from mice described in A (***= p<0.001,
688 689 690 691 692 693 694 695 696 697	Figure A. B. C.	as Log ₂ ratio. 6. Deletion of <i>Fzd7</i> rescues gastric adenoma formation following <i>Apc</i> truncation. Representative whole mount and IHC (PCNA) on wild-type (<i>Tff1Cre⁻;Apc^{fl/fl}</i>), <i>Apc</i> mutant (<i>Tff1Cre⁺;Apc^{fl/fl}</i>) and <i>Apc/Fzd7</i> mutant mice (<i>Tff1Cre⁺;Apc^{fl/fl};Fzd7^{fl/fl}</i>) 30 days following tamoxifen. Black arrows indicate gastric adenomas in top panels. Scale bars = 100µm. Weights of gastric adenomas from harvested mice described in A (**= p<0.005, mean ±SEM, n=7 mice, Mann-Whitney). Quantification of PCNA ⁺ cells in adenoma sections from mice described in A (***= p<0.001, mean ±SEM, n=3 mice, Mann-Whitney).
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