The role of the Wnt signaling pathway in cancer stem cells: prospects for drug development

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Abstract: Cancer stem cells (CSCs), also known as tumor initiating cells are now considered to be the root cause of most if not all cancers, evading treatment and giving rise to disease relapse. They have become a central focus in new drug development. Prospective identification, understanding the key pathways that maintain CSCs, and being able to target CSCs, particularly if the normal stem cell population could be spared, could offer an incredible therapeutic advantage. The Wnt signaling cascade is critically important in stem cell biology, both in homeostatic maintenance of tissues and organs through their respective somatic stem cells and in the CSC/tumor initiating cell population. Aberrant Wnt signaling is associated with a wide array of tumor types. Therefore, the ability to safely target the Wnt signaling pathway offers enormous promise to target CSCs. However, just like the sword of Damocles, significant risks and concerns regarding targeting such a critical pathway in normal stem cell maintenance and tissue homeostasis remain ever present. With this in mind, we review recent efforts in modulating the Wnt signaling cascade and critically analyze therapeutic approaches at various stages of development.

Keywords: β-catenin, CBP, p300, wnt inhibition

Introduction

Drug resistance, disease relapse, and metastasis constitute the central challenges in the management of advanced malignancies. Recently, cancer initiation, metastasis, and disease progression have been attributed to newly discovered subpopulations of self-renewing, highly tumorigenic, drug-resistant tumor cells termed cancer stem cells (CSCs), also known as tumor initiating cells (TICs). In many ways, CSCs behave very similarly to their normal counterparts, the somatic stem cells (SSCs), in that they have the ability to both self-renew and also to proceed on to more differentiated cell types. SSCs reside in specialized niches within tissues or organs (eg, hematopoietic stem cells, neuronal stem cells, and intestinal stem cells) and are critical for both normal tissue homeostasis and regeneration after injury. A major focus in cancer research over the past decade has been to both prospectively identify CSCs and, even more critically, to develop therapeutic strategies to safely eliminate this cell population without deleterious effects to the normal SSC populations.

A critical hurdle to safely accomplish this goal is the identification of key mechanisms that distinguish the control of self-renewal and proliferation of CSCs from their normal endogenous SSC counterparts. Not surprisingly, the same evolutionarily conserved signaling pathways that govern embryonic development are also critical to control the behavior of both normal somatic stem cells as well as cancer stem cells. The Wnt/β-catenin, Hedgehog, and Notch pathways have all
been implicated in stem cell and cancer stem cell biology. In this review, we will focus on the role of the Wnt signaling cascade in CSCs and the prospects for safely and effectively targeting this cascade to eliminate the CSC population in cancer.

Cancer stem cells and their role in tumorigenesis

Almost 150 years ago, Cohnheim proposed the concept that cancer might arise from a rare population of cells with stem cell–like properties. Today, increasing evidence has confirmed the existence of a small subgroup of cells in cancer, termed CSCs or, alternatively, TICs. The presence of CSCs has forced a paradigm shift from the earlier model of tumor homogeneity toward one of tumor hierarchy, where CSCs play a critical role. The cancer stem cell concept postulates that the bulk of a tumor consists of rapidly proliferating and differentiated (albeit aberrantly or only partially differentiated) cells, with a small population of CSCs that provide for the long-term maintenance of the tumor. These cells are able to self-renew, actively express telomerase, and activate antiapoptotic and multidrug resistance pathways. CSCs are relatively quiescent but can give rise to rapidly dividing progeny (so called transient amplifying cells), which form the bulk of tumor cells. Endowed with these characteristics, CSCs are thought to be responsible for tumor initiation, progression, and relapse, as well as metastasis and drug resistance. Supporting evidence exists that a stem-like signature contributes to aggressiveness and is related to poor outcome. Although CSCs resemble tissue stem cells in several characteristics, such as self-renewal and differentiation potential, it has been pointed out that the term “cancer stem cell” does not necessarily refer to the cell of origin, but can also refer to more differentiated cells that have acquired stem-like properties. Despite a still existing although decreasing controversy regarding the CSCs hypothesis, it is clear that distinct cancer cell populations have enhanced tumorigenic capacity compared with bulk tumor cells. Findings of cancer cells with enhanced tumor initiating properties were initially reported in leukemia. Bruce and van der Gaag demonstrated that only a small subgroup of cells showed extensive proliferation in vivo and in vitro. In 1997, Bonnet and Dick first isolated CSCs (known as leukemic stem cells, or LSCs) from bulk acute myeloid leukemia cells. Leukemic stem cells maintained or reacquired the ability to proliferate indefinitely, while losing the ability to properly differentiate. Subsequently, over the past decade, a large number of studies have identified CSCs in multiple solid tumors, including brain tumors, melanoma, and breast, liver, pancreatic, and colon cancer.

Wnt signaling in embryonic development and homeostasis

The Wnt/β-catenin pathway initiates a signaling cascade critical in both normal embryonic development and throughout the life of the organism in virtually every tissue and organ system. It is an enormously complex and ancient pathway that dates back to the first anaerobic metazoans. In addition to classical “canonical” Wnt activation of β-catenin/T-cell factor (TCF) transcriptional complexes, Wnt proteins can elicit a variety of alternative responses, often grouped together as “noncanonical” Wnt signaling. Wnts are secreted cysteine-rich glycoproteins that act as short-range ligands to locally activate receptor-mediated signaling pathways. The hallmark of this pathway is that it activates the transcriptional role of the multifunctional protein β-catenin. The key mediator of Wnt signaling, the armadillo protein β-catenin, dynamically localizes to multiple subcellular locations, including adherens junctions where it contributes to cell–cell contacts, the cytoplasm where β-catenin levels are tightly controlled, and the nucleus where in the canonical Wnt signaling pathway, β-catenin is involved in transcriptional regulation and chromatin modifications. The cytoplasmic pool of β-catenin is tightly regulated via phosphorylation by the “destruction complex” that includes glycogen synthase kinase 3β, casein kinase 1α, the scaffold protein Axin, and the tumor suppressor adenomatous polyposis coli (APC), among others (Figure 1A). In the absence of Wnt signaling, phosphorylation marks cytoplasmic β-catenin for ubiquitination and proteasomal degradation. A key step in the activation of Wnt target genes is the formation of a complex between β-catenin and members of the TCF/lymphoid enhancer factor family of transcription factors. To generate a transcriptionally active complex, TCF/β-catenin recruits the KAT3 transcriptional coactivator CREB-binding protein (CBP) (where CREB is an abbreviation for cAMP-response element binding protein), or its closely related homolog p300, as well as other components of the basal transcription machinery, to initiate transcription (Figure 1B).

The canonical, β-catenin-dependent Wnt signaling pathway plays crucial roles in the regulation of diverse cellular behaviors, including cell fate, proliferation, and survival. However, there exists a second noncanonical pathway, whose major effects apparently are β-catenin-independent in at least as much as that there is no apparent stabilization of cytoplasmic β-catenin. The noncanonical pathway is more
associated with differentiation, cell polarity, and migration (Figure 2) and can be further dissected into the Wnt/planar cell polarity and Wnt/calcium pathways, although these two noncanonical pathways are likely to intersect.\(^2,3\) Pathways affected by the noncanonical pathway include calcium-dependent and small GTPase-dependent signaling networks and the planar cell polarity signaling pathway, a pathway by which cells receive positional identity. Noncanonical signaling can be initiated by Wnt/frizzled receptor interactions without the help of Lrp5/6,\(^24\) or alternatively, receptor tyrosine kinase (RYK) and receptor tyrosine kinase-like orphan receptor (ROR) receptor tyrosine kinases can also act as Wnt receptors to activate β-catenin-independent signaling.\(^35\) β-catenin-independent signaling also regulates small GTPases, such as RHOA (Ras homolog gene family member A), RAC (Ras-related C3 botulinum toxin substrate) and Cdc42 (cell division control protein 42), in a dishevelled (Dsh)-dependent manner.\(^36\) Noncanonical Wnt activated calcium flux results in the activation of various kinase cascades, including protein kinase C, calcium/calmodulin-dependent protein kinase II, and JUN N-terminal kinase, which can activate NFAT (nuclear factor of activated T-cell) and AP-1-dependent transcription. Although dissection of the pathway into canonical and noncanonical may be convenient for discussion purposes, the reality is that these are interacting/intersecting pathways that can coordinately regulate and orchestrate complex processes during embryonic development, stem cell maintenance, tissue homeostasis, and wound healing. Wnt signaling plays critical roles in adults in the continuous processes of tissue homeostasis and regeneration of the hair and skin,\(^37\) maintenance of intestinal homeostasis,\(^38\) and hematopoiesis.\(^39,40\) Furthermore Wnt/β-catenin signaling is involved in liver and lung repair after injury\(^41-43\) and adult neurogenesis.\(^44\)

**The role of Wnt signaling in stem cells**

The Wnt signaling pathway has emerged as a pivotal player in the specification and maintenance of stem cell lineages and has been shown to have an important role in multiple stem cell compartments in a wide array of tissues and organs.\(^45-50\) The small intestine is organized into villi (apical) and crypts (basal). Intestinal stem cells reside in intestinal crypts\(^45\) and their maintenance and proliferation

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**Figure 1** Canonical Wnt signaling.

**Notes:** (A) Wnt off. In the absence of Wnt (Wingless) ligands, a multi-protein destruction complex in the cytoplasm consists of Axin-1 and its interacting partners tumor suppressor adenomatous polyposis coli (APC), glycogen synthase kinase 3β (GSK3β), and casein kinase 1α (CK1). The destruction complex degrades β-catenin by proteosomal degradation via phosphorylation and, thus, maintains low levels of β-catenin. (B) Wnt on. In the presence of Wnt ligand binding to the frizzled/Lrp5/6 (low density lipoprotein receptor-related proteins 5 or 6) receptors a negative regulator of the destruction complex, dishevelled (Dsh) is recruited leading to degradation of Axin and inactivation of GSK3β (glycogen synthase kinase 3β) and CK1α (casein kinase 1α), thereby inhibiting their interaction with other components of the destruction complex. As β-catenin accumulates in the cytoplasm, it translocates to the nucleus, where it forms a transcriptionally active complex with transcription factors of the T-cell factor (TCF)/lymphoid enhancer factor (LEF) family and coactivators, such as CBP (cyclic AMP response element-binding protein) and p300, driving the expression of Wnt target genes. Arrows indicate activation/induction; blunt ended lines indicate inhibition/blockade.

**Abbreviation:** APC, adenomatous polyposis coli.
is Wnt dependent.\textsuperscript{51} The loss of positive Wnt regulators, such as TCF4 or β-catenin, as well as the overexpression of negative Wnt regulators, such as Dickkopf1, dramatically decreases the proliferation capacity of this stem cell compartment.\textsuperscript{52,53} Two distinct intestinal stem cell populations have been described. The first population is made up of +4 label retaining cells,\textsuperscript{54} which are highly quiescent and are activated apparently only after injury.\textsuperscript{54} This population is characterized by the stem cell marker Bmi1.\textsuperscript{55} A second population is made up of the crypt basal columnar cells (CBCs), which are interspersed between Paneth cells and express the surface marker Lgr5.\textsuperscript{55,56} CBCs continuously cycle and are responsible for sustained tissue homeostasis. Lgr5 is a Wnt/β-catenin target gene, which can amplify Wnt signaling in CBCs via its R-spondin ligand.\textsuperscript{57} Paneth cells, are an important source for Wnt ligands (ie, Wnt3, 6, 9b),\textsuperscript{58} which appear crucial for the maintenance of intestinal stem cells.\textsuperscript{59} Supporting this notion is the fact that depletion of Paneth cells leads to a decrease in the number of intestinal stem cells.\textsuperscript{59} Wnt signaling is also critical for expression of the gene Sox9, which is important for Paneth cell lineage commitment.\textsuperscript{50,61}

In the hematopoietic system, Wnt3a has been implicated in self-renewal and proliferation.\textsuperscript{39,62} Regulation of hematopoietic stem/progenitors, as well as lineage commitment of progenitors during hematopoiesis is highly Wnt dependent.\textsuperscript{39,40} Expression of survivin, a member of the inhibitor of apoptosis protein family, is important during hematopoiesis and is prominently upregulated in CD34\textsuperscript{+} hematopoietic stem/progenitor cells upon growth factor treatment.\textsuperscript{63} Inducible deletion of survivin leads to loss of hematopoietic progenitors and bone marrow ablation, whereas heterozygous deletion of survivin leads to defects in erythropoiesis.\textsuperscript{63,64} We previously demonstrated that survivin is a Wnt/β-catenin/CBP dependent target gene in a variety of cancer cell types.\textsuperscript{65} More recently, we also

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**Figure 2** Noncanonical Wnt signaling. 

**Notes:** (A) Noncanonical Wnt/calcium pathway. Wnt ligand binding to frizzled receptors activates Dsh and trimeric G-proteins (G\textsubscript{α/β,γ}), leading to the generation of IP3 (inositol 1,4,5-triphosphate) and DAG2 (dialcylglycerol) by PLC (phospholipase C)-mediated conversion of PIP2 (phosphatidylinositol bisphosphate) and triggering release of calcium ions (Ca\textsuperscript{2+}). Subsequent calcium release activates CAMKKII (calcium/calmodulin-dependent kinase II), TAK-1 (TGF-β activated kinase 1) and NLK (nemo-like kinase). Calcium release also activates PKC (protein kinase C) and Cdc42 (cell division control protein 42), and, thereafter, actin cytoskeleton is rearranged. In addition, calcium release results in activation of calmodulin, calcineurin, and NFAT (nuclear factor of activated T-cell), which is critical for control of tissue polarity. Arrows indicate activation, blunt ended lines indicate inhibition/blockade. (B) Noncanonical Wnt/PCP (planar cell polarity) pathway. Wnt ligand binding to frizzled receptors leads to activation of dishevelled (Dsh), triggering stimulation of GTPases RAC (Ras-related C3 botulinum toxin substrate), Profilin and RHOA (Ras homolog gene family member A) through activation of DAAM1 (dishevelled associated activator of morphogenesis 1). Subsequently, actin cytoskeleton is rearranged. In addition, Dsh activates RAC and finally activates JNK (c-Jun N-terminal-kinase) and AP1, which has been implicated in cell migration. 

**Abbreviations:** RYK, receptor tyrosine kinase; ROR, receptor tyrosine kinase-like orphan receptor; ROCK, Rho-associated protein kinase; MAPKKK, mitogen-activated protein kinase kinase kinase; MAPKK, mitogen-activated protein kinase kinase; MAPK, mitogen-activated protein kinase; PKC, protein kinase C; CAMKII, calcineurin, calcium/calmodulin-dependent kinase II; TAK1, transforming growth factor-β-activated kinase 1; NLK, nemo-like kinase; NFAT, nuclear factor of activated T-cell; G\textsubscript{α/β,γ}, trimeric G-proteins; IP3, inositol 1,4,5-triphosphate; DAG, diacylglycerol; PLC, phospholipase C; PIP2, phosphatidylinositol bisphosphate; RAC, Ras-related C3 botulinum toxin substrate; Profilin, profilin; RHOA, Ras homolog gene family member A; DAAM1, dishevelled associated activator of morphogenesis 1; JNK, c-Jun N-terminal-kinase; AP1, activator protein 1.
**Figure 3** Schematic of Wnt inhibitors currently in clinical trials.

**Notes:** LGK974 is a porcupine inhibitor that inhibits Wnt posttranslational acylation. OMP-18R5 is a fully humanized monoclonal antibody that binds to multiple frizzled (Fzd) receptors. OMP-54F28 is an Fc fusion protein with Fzd8, which binds all Wnt ligands. PRI-724 binds specifically to the coactivator CBP blocking its interaction with β-catenin. Arrows indicate activation/induction, blunt ended lines indicate inhibition/blockade.

**Abbreviations:** Dsh, dishevelled; CK1α, casein kinase 1α; GSK3β, glycogen synthase kinase 3β; APC, adenomatous polyposis coli; TCF/LEF, T-cell factor/lymphoid enhancer factor; CBP, cyclic AMP response element-binding protein.

Demonstrated that survivin is critical in drug resistance in leukemia. We have demonstrated that inhibition of the Wnt pathway by disrupting the CBP/β-catenin interaction in pre-B acute lymphoblastic leukemia (ALL) represents a powerful mechanism to eradicate drug resistant subclones, which was associated with downregulation of survivin.

Wnt signaling has also been implicated in mammary gland development and cell transformation. Ectopic expression of ΔNβ-catenin or Wnt1 leads to ductal hyperplasia, while loss of function in β-catenin (using a dominant negative variant) has been shown to exert a negative effect on breast tissue development during pregnancy, in particular, lobuloalveolar proliferation. Overexpression of inhibitors (such as Axin) or loss of lymphoid enhancer factor 1 function inhibits mammary differentiation of precursor cells. The bilayered mammary epithelium consists of luminal cells (Ck8+, Muc1+) and basal cells (Ck5+, p63+). Of these two cell types, the basal cells have been shown to express both Lrp5 and 6, obligate canonical Wnt signaling receptors.

Ductal mammary stem cells comprise a sub-population of basal epithelial cells and are capable of regenerating cleared mammary fat pads. Knockout studies for Lrp5 and loss of function mutations for the Lrp6 receptor showed significantly reduced activity in this cell compartment and impaired gland branching, suggesting impaired stem cell function. Wnt signaling has also been implicated in neuronal development and neuronal stem cell cells.

**The role of Wnt signaling in cancer stem cells**

Considering the importance of the Wnt pathway in stem cell biology, it is not surprising that aberrant Wnt signaling has been implicated in the tumorigenic potential of stem cells. A typical approach to prospectively identify putative cancer stem cells is via cell surface markers; however, these are also expressed on normal somatic stem cells. Many of these markers are in fact direct Wnt target genes (including LGR5/GPR49, CD44, CD24, CD133, ABC...).
cassette genes. The first report for the existence of CSCs in solid tumors emerged from studies in breast cancer by Al Hajj et al., who showed that cells that are CD44<sup>high</sup>CD24<sup>low</sup> possess tumor initiating capacity. It has long been known that misexpression of Wnt ligands induces mammary adenocarcinomas. A role for the Wnt signaling pathway in glioblastoma stem cells has also recently been described. Another hallmark of aggressive breast cancers is an enrichment in an epithelial–mesenchymal transition (EMT)-like gene expression signature. For example, Wnt1 has been shown to upregulate Twist, thereby favoring EMT-like processes in breast cancer cells. Loss of E-Cadherin associated β-catenin in breast cancer cells leads to disruption in cell polarity resulting in an epithelial-mesenchymal transition, a CSC-like phenotype with a significant increase in the CD44<sup>high</sup>, CD24<sup>low</sup> population and increased Wnt signaling. The process of EMT has also been associated with activated β-catenin signaling. Overexpression of the EMT inducing factors twist and snail (both putative Wnt target genes) increases the expression of CSCs markers. A connection between enhanced nuclear β-catenin signaling and EMT is consistent with the large number of β-catenin target genes (eg, S100A4, fibronectin, L1CAM, CD44, MMP7, and uPAR) whose expression is associated with invasion, migration, and metastasis. The cell surface protein CD133, is expressed by normal progenitor cells of the neural, hematopoietic, epithelial, and endothelial cell lineages. Recently, enrichment of CD133<sup>+</sup> cells in colorectal cancer samples has been shown to enrich for a population of CSCs/TICs. These cells also express high levels of nuclear β-catenin. Furthermore, Lrg5/GPR49 is overexpressed in the majority of colorectal tumors compared with normal control tissue. Several studies have revealed that MDR-1, ABCG2, ABCA3, and BRCP1 are expressed in stem/progenitor cells from multiple adult tissues and that they contribute to the side population phenotype of stem cells. The expression of these so-called multidrug resistance genes has been shown to be also associated with cancer stem cells and partially responsible for poor therapeutic responses. Wnt/β-catenin signaling appears to play an important role in ABCB1/MDR-1 transcription. Multiple putative TCF binding elements were identified in the ABCB1 promoter (−1813 to −275 bp). The side population assay has been utilized to identify rare drug resistant hematopoietic CSCs/TIC populations. Hematopoietic CSCs/TIC populations have been shown to be Wnt/β-catenin dependent. Furthermore, many Wnt signaling genes are upregulated in hematopoietic malignancies and epigenetic silencing of negative regulators of the Wnt signaling cascade is frequently associated with leukemias, including chronic myeloid leukemia.

**Wnt signaling pathway as a potential oncotarget**

Aberrant regulation of Wnt signaling has emerged as a recurrent theme in cancer biology. The discovery in 1991 that mutations in the tumor suppressor adenomatous polyposis coli (APC) were associated with the vast majority of sporadic colorectal cancers via aberrant activation of Wnt signaling provided significant impetus to attempt to therapeutically target this pathway. Germline defects in APC are the cause of familial adenomatous polyposis. Affected individuals develop hundreds of polyps in the large intestine at an early age and ultimately progress to colorectal cancer with 100% penetrance. Loss of function in both alleles of APC is required for tumorigenesis and is connected to the protein’s ability to regulate β-catenin protein stability as well as chromosomal stability. APC is the most frequently mutated gene in human cancers. Wnt pathway mutations are, however, not limited to colon cancer. Loss-of-function mutations in Axin have been found in hepatocellular carcinomas. Moreover, oncogenic β-catenin mutations, first described in colon cancer and melanoma, have also been found to occur in a wide variety of solid tumors including hepatocellular carcinomas, thyroid tumors, and ovarian endometrioid adenocarcinomas. Additionally, epigenetic silencing is frequently observed to alter levels of expression of Wnt/β-catenin pathway negative regulators. For example, methylation of genes that encode extracellular Wnt antagonists, such as secreted frizzled-related proteins, has been described in colon, breast, prostate, lung, and other cancers. Increased expression of Wnt ligands or effector proteins (eg, Dsh) has also been reported. Clearly, the ability to target the Wnt signaling pathway offers enormous promise as an oncological target. However, significant risks and concerns regarding targeting such a critical pathway in normal stem cell maintenance and tissue homeostasis are ever present.

**Current Wnt inhibiting molecules: small molecules and biologics**

Despite a wealth of information and significant investment in research and development, only recently have a few therapeutic agents that specifically target the Wnt pathway been
have been implicated as nonspecific inhibitors of the Wnt/β-catenin signaling pathway, although the mechanisms of action of these agents are not clear.149–157 Through screening a library of FDA-approved drugs, the anthelmintic agent pyrvinium was identified. This agent was shown to potentiate the activity of the casein kinase 1 alpha leading to enhanced degradation of β-catenin and the coactivator Pygo and thereby reduction of Wnt/β-catenin signaling.154

A number of molecularly targeted agents have been reported, which can be classified basically into several classes, ie, β-catenin/TCF-antagonists, PDZ (postsynaptic density protein 95, Drosophila disc large tumor suppressor, zonula occludens-1 protein),155 the domain of Dsh binders, and other mechanism-based inhibitors, principally enzymes (eg, kinases, tankyrases,156 Porcupine,157 and biologics). To date, most of these have only been evaluated preclinically and for several recent reviews the reader is referred to the following.6,114,158,159 Ongoing clinical trials of Wnt inhibitors/modulators are summarized in Table 2 and Figure 3. Recently, Novartis International AG, (Basel, Switzerland) initiated a Phase I trial of the Porcupine inhibitor LGK974 (NCT01351103)160 to treat a variety of malignancies (melanoma, breast cancer, and pancreatic adenocarcinoma) associated with aberrant Wnt signaling. Porcupine is a member of a family of O-acyltransferases that is apparently dedicated to Wnt posttranslational acylation.161 This trial has not been completed and to date no public information is available.

Two Wnt-targeting biology developed by OncoMed Pharmaceuticals, Redwood City, CA, USA have recently entered clinical trials. OMP-18R5 is a fully humanized monoclonal antibody that binds to multiple frizzled receptors.162 An open label Phase Ia study for solid tumors was recently completed (NCT01345201). The results of this trial were recently reported at the American Society of Clinical Oncology conference in June 2013.163 A total of 18 patients were treated in 5 dose escalation cohorts (0.5 and 1 mg/kg once per week; 0.5 mg/kg every two weeks; 1 and 2.5 mg/kg every three weeks). The most common related adverse events included grade 1 and 2 fatigue, vomiting, abdominal pain,
constipation, diarrhea, and nausea. The only related grades greater than or equal to 3 were dose limiting toxicities of grade 3 diarrhea and vomiting in one patient at 1 mg/kg/week. 1 patient at 0.5 mg/kg per week suffered a therapy-related bone fracture on day 110. Further clinical trials using OMP-18R5 in combination with other agents in solid tumors (NCT01957007) and breast cancer (NCT01973309) are ongoing. The second agent, OMP-54F28, is an Fc fusion protein with frizzled family receptor 8, which binds all Wnt ligands. A trial in solid tumors was initiated last year (NCT01608867) with the primary end point being safety.\(^\text{164}\) Potential deleterious effects of this agent on bone formation/turndover have been prospectively designed into the trial. Subjects will be monitored throughout the study for effects on bone density and turndover. Dose escalation studies in combination with other agents are currently ongoing in hepatocellular cancer, liver cancer (NCT02069145), ovarian cancer (NCT02092363), and pancreatic cancer (NCT02050178).

Our group used a Topflash reporter gene screen to identify inhibitors of Wnt signaling in SW480 colon carcinoma cells. This led to the identification of ICG-001 from a library of secondary structure mimetics.\(^\text{165}\) In this assay, ICG-001 had an IC50 value of 3 μM. We subsequently identified and validated, using a gain-of-function/loss-of-function strategy, that ICG-001 binds specifically and with high affinity (−1 nM) to the coactivator CBP, but importantly, not to its closely related homolog p300, despite the fact that these two coactivators are up to 93% identical, with even higher homology, at the amino acid level.\(^\text{165,166}\) PRI-724, a second generation specific CBP/catenin antagonist, developed by Prism Pharma Co., Ltd. Yokohama, Japan entered an open label Phase Ia safety study in subjects with solid tumors, where the expression of the biomarker survivin/BIRC5 was measured by immuno-magnetic RT-PCR in circulating tumor cells. Trial results were reported at the American Society of Clinical Oncology conference in June 2013 (NCT01302405).\(^\text{167}\) Eighteen patients were treated (dose escalation from 40-1,280 mg/m²/day) via continuous infusion for 7 days. PRI-724 had a low toxicity profile: one dose limiting toxicity of grade 3 hyperbilirubinemia was reported. Reported grade 2 adverse events were diarrhea (2 patients, 11%), bilirubin elevation (2 patients, 11%), hypophosphatemia (2 patients, 11%); nausea (1 patient, 6%), fatigue (1 patient, 6%), anorexia (1 patient, 6%), thrombocytopenia (1 patient, 6%), and alkaline phosphatase elevation (1 patient, 6%). There was no maximum tolerated dose at the doses tested. Three patients with colon cancer had stable disease for 8, 10, and 12 weeks. Down regulation of survivin/BIRC5 in circulating tumor cells was dose dependent.\(^\text{167}\) Additional trials with PRI-724 in myeloid malignancies (NCT01606579) and in combination with gemcitabine in pancreatic adenocarcinoma (NCT01764477) are ongoing.

### Development of novel Wnt inhibitors: challenges and prospects

More than 30 years after the groundbreaking discovery of Wnt signaling and extensive investigation into this fundamental and highly evolutionarily conserved pathway, there is still no FDA approved agent that specifically targets aberrant Wnt signaling in cancer. Very recently, a number of small molecules and biologics have entered human clinical trials. Despite exciting preclinical data in a variety of tumor models, it is still too early to know if any of these therapeutic agents will be efficacious with an acceptable safety profile. However, it is already clear that successfully targeting Wnt signaling in cancer will require a fine balancing act, whereby the “dark side” of Wnt signaling in cancer can be abrogated without interfering with the critical role of Wnt signaling in tissue homeostasis (eg, intestinal epithelium, blood, and bone) and repair. Numerous potential concerns arise in the development of therapeutic strategies that antagonize the Wnt pathway. Therapeutic agents that target critical developmental signal transduction pathways (eg, Wnt) are likely to have devastating effects on embryonic patterning. Further concerns about on-target toxicity include effects on intestinal stem cells, bone turnover, and hematopoiesis. For example, the Porcupine inhibitor LGK974 exhibited 63% tumor growth delay when administered at 3 mg/kg/day.\(^\text{168}\) However, at a dose of 20 mg/kg/day, significant loss of intestinal epithelium was observed. Concerns about potential deleterious effects on bone formation/turndover have been prospectively addressed in the OMP-54F28 trial design (NCT01608867), as all subjects receive 30 days of vitamin D3 and calcium carbonate after discontinuation of OMP-54F28 and are monitored during the study for effects on bone density and turnover. It remains to be seen if toxic side effects occur with inhibitors of Wnt signaling, and, if there are none, it would be critical to understand why.

Despite all of these potential concerns regarding targeting Wnt signaling in cancer, there is also tremendous excitement as our knowledge of this pathway continues to increase and our clinical experience with novel Wnt-targeting therapeutic agents expands.
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