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Parallel states of pathological Wnt signaling in neonatal brain injury and colon cancer

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Abstract

In colon cancer, mutation of the Wnt repressor *Adenomatous polyposis coli (APC)* leads to a state of aberrant and unrestricted "high-activity" signaling. However, relevance of high Wnt tone in non-genetic human disease is unknown. Here we demonstrate that distinct Wnt activity functional states determine oligodendrocyte precursor (OPC) differentiation and myelination. Murine OPCs with genetic Wnt dysregulation (high tone) express multiple genes in common with colon cancer including Lef1, SP5, Ets2, Rnf43 and Dusp4. Surprisingly, we find that OPCs in lesions of hypoxic human neonatal white matter injury upregulate markers of high Wnt activity and lack expression of APC. Finally, we show lack of Wnt repressor tone promotes permanent white matter injury after mild hypoxic insult. These findings suggest a state of pathological high-activity Wnt signaling in human disease tissues that lack pre-disposing genetic mutation.

Author Contributions

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S.P.J.F. conceived of and performed all experiments and analysis, with the exception of the following: E.P.H performed and analyzed all experiments related to *in vitro* OLP cultures. J.C.S., T.J.Y. and L.S. helped analyze Wnt pathway activation in murine hypoxic injury. S.E.B. performed bioinformatics. E.J.H. procured human brain developmental tissue. S.L. provided advice on ChIP protocol and analysis. D.H.R conceived the experiments and oversaw all aspects of the analysis. The paper was written by S.P.J.F. and D.H.R.

Introduction

Activation of canonical Wnt signaling results in the stabilization of β -catenin and its translocation to the nucleus to associate with TCF family transcriptional protein partners. In the intestine, low activity signaling mediated by TCF4- β -catenin is necessary for normal turnover of the epithelium throughout life and maintenance of Lgr5+ crypt stem cells (1). A fine balance exists in the colon between Wnt pathway activation and inhibition in the determination of this low activity state, and numerous control mechanisms therefore exist to limit Wnt signaling levels (2). Germ line mutation of adenomatous polyposis coli (APC) causes familial predisposition to colon cancer (3). In the absence of APC, failure to degrade β-catenin results in unrestricted Wnt signaling, which reaches a threshold to elicit expression of LEF1(4). This high LEF1- β -catenin activity state drives uncontrolled epithelial growth and colon adenocarcinoma (5, 6). Similar low/high Wnt signaling states also distinguish normal hematopoies versus leukemias with mutations in Wnt pathway repressor genes (7). Many other markers have been identified from expression profiling that distinguish colon cancer from normal intestinal stem cell populations (3). Some of these factors have been shown to function as suppressors of the Wnt pathway, such as Rnf43, an E3 ligase that ubiquitinates frizzled receptors (8). Although in principle, such markers might be useful in detecting a high-activity state in a variety of human diseases, it has yet to be shown whether pathological Wnt signaling can be achieved in non-genetically-based conditions.

Oligodendrocytes are the myelinating cells of the central nervous system (CNS) and are cellular targets of injury in human disease such as multiple sclerosis (MS) (9) and neonatal white matter injury (WMI) leading to cerebral palsy (CP) (10, 11). Pathological analysis of these demyelinated lesions has revealed the presence of oligodendrocyte precursor cells (OPCs) arrested at an immature/premyelinating stage (12–14). As remyelination of experimental rodent demyelinated lesions is reproducibly efficient, why do human lesions fail to myelinate? This question prompted us to investigate whether a high pathological Wnt tone could be demonstrated in human neonatal subcortical white matter affected by hypoxic ischemic encephalopathy (HIE), despite the fact that this is an injury-induced, non-genetic condition.

Results

Common markers of high Wnt tone across diverse tissue types

Although Wnt signaling can delay timing of OPC differentiation (15–18), it is unclear whether signaling can achieve pathological-high levels to account for fixed demyelinated lesions in human injury. Indeed, the notion of a high-activity Wnt signaling state in human WMI, or other non-genetic disorders, has not been investigated. To define markers for high Wnt signaling across various tissues, we first compared mRNA transcripts upregulated in mouse Wnt-activated OPCs (*Olig2cre/DA-Cat*: (15)) and human colon carcinomas that are commonly driven by mutations of the Wnt pathway such as loss of *APC* function (Suppl. Tab. 1; (2, 3)). This revealed over 47 genes with significant upregulation (Suppl. Tab. 1). We reasoned that such information might identify both novel targets of the Wnt pathway in colon cancer and new high-activity pathway markers in OPCs (Fig. 1a). We focused on LEF1, SP5, RNF43, DUSP4 and ETS2 based on the following criteria: (1) upregulation

associated with high-activity Wnt signaling in other studies, (2) known functions in colon cancer and (3) markers available for immunohistochemistry (IHC) in human and rodent samples. All five markers were expressed in adenocarcinoma (Fig. 1b). DUSP4, a MAP-kinase phosphatase protein has not previously been reported as a Wnt-activated target; thus, our approach serves as a gene discovery tool.

High activity Wnt elicits Lef1 and OPC maturation arrest

In the adult mammalian CNS, APC expression is specific to OPCs upon cell cycle exit and mature oligodendrocytes (Suppl. Fig. 1). We generated *Olig2cre, APC*^{fl/fl} mice, which resulted in ~85% efficiency of APC protein loss in OPCs and ~15% "escapees" that failed to delete *APC* (Suppl. Fig. 2). While numbers of OPCs were normal in brain and spinal cord (Suppl. Fig. 2b), *APC*-deficient OPCs showed irreversible maturation arrest and failed to express myelin regulatory factor (Fig. 2a)(19), or the mature oligodendrocyte markers *Cnp, Mag* (Fig. 2a), *Mal, Plp* and *Fa2h* (Suppl. Fig. 2b, Suppl. Fig. 3b). *APC*-deficient OPCs expressed greatly elevated levels of the Wnt transcriptional target *Axin2* (Fig. 2b, Suppl. Fig. 4). Only "escapees" of *Olig2-cre* activity that expressed APC were capable of differentiation to NOGO-A-positive mature oligodendrocytes (Suppl. Fig. 3a).

High-activity Wnt signaling in colon cancer reaches a threshold to elicit the expression of LEF1, a reported target of the Wnt pathway, and involves a subsequent switch from a TCF4- β -catenin to a LEF1- β -catenin complex (Fig. 3a; (4)). Once activated, LEF1 participates in a feed-forward loop driving its own expression, which is critical for the high-activity state. As shown (Fig. 3b), Lef1 is not expressed during normal oligodendrocyte development; however, we observed robust Lef1 expression in the corpus callosum and spinal cord of *Olig2-cre, APC*^{fl/fl} animals, suggesting a high Wnt signaling threshold had been reached. Such *APC*-deficient OPCs expressed high levels of *Axin2* and remained permanently undifferentiated (Fig. 2b, Suppl. Fig. 5), in keeping with other recent findings (20). Conversely, *Axin2* expression was undetectable in *APC*-intact, Tcf4-positive OPCs (Fig. 3b). A fascinating finding is that such OPCs express the high activity markers SP5, Dusp4 and Ets2 (Fig. 4a and 4b). However, unlike Wnt-driven cancers of gut and hematopoetic systems, which respond to mitogenic Wnt signaling, we did not observe hyperproliferation of OPCs or glioma (Suppl. Fig. 6). In summary, high Wnt tone in OPCs is marked by the expression of Lef1 and acts purely to block differentiation (Fig. 3a).

SP5 is a Wnt target that inhibits myelin gene expression

We investigated *SP5*, encoding a zinc finger type transcription factor of the SP/KLF family, which was the most highly upregulated transcription factor in Wnt-activated OPCs. We first asked whether *SP5* was a direct Wnt target. As shown (Fig. 5a), Tcf4 binds to consensus binding sites at the *SP5* locus in OPCs as shown by chromatin immunoprecipitation (ChIP). *SP5* luciferase reporter gene activation was observed in the presence of β -catenin and endogenous Tcf4 in OPCs, and was inhibited by dominant-negative Tcf4 (Fig. 5b). The related factor SP1 binds to the *myelin basic protein (MBP)* enhancer to activate expression (21). As shown (Fig. 5d, Suppl. Fig. 7), SP5 directly bound multiple myelin gene loci in OPCs, but not mature oligodendrocytes while the converse was the case for SP1, which showed preferential binding at myelin genes in mature oligodendrocytes. SP5 expression

inhibited baseline *MBP* levels, while SP1-induced activation of *MB*P enhancer activity (Fig. 5c). These findings indicate direct activation of *SP5* by Wnt signaling within OPCs, and that SP5 proteins inhibit myelin gene expression through competition with SP1.

SP5-null animals are viable without known gut or white matter abnormalities (22). Consistent with this, we found that oligodendrocyte development proceeds normally in $SP5^{-/-}$ animals (Fig. 5e, Suppl. Fig. 8). We next used focal injection of lysolecithin into adult spinal cord white matter to assess kinetics of remyelination. Compared to low *SP5* expression levels in developing WT OPCs (Fig. 4a), we observed increased *SP5* expression levels in OPCs during remyelination (Fig. 5f). Strikingly, $SP5^{-/-}$ animals showed acceleration of OPC differentiation during remyelination and precocious expression of myelin genes *Mag, Cnp, Fa2h, Mbp* and *Plp* at 7-days post-lesion (dpl)(Fig. 5f and 5g, Suppl. Fig. 9). These findings indicate that *SP5* is a direct Wnt target and the first transcription factor identified whose function is required to delay kinetics of normal remyelination. Further, they suggest remyelination achieves an "intermediate" Wnt signaling state characterized by increased expression of *SP5* and requirement for *SP5*-function.

OPCs in human WMI express multiple novel genes in common with colon cancer

Subcortical WMI occurs in term infants that suffer severe hypoxic-ischemic encephalopathy (HIE; Fig. 6a). As shown (Fig. 6b–d, and g), immature oligodendrocyte lineage cells in such lesions expressed markers of high activity Wnt signaling (*i.e.*, LEF1, SP5, ETS2, DUSP4, RNF43). Surprisingly, such OPCs (Olig2+, Nkx2.2+) that co-expressed LEF1 and SP5 (Fig. 6c and d) in these lesions were uniformly APC-negative (n = 300 Lef1+ cells, three independent HIE cases)(Fig. 6e), suggesting deficient Wnt repressor tone. Indeed, RNF43 was expressed in cells with the simple bipolar morphology of OPCs in neonatal WMI (Fig. 6f), and in a chronic white matter lesion from a 12-year old child with CP (Fig. 6h), consistent with the possibility of a long-standing high-activity Wnt signaling state.

Loss of Wnt repressor tone determines transition to high activity signaling

Given these findings, we further investigated the function of Wnt repressor tone during remyelination, and compared Lef1 expression in animals that were either wild type, Axin2-/ – or Olig2-cre, APCfl/fl (Fig. 7a). Whilst Axin2 is a direct target of the Wnt pathway (Fig. 2b), Axin2-/– mice also have a hyperactive Wnt tone, as the Axin2 protein is a known repressor of the pathway through functions in the β –catenin degradation complex (16, 23). In the lysolecithin gliotoxic model, white matter oligodendrocytes are initially killed off and must be replaced by newly generated OPCs. As expected, based on findings from development (Fig. 2), APC-null oligodendrocytes show a permanent block in remyelination with persistence of Lef1 and Axin2 expression for as long as 90 dpl (Fig. 7c, Suppl. Fig. 10). Moreover, we observed that numbers of Lef1+ cells were increased in lesions inversely to Wnt repressor tone (Fig. 7b), suggesting that the percentage of Lef1+/Olig2+ OPCs comprises an index of Wnt activity levels.

To assess whether Wnt repressor function regulates neonatal WMI, we tested effects of mild/chronic neonatal hypoxemia (10% FiO2, P3–P7) on white matter volume throughout the brain. In wild type mice, this treatment causes only transient delay of myelination at P11

(data not shown) with full recovery of white matter volume by P40 (Fig. 7d and e). In contrast, *Axin2*–/– mice demonstrated selective white matter volume deficits, while grey matter volume was not significantly affected. Together these findings suggest that a deficient Wnt repressor tone and hypoxia is sufficient for permanent effects on white matter.

Discussion

Pathological Wnt signaling in cancers of colon and blood are associated with a high-activity state in genetically predisposed cells. We demonstrate for the first time that various Wnt pathway activity states can exist in oligodendrocyte lineage, and distinguish the normal process of myelination from pathological OPC differentiation block, in a dose-dependent fashion. We show that developing OPCs require APC function to prevent transition to a high-activity Wnt signaling state that permanently blocks their differentiation. Considering the requirement for APC in both colonic stem cells and OPCs, it is possible that individuals with familial adenomatous polyposis (FAP) syndrome, who carry heterozygous loss-offunction APC mutations, who are reported to suffer from intellectual disability (24), might be prone to WMI. OPCs with high-activity Wnt signaling express many known markers in common with colon cancer including Lef1, SP5, Ets2, Rnf43 (4, 8, 25, 26) as well as newly identified Dusp4 (this study). Further studies are needed to define the tissue-specific functions of such putative pathway components downstream of Wnt signaling in OPCs. As one example, we found that SP5 is a direct Tcf4 target, which acts in a competitive balance with the related transcription factor SP1 for binding at mature myelin gene loci, and is essential to inhibit kinetics of OPC differentiation after demyelinating injury.

Akin to the colon, high activity Wnt signaling in OPCs reaches the threshold to induce the expression of Lef1, with a subsequent switch from a TCF4- β -catenin to a LEF1- β -catenin dependent complex. We show that remyelination in a simple murine model and wild type genetic background does not result in high activity Wnt signaling, as Lef1-expressing OPCs are not observed. Nor did we observe Lef1 expression when other factors such as hypoxia or LPS-induced inflammation were added to gliotoxic injury (Fig. 7a and data not shown). In contrast, OPCs that carry heterozygous or homozygous mutations of *Axin2* or *APC* repressor genes show Lef1 expression in OPCs in a dose dependent manner. Thus, Wnt repressor tone is the critical factor that regulates transition to a high activity state after injury. Indeed, *Axin2*-/- mice subjected to mild hypoxia develop fixed deficits in white matter volume at P40 in dramatic contrast to wild type controls that show no injury whatsoever.

Genetic alterations in Wnt pathway repressors lead to certain cancers as well as noncancerous pathology (*e.g.*, bone hypertrophy) (27). Surprisingly, we found that OPCs in human neonatal HIE (*i.e.*, a non-genetic lesion) expressed high-activity markers similar to colon cancer, suggesting a parallel state of pathological Wnt signaling. This pathological Wnt tone in OPCs in human HIE goes some way to explaining why human lesions may fail to myelinate. However, further investigation is needed to determine how this high-activity state is achieved. One possibility is that hypoxic and/or inflammatory factors result in epigenetic dysregulation of Wnt repressor tone. In this regard, it is noteworthy that OPCs with high-Wnt tone consistently lacked APC expression. APC might be regulated at the transcriptional, translational level (28), or could be mis-localized within OPCs (29) in the

setting of injury. For example, hypoxia in colon cancer cell lines affects APC levels via a direct effect of hypoxia inducible factors (HIFs) at the APC promoter (30). Secondly, several previous studies indicate that epigenetic silencing of Wnt repressors in certain colon cancers may contribute to disease progression (31, 32), which could represent a pre-cancerous 'kindling state' that selects for somatic mutations. Further research is needed to establish epigenetic mechanisms that mitigate against Wnt repressor tone in human WMI and the extent to which a high-activity Wnt signaling state exists in other non-genetically-based human diseases.

Methods

Human HIE neonatal tissue and human colon cancer tissue

All human HIE tissue was collected in accordance with guidelines established by the University of California San Francisco Committee on Human Research (H11170-19113-07) as previously described (16, Fancy et al., *Nat Neurosci* 14, 1009 (2011)). Immediately after procurement, all brains were immersed in phosphate buffered saline with 4% paraformaldehyde for three days. On day 3, the brain was cut in the coronal plane at the level of the Mamillary Body and immersed in fresh 4% paraformaldehyde/PBS for an additional three days. Post fixation, all tissue samples were equilibrated in PBS with 30% sucrose for at least 2 days. Following sucrose equilibration, tissue was placed into molds and embedded with OCT for 30 minutes at room temperature followed by freezing in dry icechilled ethanol. UCSF neuropathology staff performed brain dissection and its evaluation. The diagnosis of hypoxic ischemic encephalopathy (HIE) requires clinical and pathological correlation; no widely accepted diagnostic criteria is present for the pathological diagnosis of HIE. None of the HIE cases evaluated in this study were identical but did show consistent evidence of diffuse white matter gliosis, as evaluated by the qualitative increase in the number of GFAP positive cells in addition to the increased intensity of GFAP staining.

Human colon cancer samples were from US Biomax, T055 colon cancer tissue array ($4 \times (2 \times 3)$ array) with 6 cases/24 cores per slide.

Comparison of transcripts in Wnt-activated OPCs with human colon carcinomas

We retrieved log(2) fold-change expression values from a previous study using Olig2-Cre/DA-Cat mice (15, Fancy et al., *Genes Dev* 23, 1571 (2009)) and another on colon cancer RNA profiling (3, *Nature* 487, 330 (2012)). Transcripts elevated in either data set or with a positive normalized "domain knowledge score" (DKS) were selected (Suppl Table 1). Normalized DKS was defined as the ratio of the number of articles in PubMed supporting upregulation vs. downregulation of each transcript. The PubMed query terms for upregulation were: ("elevated expression" OR "upregulated" OR "increase" OR "overexpression" OR "highly expressed" OR "enhanced expression") AND ("colorectal" OR "colon" OR "rectal"). The PubMed query terms for downregulation were: ("reduced expression" OR "downregulated" OR "decrease") AND ("colorectal" OR "rectal").

Mice

Animal husbandry and procedures were performed according to UCSF guidelines under IACUC approved protocols.

APC-fl/fl—A conditional (floxed) allele of APC provided by Dr. R. Fodde (Leiden University) (Robanus-Maandag, E.C., et al., *Carcinogenesis*, 31(5): p. 946–52 (2010)). Through intercrosses with an early Olig2 pan lineage cre-recombinase we can achieve conditional knockout of APC in OPCs.

SP5-LacZ—The SP5-LacZ mouse has been described previously (22, Harrison et al., *Developmental biology* 227, 358 (2000)) and has a LacZ knocked into the SP5 locus.

Axin2-LacZ—The Axin2-LacZ mouse has been described previously (Lustig, B, et al. *Mol Cell Biol* 22:1184–1193 (2002)). The LacZ insert mimics the expression pattern of Axin2 but does not lead to a detectable phenotype in the heterozygous state. In the homozygous state, this effectively acts as an Axin2 null animal.

Olig2-tva-cre—A multi-functional mouse line was constructed previously (Schüller, U, et al. *Cancer Cell* 14:123–134 (2008)) by inserting tva, an avian-specific retroviral receptor, and an IRES-cre recombinase cassette into the endogenous Olig2 locus by homologous recombination. This Olig2-cre allowed for cre mediated activity in oligodendrocyte lineage cells.

Immunohistochemistry and antibodies

The primary antibodies used were as follows: APC (1:100 mouse monoclonal OP80, Calbiochem), Lef1 (1:200 Rabbit mAb C12A5, Cell Signaling), anti-human Lef1 (1:100 rabbit polyclonal Ab22884, Abcam), RNF43 (1:200 rabbit polyclonal Ab84125, Abcam), ETS2 (1:200 chicken polyclonal Ab17901, Abcam), DUSP4 (1:200 rabbit polyclonal Ab60229, Abcam), SP5 (1:200 rabbit polyclonal Ab36593, Abcam), Tcf4 (1:100 mouse monoclonal 6H5-3, Upstate), Olig2 (1:3000 rabbit polyclonal from CD Stiles, Harvard), Nkx2.2 (1:100 mouse monoclonal, Developmental Studies Hybridoma Bank), PDGFRa (1:200 rat 558774, BD Biosciences), NOGO-A (1:200 rabbit, Millipore), and GFAP (1:500 mouse, Sigma).

Murine lysolecithin focal demyelination model

Demyelinated lesions were produced in the ventrolateral or dorsal spinal cord white matter of 8–10 week old *Axin2 null, Olig2cre:APCfl/fl* and wild-type littermate mice. Anaesthesia was induced and maintained with inhalational isoflurane/oxygen, supplemented with 0.05 ml of buprenorphine (Vetergesic 0.05 mg/ml) given subcutaneously. Having exposed the spinal vertebrae at the level of T12/T13, tissue was cleared overlying the intervertebral space, and the dura was pierced with a dental needle just lateral to midline. A Hamilton needle was advanced through the pierced dura at an angle of 45° and 0.5µl 1% lysolecithin (Lα-lysophosphatidylcholine) was injected into the ventrolateral white matter.

Murine developmental Chronic Hypoxia and statistics

Chronic hypoxic rearing was performed as previously described (Weiss et al., *Experimental Neurology*, 189: pp. 141–149 (2004)). Briefly, C57/B6 litters were culled to a size of 8 pups, cofostered with a CD1 dam and reared from P3 to P11 in a ventilated hypoxia chamber set to 10% 02 (A-Chamber, Biospeherix, Ltd. Lacona, NY). Following hypoxic rearing animals were returned to normal housing, weened at P28, and sacrificed at P40 as were normoxic controls. Statistical significance for the interaction effect of chronic hypoxic rearing and Axin2 deletion was determined by Two-Way ANOVA tests using the StatPlus software plugin (AnalystSoft) for Microsoft Excel. P-values of less than 0.05 are reported as significant.

Cell counts and statistical measures

For all cell counts, at least four animals per genotype were used for each time point indicated. Cells were counted on three or more non-adjacent sections per mouse and presented as an average +/- standard deviation. Statistical comparisons of cell counts were established using Student t test. No statistical methods were used to pre-determine sample sizes but our sample sizes are similar to those reported in previous publications (15, Fancy et al., *Genes Dev* 23, 1571 (2009)(16, Fancy et al., *Nat Neurosci* 14, 1009 (2011). Data collection and analysis were performed blind to the conditions of the experiment. Data were collected and processed randomly, and animals were randomly assigned to treatment groups.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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Fig. 1.

b

Common markers of high activity Wnt signaling across diverse tissue types. (a) Schematic to show strategy for comparison of mRNA transcripts upregulated in Wnt-activated OPCs (Olig2cre/DA-Cat)(15) with those upregulated in human colon carcinomas (3). (b) The markers LEF1, SP5, RNF43, ETS2 and DUSP4 are expressed in human colon adenocarcinoma. Scale bar represents 15 µm.



Fig. 2.

High activity Wnt signaling causes permanent OPC maturation arrest. (a) OPCs lacking APC fail to undergo a differentiation program during development and fail to express mRNAs for multiple mature oligodendrocyte markers, including *MRF*, *CNPase*, and *MAG*, as seen at postnatal day 15 (P15) in the spinal cord (SC) of *Olig2-cre*, $APC^{fl/fl}$ mice. Scale bar represents 100 µm. (b) *APC*-deficient OPCs expressed greatly elevated levels of the Wnt transcriptional target *Axin2* in P9 SC (which remains elevated at P120), and remain persistently undifferentiated as late as P650. Scale bar represents 300 µm (left panel), 10 µm (right panels).



Fig. 3.

The switch to high-activity Wnt signaling in colon cancer (4) involves a switch from a TCF4- β -catenin to a LEF1- β -catenin complex (a). Once activated, LEF1 (absent in normal colon) participates in a feed-forward loop driving its own expression. (b) Lef1 is not expressed during normal oligodendrocyte development in *Olig2-cre, APC*^{fl/+} heterozygous mice (which have lost one allele of APC in oligodendrocyte lineage), but there is robust expression of *Lef1* mRNA in the corpus callosum (CC)(Scale bar represents 70 µm) and Lef1 protein in spinal cord (SC)(Scale bar represents 30 µm) of *Olig2-cre, APC*^{fl/fl} animals at postnatal day 15 (P15) and P9 respectively, and these cells expressed high levels of *Axin2* (*black arrows inset*). Conversely, *Axin2* was not detected in the *APC*-intact, Tcf4-positive OPCs (arrowheads)(Scale bar represents 10 µm).



Fig. 4.

Novel markers of high activity Wnt signaling in OPCs. (a) OPCs in *Olig2-cre, APC*^{fl/fl} spinal cord at P9 and P15 express the high threshold Wnt signaling markers SP5, Dusp4 and Ets2, whereas these are undetectable in mice with heterozygous loss of one APC allele in oligodendrocyte lineage. Scale bar represents 60 μ m. (b) High threshold Wnt signaling markers are expressed in Nkx2.2+ and Olig2+ OPCs in SC and corpus callosum (CC) of *Olig2-cre, APC*^{fl/fl} mice. Scale bar represents 15 μ m.



Fig. 5.

Transcription factor SP5 is a Wnt target that inhibits mature oligodendrocyte gene expression. SP5 was the most highly upregulated transcription factor in Wnt-activated OPCs. (a) Sp5 is a target of the Wnt pathway in OPCs in vitro. There are predicted Tcf4 binding sites at the SP5 promoter locus, and ChIP demonstrates Tcf4 binding at these predicted sites in OPCs. (b) Sp5 transcription is increased with Wnt pathway activation, as SP5 luciferase reporter gene activation was observed in the presence of β -catenin (dominant active SY33) and endogenous Tcf4 in OPCs, and was inhibited by transfection of dominantnegative Tcf4 (dnTCF4) (t test * p < 0.05). (c) *MBP* luciferase reporter gene activation in vitro in OPCs is inhibited by transfection with SP5, and this also inhibited SP1-induced activation of *MBP* enhancer activity (*t* test * p < 0.05, ** p < 0.01). (d) ChIP showed that SP5 directly bound multiple myelin gene promoter loci (MBP, MAG, MAL, FA2H, CNPase) in OPCs but not mature oligodendrocytes while the converse was observed for SP1. (e) Oligodendrocyte development proceeds normally in $SP5^{-/-}$ animals with normal expression of CNPase and MBP mRNA at P1 and P9 respectively in the SC, and MBP protein in SC and CC. Scale bar represents 100 µm. (f) SP5 expression is seen in a subset of Olig2+ cells during wildtype mouse remyelination at 7dpl (Scale bar represents 10 μ m). SP5^{-/-} animals showed acceleration of OPC differentiation in the mouse model of remyelination using focal injection of lysolecithin into adult spinal cord white matter, with significant increases in number of cells expressing PLP mRNA at 7 days post lesioning (7dpl) and 14dpl (t test * p = 0.0002, ** p = 0.04, n=4). (g) SP5^{-/-} animals showed acceleration of OPC differentiation during remyelination and precocious expression of myelin genes Plp, Cnp, Mag, Mbp and Fa2h at 7dpl compared to wildtype (WT) animals. Scale bar represents 100 µm. Error bars represent s.d.

Fig. 6.

OPCs in human neonatal WMI express multiple genes in common with colon cancer. (a) Subcortical WMI occurs in human term infants that suffer severe hypoxic-ischemic encephalopathy (HIE). (b) High threshold Wnt signaling marker LEF1 is expressed in a subset of Olig2+ oligodendrocyte lineage cells in human HIE. (c) A proportion of immature OPCs (Nkx2.2+) co-expressed LEF1 (c) and SP5 (d) in these lesions and were *always* APC-negative (e). (g) Immature OPCs in such lesions expressed additional markers of high threshold Wnt signaling (ETS2, DUSP4). RNF43 was expressed in cells with the simple bipolar morphology of OPCs in neonatal WMI (f), and in a chronic WMI lesion from a 12-year old child with cerebral palsy (h). LEF1 and SP5 were not expressed by either GFAP+

astrocytes (i) or inflammatory cells expressing CD45 or CD24 markers (j) in lesions of human HIE. Scale bars represent 10 $\mu m.$

Fig. 7.

Loss of Wnt repressor tone determines transition to pathological Wnt signaling. (a) High threshold wnt marker Lef1 is expressed at very low levels in wildtype (WT) mouse remyelination (lysolethicin focal injection spinal cord)(Scale bar represents 100 μ m) at 14dpl and is not increased by intraperitoneal injection of LPS during remyelination, but Lef1 expression increased very significantly in lesions inversely to wnt tone in *Axin2*–/– or *Olig2-cre*, *APCfl/fl* mice (a and b). (c) *APC*-null oligodendrocytes show a permanent block in differentiation to a remyelinating stage with persistence of Lef1 and *Axin2* expression at 90 days after injury (90dpl). Scale bar represents 10 μ m. (d) Wnt repressor function regulates neonatal WMI, as demonstrated using mild/chronic neonatal hypoxemia in

developing mice (10% FiO2, postnatal day 3–7). Hypoxia in wildtype mice (WT HX) causes only transient delay of myelination at P11 (data not shown) with full recovery of corpus callosum thickness (d) and striatal thickness (e) to wildtype normoxic levels (WT NX) by P40 (Scale bars represent 50 μ m). *Axin2–/–* mice kept in normoxic conditions (Axin2 NX) have similar white matter volumes to WT animals at P40. *Axin2–/–* mice kept in hypoxic conditions (Axin2 HX) demonstrate significantly reduced corpus callosum thickness (2 way ANOVA *p = 0.023, n=4) and striatum white matter volumes (2 way ANOVA *p = 0.0009, n=4) at P40 compared with WT HX. Error bars represent s.e.m.