

Wnt/ β -Catenin Signaling during Early Vertebrate Neural Development

David Brafman,¹ Karl Willert ²

¹ School of Biological and Health Systems Engineering, Arizona State University, Tempe, Arizona 85287

² Department of Cellular & Molecular Medicine, University of California San Diego, La Jolla, CA 92093-0695

Received 2 May 2017; revised 24 July 2017; accepted 9 August 2017

ABSTRACT: The vertebrate central nervous system (CNS) is comprised of vast number of distinct cell types arranged in a highly organized manner. This high degree of complexity is achieved by cellular communication, including direct cell-cell contact, cell-matrix interactions, and cell-growth factor signaling. Among the several developmental signals controlling the development of the CNS, Wnt proteins have emerged as particularly critical and, hence, have captivated the attention of many researchers. With Wnts' evolutionarily conserved function as primordial symmetry breaking signals, these proteins and their downstream effects are responsible for simultaneously establishing cellular diversity and tissue organization. With their expansive

repertoire of secreted agonists and antagonists, cell surface receptors, signaling cascades and downstream biological effects, Wnts are ideally suited to control the complex processes underlying vertebrate neural development. In this review, we will describe the mechanisms by which Wnts exert their potent effects on cells and tissues and highlight the many roles of Wnt signaling during neural development, starting from the initial induction of the neural plate, the subsequent patterning along the embryonic axes, to the intricately organized structure of the CNS. © 2017 Wiley Periodicals, Inc. *Develop Neurobiol* 77: 1239–1259, 2017

Keywords: Wnt; cell signaling; neural development; embryonic development

INTRODUCTION

The formation of the central nervous system (CNS) begins during gastrulation with the emergence of the ectoderm from the epiblast (human embryonic development days 14–18 [E14–E18]) (Sanes et al., 2012). During neural induction (E16–E18), naïve ectoderm becomes committed to the neural lineage as the neural plate forms along the dorsal midline of the embryo (Andoniadou and Martinez-Barbera, 2013; Kicheva and Briscoe, 2015). The neural plate continues to fold into the underlying mesoderm resulting in the formation of the neural groove surrounded on each side by the neural

folds (E18–E20; Andoniadou and Martinez-Barbera, 2013; Kicheva and Briscoe, 2015). As the neural groove elongates along the anterior-posterior axis the neural folds fuse to create the neural tube (E20–E24; Andoniadou and Martinez-Barbera, 2013; Kicheva and Briscoe, 2015). Concomitantly with the formation of neural tube, the neural crest arises from non-neuronal ectodermal cells along the lateral edge of the neural plate (Huang and Saint-Jeannet, 2004). The neural crest will eventually give rise to a myriad of cell populations, including the neural cells of the sensory, sympathetic, and parasympathetic nervous system as well as various craniofacial musculoskeletal cell types (Huang and Saint-Jeannet, 2004; Sanes et al., 2012; Liu and Cheung, 2016). By the fourth week of development, the neural tube becomes further subdivided into three primary vesicles—the prosencephalon (forebrain), the mesencephalon (midbrain), and the rhombencephalon (hindbrain, HB)—and the emerging spinal cord (SC;

Correspondence to: D. Brafman (David.Brafman@asu.edu) or K. Willert (kwillert@ucsd.edu).

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Published online 11 August 2017 in Wiley Online Library (wileyonlinelibrary.com).

DOI 10.1002/dneu.22517

Sanes et al., 2012). The prosencephalon will be further divided into the anterior telencephalon (which eventually matures into the cerebrum) and the posterior diencephalon (which will form the thalamic and hypothalamic regions; Wilson and Houart, 2004). The rhombencephalon further separates into eight defined regions called rhombomeres (Guthrie and Lumsden, 1991; Glover, 2001). The anterior rhombomeres (R1–R3), which comprise the metencephalon, will differentiate into the pons and cerebellum (Carlson, 2014). The posterior rhombomeres (R4–R8), which constitute the myelencephalon, will generate the structures of the medulla oblongata (Carlson, 2014).

A variety of signaling molecule pathways act in a tightly regulated spatial and temporal manner to instruct the specification, development, and maturation of the CNS. In particular, the Wnt signaling pathway plays critical roles in every aspect of CNS development. In this review, we will focus of the role of Wnt signaling in the earliest stages of neurodevelopment and patterning from gastrulation to approximately week 5 of human *in utero* development (Fig. 1). The mechanisms by which Wnt signaling regulates later CNS developmental processes including corticogenesis (Bielen and Houart, 2014; Inestrosa and Varela-Nallar, 2015; Fortress and Frick, 2016), neural crest emergence (De Calisto et al., 2005; Pegoraro and Monsoro-Burq, 2013; Mayor and Theveneau, 2014), eye morphogenesis (Cavodeassi, 2014; Fujimura, 2016), axon growth and guidance (Guan and Rao, 2003; Clark et al., 2012; Onishi et al., 2014; Petrova et al., 2014), synaptic formation and function (Sanes and Lichtman, 2001; Burden, 2002; Salinas, 2012; Dickins and Salinas, 2013; Poon et al., 2013; Stamatakou and Salinas, 2014), and adult neurogenesis (Bielen and Houart, 2014) have been reviewed extensively elsewhere.

Overview of Wnt Signaling

The importance of Wnt signaling in the biology of multicellular life was first appreciated in two independent lines of investigation: in a mouse model for breast cancer and, separately, in genetic analysis of *Drosophila* developmental mutants. Work by Nusse and Varmus demonstrated that proviral insertion of the mouse mammary tumor virus in a specific locus of the genome, termed *int1*, promoted tumorigenesis (Nusse and Varmus, 1982). A polyadenylated RNA transcribed from within this locus was expressed in tumors only, and subsequent studies (Nusse et al., 1984; Fung et al., 1985) showed this RNA to encode a protein with two of the key hallmarks of all Wnt proteins: a hydrophobic amino terminus signifying

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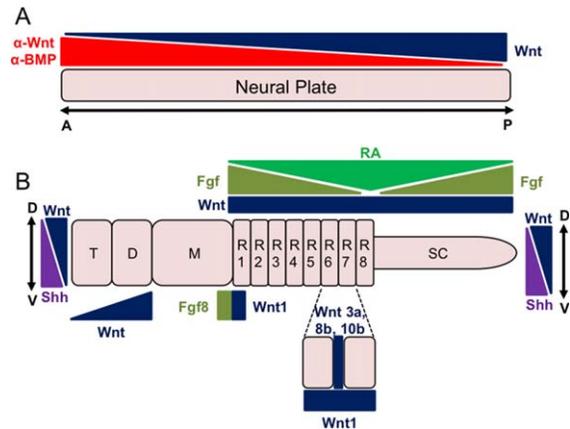


Figure 1 Summary of early of neural development events regulated by Wnt signaling. A: The neural plate initially emerges from the ectoderm through the action of BMP antagonists (α -BMPs). Opposing gradients of Wnt antagonists (α -Wnt) and Wnt ligands pattern the neural tube along the anterior-posterior (A/P) axis. B: As the neural tube continues to mature, it becomes further subdivided into primary vesicles (prosencephalon, mesencephalon [M], and the rhombencephalon) and the developing SC. The prosencephalon will be further divided into the telencephalon (T) and diencephalon (D). In addition, the rhombencephalon divides into eight rhombomeres (R1–R8). Wnt signaling along with other signaling molecules (e.g., FGFs, Shh, RA) regulates the further patterning and maturation of these regions along both the A/P as well as dorsal-ventral (D/V) axis. [Color figure can be viewed at wileyonlinelibrary.com]

targeting to the secretory pathway and an unusually high content of cysteine residues.

Predating these mouse studies, groundbreaking genetic screens in *Drosophila* led to the identification of a group of genes that controlled embryonic patterning (Nusslein-Volhard and Wieschaus, 1980). One of the groups of genes, called segment polarity genes, included *wingless* (*wg*), previously named after a hypomorphic allele that produced flies lacking wings (Sharma and Chopra, 1976). Molecular cloning of the *Wg* gene revealed it to be homologous to the mouse gene transcribed from the *int1* locus. With the recognition that *Wg* and *int1* represented prototypes for a much larger family of related genes, the mnemonic Wnt was coined (Nusse et al., 1991).

Wnt genes have been identified in all metazoan species, with the mammalian genome containing 19 distinct *Wnt* genes, some of which encode alternative transcripts. Because Wnt proteins and their signaling pathways govern a myriad of biological processes—from patterning of the fly embryo to cancer development in mammals—it may, at times, be difficult to discern a theme that underlies and unifies Wnt biology. Nonetheless, from an evolutionary perspective,

it is clear that Wnt proteins are pivotal symmetry breaking signals that establish the primary body axis during embryonic development (reviewed in Loh et al., 2016). Consistent with this universal role for Wnts in organismal development, and as we will discuss in this review, Wnt signaling regulates anterior-posterior and dorsal-ventral patterning of the developing CNS. The study of Wnt signaling has extended into virtually all aspects of biology and the reader is encouraged to visit the Wnt homepage (wnt.stanford.edu) for additional information on Wnt across the spectrum of developmental biology.

Wnt Proteins. Wnt genes encode secreted lipid-modified glycoproteins (Brown et al., 1987; Papkoff et al., 1987; Mason et al., 1992; Willert et al., 2003). A nearly invariant pattern of 22 cysteine residues, all of which are engaged in disulfide linkages, are essential for the proper folding of this growth factor into a two-domain hand-like structure, which was recently elucidated by X-ray crystallography (Janda et al., 2012). A covalently attached lipid extending from the tip of the “thumb” is critically important for Wnt function and renders it highly hydrophobic (Willert et al., 2003; Takada et al., 2006). The process of attaching this lipid, palmitoleic acid (a 16-carbon monounsaturated lipid), on a conserved Serine residue of the Wnt polypeptide chain, occurs in the endoplasmic reticulum and requires the activity of Porcupine (Porcn), an acyl-transferase (Hofmann, 2000; Tanaka et al., 2000). Since Porcn likely processes all Wnt proteins, blocking its function, either by mutation (Barrott et al., 2011; Biechele et al., 2011; Liu et al., 2012) or treatment with Porcn inhibitors (Chen et al., 2009; Liu et al., 2013; Proffitt et al., 2013), provides a powerful tool to interfere with all Wnt signaling activity and create an all-Wnt deficient state. Following processing by Porcn, Wnts transit through the secretory pathway and are eventually secreted from the cell, a process that requires an additional protein encoded by the *wntless* (Wls) gene (Banziger et al., 2006; Bartscherer et al., 2006; Herr and Basler, 2012).

The lipid renders the Wnt protein highly hydrophobic and, hence, insoluble in what is largely an aqueous environment. This feature serves to limit Wnt diffusion in the extracellular space, and for the majority of cases, Wnt proteins remain closely associated with the Wnt expressing cells. Of note, engineering a Wnt protein such that it remains tethered to the cell surface via a transmembrane domain does not significantly limit its signaling activity: flies expressing such an allele of *wg* are patterned normally and exhibit no major developmental defects other than a slight reduction in body size (Alexandre et al., 2014).

These observations indicate that initiation of a Wnt signal through receptor engagement involves close cell-cell proximity, as in the case of Notch signaling, rather than the action of a diffusible growth factor that can act at a long distance from its original site of secretion. Nonetheless, Wnt proteins, in particular Wg, have been observed at significant distances from Wnt-secreting cells, which may be mediated by transport through cells via a transcytosis-like process (Dierick and Bejsovec, 1998; Yamazaki et al., 2016), by long membrane protrusions, such as cytonemes or filipodia (Hsiung et al., 2005; Stanganello et al., 2015), or by exosomes (Korkut et al., 2009; Gross et al., 2012).

Wnt Receptors. The main class of Wnt receptors encoded by the Frizzled (Fzd) gene family is comprised of ten independent genes (Fzd1–10) in mammals. Wnt engages Fzd by grasping with its thumb and finger extensions, with the lipid cradled in a hydrophobic groove of the extracellular cysteine rich domain (CRD) of Fzd. Binding specificities between the 19 Wnts and 10 Fzds are poorly characterized, but it is generally accepted that interactions exhibit a certain degree of promiscuity. Specificity in binding and subsequent signaling is conferred through the interaction with co-receptors, in particular Lrp5/6. Wnt signaling is partially regulated by the availability of Fzd proteins on the cell surface: in particular, membrane spanning RING-type E3 ubiquitin ligases, such as Rnf43 and Znf3, ubiquitinate Fzd thereby leading to reduced Fzd cell surface expression. The activity of these ubiquitin ligases is in turn regulated by association with a complex comprised of the transmembrane Lgr4/5 receptors and the R-spondin (Rspo1–4) ligands: formation of a Znf3, Lgr, Rspo complex promotes Znf3 turnover, thereby stabilizing Fzd levels and enhancing WNT signaling output (Hao et al., 2012; Chen et al., 2013; Xie et al., 2013; Zebisch et al., 2013). Therefore, even though Rspo by itself has no inherent Wnt signaling activity, it potently augments basal levels of Wnt signaling through increased Fzd receptor availability.

In addition to the Fzd/Lrp receptor complexes, several other Wnt receptors have been identified, including the receptor tyrosine kinase like orphan receptors 1 and 2 (Ror1 and 2) and the receptor-like tyrosine kinase Ryk (Green et al., 2014). Despite their homologies, tyrosine kinase activity has only been demonstrated for Ror2 (Mikels et al., 2009), with Ror1 and Ryk most likely functioning as pseudokinases. Ror1 and 2 carry an extracellular CRD such that Wnt likely engages these receptors in a manner similar to its binding to Fzd. In contrast, Ryk carries an extracellular Wnt inhibitory factor (WIF) domain that directly

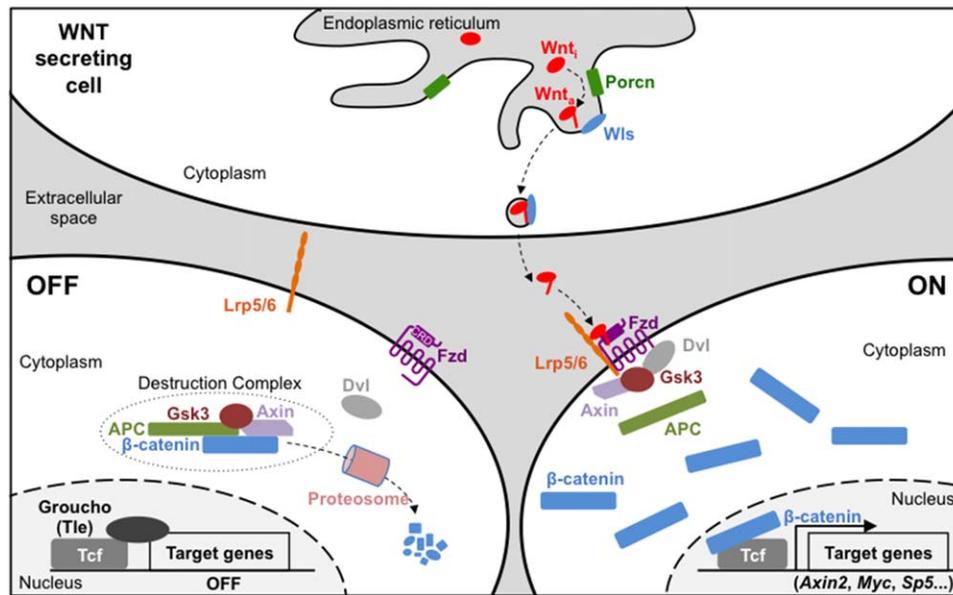


Figure 2 Schematic of Wnt signaling pathway. Upon translation and entry into the secretory pathway, Wnt proteins are acylated by Porcn and ushered out of the cell via Wls. In the off-state, a destruction complex composed of multiple proteins targets β -catenin for degradation by the proteasome. In the on-state, Wnt engages a Fzd/Lrp receptor complex leading to the disassembly of the destruction complex and relocalization of some of its components to the cell membrane. As a result, β -catenin is no longer degraded, accumulates in the cytoplasm and enters the nucleus where it associates with the Tcf transcription factors to activate expression of target genes. [Color figure can be viewed at wileyonlinelibrary.com]

binds Wnt. Signaling mechanisms of these receptors are poorly understood, but is thought to involve formation of various types of receptor complexes, including oligomerization of Ror1 and Ror2 (Yu et al., 2016). In certain contexts Ryk signaling involves proteolytic cleavage to release a C-terminal fragment that translocates to the nucleus where it affects gene expression (Lyu et al., 2008), in a process reminiscent of signaling by the Notch intracellular domain. Similar proteolytic processing upon ligand binding may also occur with other Wnt receptors, including Fzd (Mathew et al., 2005; Mosca and Schwarz, 2010) and ROR1 (Tseng et al., 2010, 2011). Ryk plays critical functions in Wnt-mediated axon guidance, as first demonstrated for the Derailed receptor, the *Drosophila* homolog of Ryk (Yoshikawa et al., 2003). Studies in vertebrate systems have cemented Ryk's critical role as a regulator of neurite outgrowth, acting largely as repulsive cue that inhibits axon growth (Liu et al., 2005; Schmitt et al., 2006; Liu et al., 2008; Hollis et al., 2016) and dendritic arborization (Lanoue et al., 2017). In addition, Ryk has also been implicated in mediating neurite outgrowth (Lu et al., 2004).

Wnt Signaling Pathways. Wnt signaling is generally divided into two categories—commonly referred to Developmental Neurobiology

as canonical and noncanonical Wnt signaling—that are portrayed as independent signaling pathways. Although it is clear that Wnt regulates multiple downstream effects, the distinction between canonical and noncanonical Wnt signaling is an artificial one reflective of experimental settings in which these pathways have been interrogated. A recent review by Nusse and coworkers (Loh et al., 2016) presents a more unified view and posits that integration of these pathways lies at the heart of Wnt's role as a symmetry-breaking signal common to all metazoan life forms: by simultaneously regulating cell fate (= canonical Wnt signaling) and cell polarity (= noncanonical Wnt signaling), Wnts control not only the diversity of cell types but also their organization within the developing organism.

The Wnt cell fate signaling pathway (Fig. 2) has garnered the most attention and interest. In this pathway, Wnt engages a Fzd-Lrp5/6 receptor complex leading to the inactivation of cytoplasmic protein complex, which, in the absence of Wnt signal input, acts to target β -catenin for ubiquitination and proteasomal degradation. Consequently, Wnt signaling stabilizes cytosolic β -catenin, which in turn enters the nucleus and displaces transcriptional corepressors of the Groucho family (TLE) to interact with the

Tcf/Lef family of transcription factors (including Tcf7/Tcf1, Tcf711/Tcf3, Tcf712/Tcf4, and Lef1) and activate transcription of target genes. Some of the main components of the β -catenin destruction complex include the protein products of Adenomatous Polyposis Coli (APC), Axin, and Glycogen Synthase Kinase 3 (Gsk3). Another key component of this Wnt signaling pathway is Dishevelled (Dvl1–3), which associates with the intracellular C-terminus of Fzd and serves to recruit components of the destruction complex to the membrane, thereby releasing β -catenin from the destruction complex. Gsk3, together with the priming kinase Ck1, phosphorylates a series of conserved serine, and threonine residues in the amino terminus of β -catenin to generate a binding site for the β -transducin repeat containing E3 ubiquitin protein ligase (Btrc). Mutation of these phosphorylation sites produces a constitutively active (CA) form of β -catenin; such mutations have been identified in various types of cancer, and CA- β -catenin has served as a powerful tool to study Wnt-independent activation of the pathway.

Wnt/ β -catenin target gene signatures are cell-type specific, however, a common class of target genes encode Wnt antagonists that act at multiple levels of the pathway, including directly on the Wnt protein (e.g., Sfrp and Notum), the receptors (Dkk, Znr3, Rnf43), and the β -catenin destruction complex (Axin2, Nkd). In pluripotent stem cells (PSCs), activation of Wnt signaling induces the transcription of genes associated with endodermal and mesodermal lineages (ten Berge et al., 2008; Davidson et al., 2012), consistent with Wnt's role in primitive streak formation and vertebrate axis formation (Liu et al., 1999; Huelsken et al., 2000). In contrast, ectopic Wnt pathway activation during the derivation of neural progenitor cells from human PSCs induces the expression of genes associated with posterior fates (Moya et al., 2014), again consistent with Wnt's role in establishing the anterior-posterior identity.

Wnt signaling is under exquisitely tight control in the extracellular environment (Fig. 3). Secreted Wnt antagonists can either bind Wnt to preclude receptor binding [including Secreted Frizzled Related Proteins (Sfrp1–5)] and WIF-1 (Hsieh et al., 1999) or bind receptors to preclude Wnt-receptor interactions, as is the case for Dickkopf (Dkk1–4) and Wise (SOSTDC1), which bind Lrp5/6. Additionally, Wnt proteins can be modified by enzymatic mechanisms, such as by Notum, which catalyzes the deacylation of WNT proteins, thereby rendering Wnt unable to bind the CRDs of Fzd. Another protein, Tiki encoded by the TRABD2A and B genes, inactivates Wnt by cleaving the amino-terminal region (Zhang et al., 2012; Zhang et al., 2016). The activity and

distribution of Wnt proteins is additionally influenced by the composition of the extracellular environment, and several proteoglycans, such as Syndecan and Glypican, as well as a host of diverse extracellular matrix proteins, physically interact with Wnt and regulate their signaling range (reviewed in Perrimon and Bernfield, 2000; Zhu and Scott, 2004).

The Wnt/ β -catenin pathway is frequently depicted as a two-state model with an OFF (no Wnt) and ON (Wnt binding its receptor), as shown in Figure 2. However, such a binary system is insufficient in explaining the complex and extensive downstream effects of Wnt signaling. Rather, multiple states of Wnt signaling are essential in generating cellular diversity and tissue patterning. Two mechanisms influencing such a multistate system are likely at work. First, Wnts achieve distinct outcomes depending on their concentration. A concentration gradient of Wnt activity may be achieved by varying the amounts of the Wnt agonist, the availability of receptor(s), or the amounts of Wnt antagonists, such as Sfrp, Dkk, and Notum. Wnt gradients have been visualized in several settings, such as in the *Drosophila* wing (Zecca et al., 1996), *Xenopus* anteroposterior neural patterning (Kiecker and Niehrs, 2001), mouse intestinal crypts (Farin et al., 2016), in caudalization of neural tissue in the chick embryo (Nordstrom et al., 2002), and the ventricular zone of the chick tectum (Schmitt et al., 2006). A second mechanism that imparts multistate complexity on Wnt is through the integration with other developmental signaling systems, including TGF β , Shh, FGF, RA, and Notch. For example, the process of specifying the anteroposterior axis during gastrulation, as first described by Nieuwkoop (1952), requires the cooperative actions of Wnt, BMP, and FGF signaling (reviewed in Hikasa and Sokol, 2013). Later sections will highlight examples of such cooperativity during neural development.

In contrast to the cell fate pathway, very few if any robust transcriptional target genes have been identified downstream of the Wnt cell polarity signaling pathway. Effects of this pathway are generally difficult to model in cell culture systems and, rather, are most clearly observed during organismal or tissue development and involve complex cell movements, as occur, for instance, during gastrulation, gut tube elongation, neural tube closure and neural crest migration. Certain signaling components, including Wnt, Fzd, and Dvl are shared between cell fate and cell polarity pathways, however, some pathways are quite distinct, utilizing distinct receptors (e.g., Ror, Ryk, and Ptk7) and signaling components, such as Vangl, Celsr, and Prickle that act in concert to organize cells within the plane of a tissue.

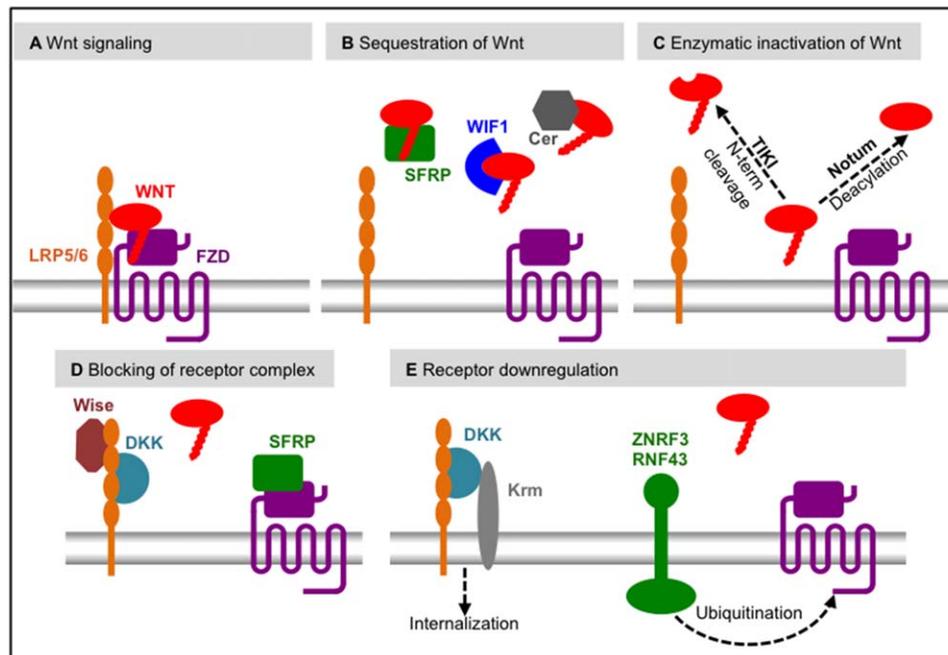


Figure 3 Extracellular regulation of Wnt signaling. A: Wnt signaling is activated when a Wnt protein engages the receptor complex comprised of FZD and LRP. B: Wnt proteins can be sequestered by binding proteins, such as Cerebus (Cer), SFRP, and WIF1, thereby preventing their interaction with the receptor complex. C: Wnt proteins can be inactivated irreversibly, either by deacylation by Notum or by amino-terminal cleavage by TIKI. D: Several secreted molecules interact directly with either FZD (e.g., SFRP) or LRP (DKK and Wise), thereby preventing Wnt-mediated receptor complex formation. E: Wnt receptors are subject to downregulation, thus reducing Wnt signaling. FZD can be ubiquitinated by the E3 ligases ZNRF3 or RNF43 thereby leading to their degradation. LRP is internalized when in complex with the DKK coreceptor Kremen. [Color figure can be viewed at wileyonlinelibrary.com]

The literature is rich with descriptions of distinct Wnt pathways collectively referred to as “non-canonical” Wnt pathways. A variety of downstream signaling components mediate these non-canonical Wnt signals, including G proteins, small GTPases (e.g., Rac and Rho), kinases (e.g., Jnk, Tak1, and Nlk), second messengers (e.g., Calcium), transcription factors (e.g., NF κ B), and more. Assays for these downstream effectors of noncanonical Wnt signaling are scarce and highly cell- and context-dependent. One robust downstream assay of a noncanonical pathway is the inhibition of the Wnt/ β -catenin, as demonstrated for Wnt5a-Ror2 signaling (Ishitani et al., 2003; Mikels and Nusse, 2006; Mikels et al., 2009). Of note, although many studies refer to individual Wnts as either “canonical” or “noncanonical,” the downstream effects of any Wnt is dependent on the expression of receptors and signal transducers rather than by an intrinsic property of the Wnt. For example, the so-called noncanonical Wnt5a can act “canonically” (i.e., through β -catenin) in certain contexts (He et al., 1997; Mikels and Nusse, 2006). This

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review will largely focus on the role of the cell fate pathway, depicted in Figure 2, in neural development. However, it should be stressed that coordinated actions of Wnt in the cell fate and cell polarity pathways are critical in establishing the complexity of the neural system, both in terms of cellular diversity and tissue organization.

Activation or Repression? The Unresolved Role of Wnt Signaling in Neural Specification of Ectoderm

Wnt signaling is highly dynamic throughout all stages of human development with multiple Wnt ligands and antagonists expressed in a temporal and regionally specific manner during early embryonic neurodevelopment (Table 1). The first principal event of neurodevelopment that is influenced by Wnt signaling is the neural specification of embryonic ectodermal cells and the subsequent formation of the neural plate. Pioneering work conducted by Mangold and Spemann in the 1920s led to a proposed

Table 1 Function of Wnt Ligands in Early Neurodevelopment Processes

Ligand	Neurodevelopment Process
Wnt1	AP (Hollyday et al., 1995), FB (Molven et al., 1991; Hollyday et al., 1995; Lagutin et al., 2003), MHB (Wolda et al., 1993; Hollyday et al., 1995; Lagutin et al., 2003; Buckles et al., 2004), HB (Riley et al., 2004), SC (Parr et al., 1993)
Wnt2b	MHB (Kunz et al., 2004)
Wnt3	AP, SC (Roelink and Nusse, 1991), FB (Braun et al., 2003)
Wnt3a	AP, FB (Hollyday et al., 1995), MHB (McMahon et al., 1992; Buckles et al., 2004), HB (Riley et al., 2004), SC (Parr et al., 1993)
Wnt4	AP, FB, MHB (Hollyday et al., 1995), HB (McGrew et al., 1992)
Wnt5a	FB (Hollyday et al., 1995)
Wnt7b	FB (Hollyday et al., 1995)
Wnt8a	NE (Baker et al., 1999), HB (Hume and Dodd, 1993)
Wnt8b	FB (Hollyday et al., 1995; Houart et al., 2002), MHB (Kunz et al., 2004), HB (Riley et al., 2004)
Wnt10a	FB, HB (Kelly et al., 1993)
Wnt10b	MHB (Buckles et al., 2004), HB (Riley et al., 2004)

Abbreviations: NE: neural specification of ectoderm; AP: anterior-posterior patterning of the neural tube; FB: patterning of the forebrain; MHB: induction of the midbrain-hindbrain boundary; HB: specification and segmentation of the hindbrain; SC: induction and patterning of the spinal cord

mechanism by which a subset of ectodermal cells acquire a neural identity through signaling supplied by a central “organizer” in the adjacent mesodermal tissue [now referred to as the Spemann-Mangold organizer; (Spemann and Mangold, 2001) published in 1923]. Subsequently, it was discovered that FGF signaling and BMP antagonism supplied by the Spemann-Mangold organizer was responsible for neural induction of ectodermal tissue (Sasai et al., 1995; Fainsod et al., 1997; Harland, 2000; De Robertis, 2009). However, the role Wnt plays in the determination of neural versus non-neural ectodermal (i.e., epidermal) cell fates has been difficult to precisely ascertain, in part, due to species-specific differences (Stern, 2005).

Early work in *Xenopus* embryos demonstrated that forced expression of Wnt8 or β -catenin inhibited the expression of BMP4, thereby allowing the ectoderm to respond to neural inducing signals, such as FGF (Baker et al., 1999) [Fig. 4(A)]. Additional studies in *Xenopus* revealed that active Wnt signaling promoted neural induction by stimulating the expression of secreted BMP antagonists (Baker et al., 1999) [Fig. 4(A)]. However, other studies implementing *Xenopus* as a model system have produced conflicting results suggesting that Wnt signaling interferes with neural induction (Heeg-Truesdell and LaBonne, 2006; Min et al., 2011). Likewise, work with chick embryos suggests that Wnt signaling promotes epidermal fates rather than neural ectodermal cell identities through the attenuation of FGF signaling and activation of BMP signaling (Wilson et al., 2001) [Fig. 4(B)]. It is

implied that only in the absence of Wnt signaling does FGF upregulation occur, thereby leading to the generation of neural cell fates (Wilson et al., 2001). Work with PSC-based models may provide an opportunity to resolve this controversy in a human system (see **Emerging Trends: Human PSC Models to Study Neurodevelopment**).

Wnt Signaling Is Required for Posteriorization of the Neural Tube

Building on the initial work of Mangold and Spemann, Nieuwkoop proposed a two-step model of neural induction and patterning in which neural tissue of anterior identity is first induced by the Spemann-Mangold organizer and then subsequently patterned along the anterior-posterior (A/P) axis. More specifically, the role of Wnt signaling is multifaceted as a precise balance of both signaling inhibition and activation is required for the development of an endogenous Wnt gradient that specifies regional neural cell identities along the A/P axis (McGrew et al., 1995; McGrew et al., 1997).

Although naïve ectodermal cells acquire an initial anterior neural identity through BMP antagonism and the absence of active Wnt signaling, the maintenance of anterior identity during A/P patterning requires the active inhibition of Wnt activity. The initial evidence of this principle was displayed in zebrafish headless (hdl) mutants, which lack Tcf711/Tcf3-based transcriptional repression of Wnt signaling (Kim et al., 2000). As such, the hdl mutants lack several anterior

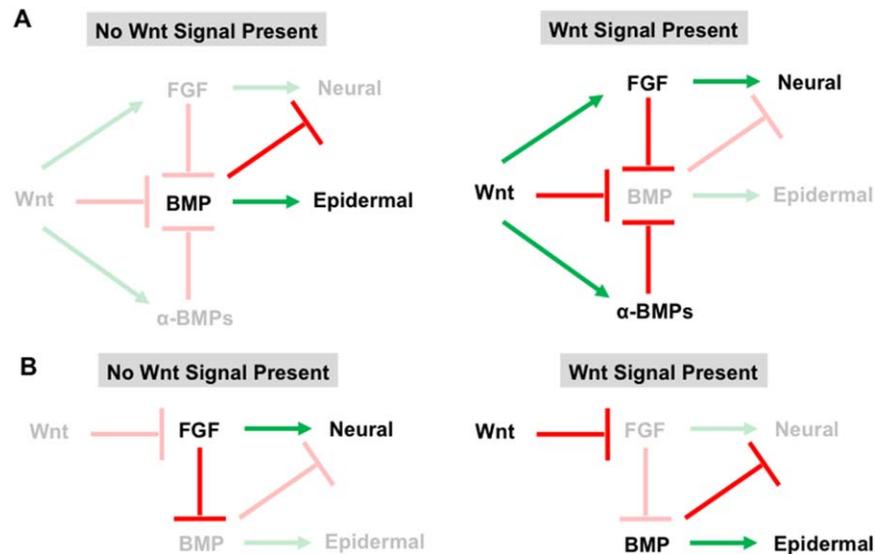


Figure 4 Two proposed models for the role of Wnt signaling in determination of neural versus non-neural ectodermal fates. A: In the first model, in the absence of Wnt signaling BMP expression promotes epidermal fate while repressing neural specification through the inhibition of FGF signaling (upper left panel). Active Wnt signaling promotes ectodermal induction through activation of FGF signaling as well as direct and indirect (through the induction of BMP antagonists [α -BMPs]) repression of BMP signaling (upper right panel). B: In an alternative model, absence of Wnt signaling allows for active FGF signaling to induce neural cell fates (lower left panel). Active Wnt signaling attenuates FGF signaling thereby providing for BMP signals to promote epidermal cell identities (lower right panel). [Color figure can be viewed at wileyonlinelibrary.com]

structures thereby demonstrating that repression of Wnt signaling pathway activity is required for vertebrate head formation and patterning (Kim et al., 2000). Several Wnt antagonists, such as Dickkopf-1 (Dkk1; (Glinka et al., 1998; Hashimoto et al., 2000; Kazanskaya et al., 2000; Mukhopadhyay et al., 2001; Kimura-Yoshida et al., 2005) and Sfrps-1, -3, -5 (Sfrps; (Leyns et al., 1997; Kemp et al., 2005; Mii and Taira, 2009), secreted by the surrounding anterior visceral endoderm and anterior mesendoderm, play critical roles in anterior neural development. Several studies have shown that abolishing expression of these Wnt antagonists during A/P patterning results in the absence or malformation of anterior structures (Glinka et al., 1998; Kimura-Yoshida et al., 2005). Additional negative modulators of Wnt activity, including Cerberus (Glinka et al., 1997), Tiki (Zhang et al., 2012; Reis et al., 2014; Zhang et al., 2016), and Notum (Zhang et al., 2015), have also been implicated in anterior neural tissue formation.

Numerous studies have revealed that the direct action of Wnt proteins on anterior neural cells is required for their posteriorization into structures that will eventually become the prospective midbrain, HB, and SC (Nordstrom et al., 2002). For example,

early *ex vivo* studies that involved the overexpression of different Wnt ligands, β -catenin, or Tcfs in *Xenopus* animal cap cells demonstrated the posteriorizing effects of Wnt signaling at the expense of the expression of anterior neural markers (McGrew et al., 1995; Domingos et al., 2001). Complementing these gain-of-function studies, disruption, or loss of Wnt1 (McMahon and Bradley, 1990; Thomas and Capecchi, 1990; Augustine et al., 1993), Wnt3 (Liu et al., 1999), Wnt3a (Augustine et al., 1993; McGrew et al., 1997; Shimizu et al., 2005), and Wnt8 (Erter et al., 2001; Lekven et al., 2001) in *Xenopus*, zebrafish, or mouse embryos led to the expansion of the forebrain compartment and the improper development of mid-brain, HB, and SC structures. More recent studies suggest that the source of this graded posteriorizing Wnt signal is the paraxial dorsolateral mesoderm, which underlies the developing neural tube (Elkouby et al., 2010).

Multiple Roles of Wnt Signaling in Patterning of the Forebrain

After the initial patterning of the neural tube, the forebrain is further partitioned along the anterior-posterior

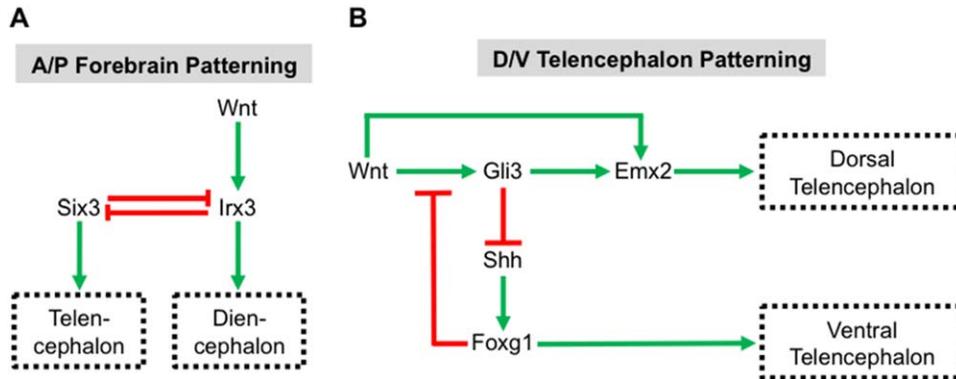


Figure 5 Wnt signaling patterns the forebrain along the anterior-posterior (A/P) and dorsal-ventral (D/V) axis. **A:** Wnt signals induce the expression of *Irx3*, which leads to the adoption of posterior diencephalic cell identities. *Six3* expression results in the generation of anterior telencephalic cell types. Cross-repressive activities of *Six3* and *Irx3* restrict expression of these transcription factors to the telencephalon and diencephalon, respectively. **B:** Wnt signaling establishes dorsal telencephalic (i.e., pallidal) cell fates through the direct and indirect (via *Gli3*) induction of *Emx2* expression and the *Gli3*-mediated repression of *Shh* signaling. Ventral telencephalic (i.e. subpallidal) cell identities are established through the *Shh* stimulated expression of *Foxg1*. In addition, *Foxg1* represses dorsal cell fates in the ventral telencephalon through direct inhibition of Wnt signaling. [Color figure can be viewed at wileyonlinelibrary.com]

(A/P) and dorsal-ventral (D/V) axis. More precisely, the anterior forebrain is patterned into the dorsal telencephalon (which will develop the cerebral cortex) and the ventral telencephalon (which will develop into the basal ganglia) while patterned posteriorly into the diencephalon (which will generate the thalamic tissues; Wilson and Houart, 2004).

As it relates to A/P patterning of the developing forebrain, a general model has emerged in which local Wnt signaling is necessary for the diencephalic development while Wnt antagonism is necessary to promote emergence of telencephalic cell types (Wilson and Houart, 2004). Early evidence for this model was identified in zebrafish with masterblind (*mbl*) mutations in *Axin1*. The absence of *Axin1*-dependent repression of Wnt signaling in *mbl* mutants leads to the anterior expansion of diencephalic cell types and the elimination of telencephalic cell identities (Masai et al., 1997; Heisenberg et al., 2001). Along similar lines, ectopic expression of Wnt signaling during forebrain patterning led to the generation of embryos phenotypically similar to *mbl* mutants (van de Water et al., 2001). Corroborating work with chick (Braun et al., 2003) and mouse (Lagutin et al., 2003) models further demonstrated that proper regionalization of the forebrain requires the repression in Wnt signaling in the anterior compartments and Wnt activation in the posterior areas. Additional studies have implicated *Wnt8b* as the inducer of posterior diencephalic fates in the

emerging forebrain whereas *Tlc*, a zebrafish *Sfrp* homolog secreted by the anterior neural ridge, antagonizes this Wnt signal in the anterior forebrain regions allowing for the development of telencephalic cell identities (Houart et al., 2002; Tian et al., 2002; Echevarria et al., 2003). Finally, several studies have provided insights into the transcriptional targets likely mediating these Wnt-inducing effects of the regionalization of the forebrain (Kobayashi et al., 2002; Braun et al., 2003; Lagutin et al., 2003; Gestri et al., 2005). Briefly, these studies have established that *Irx3* expression is induced by Wnt signaling and acts directly to repress *Six3* expression in the diencephalon, and conversely, *Irx3* represses *Six3* expression in the telencephalon [Fig. 5(A)].

Within the developing telencephalon, Wnt signaling plays a prominent role in specifying dorsal (i.e., pallidal) versus ventral (i.e., subpallidal) cell fates (Campbell, 2003). Initial experiments with chick embryos implicated *Wnt1*, *Wnt4*, and *Wnt8b* as likely inducers of the Wnt signaling response in the dorsal telencephalon (Hollyday et al., 1995). Additionally, *ex vivo* experiments with chick neural tissue explants demonstrated that soluble *Wnt3a* induced dorsal identities in ventral cells (Gunhaga et al., 2003). Conversely, dorsal telencephalic cells failed to emerge in explant cultures in which Wnt signaling was inhibited (Gunhaga et al., 2003). Similarly, targeted deletion of β -catenin in mouse embryos negatively impacted dorsal telencephalic development

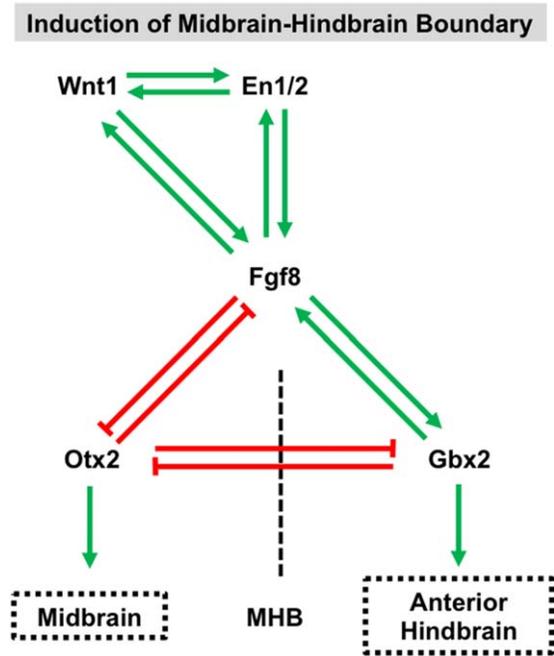


Figure 6 Wnt1 and Fgf8 signaling establishes a gene regulatory network to control the induction and maintenance of the midbrain-hindbrain boundary (MHB). A positive regulatory loop established by the reciprocal expression of Wnt1, En1/2, and Fgf8 is responsible for establishing and maintaining the MHB. Concomitantly, mutual activation of Fgf8 and Gbx2 establishes anterior HB (i.e., metencephalic) cell identities posterior to the MHB whereas mutual repression of Fgf8 and Otx2 induces midbrain cell fates anterior to the MHB. Reciprocal Otx2 and Gbx2 repression is also critical in establishing these cell types at the MHB. [Color figure can be viewed at wileyonlinelibrary.com]

(Gulacsi and Anderson, 2008). Finally, numerous studies have revealed a complex gene regulatory network by which Wnt target genes such as *Emx2* and *Gli3* exert their dorsalizing effects on the immature telencephalon through the repression of *Shh* signaling (Theil et al., 2002; Alvarez-Medina et al., 2008) [Fig. 5(B)]. Conversely, in the ventral telencephalon *Shh* target genes, such as *Foxg1*, serve as transcriptional repressors of Wnt signaling (Danesin et al., 2009) [Fig. 5(B)].

Wnt and Fgf Signaling Act Cooperatively to Induce the Midbrain-HB Boundary

The midbrain-hindbrain boundary (MHB) is a key organizing center that is critical for the proper formation of midbrain and anterior HB (Rhinn and Brand, 2001; Harada et al., 2016). Examination of developing *Xenopus* (Schohl and Fagotto, 2002) and mouse (Maretto et al., 2003) embryos revealed a strong

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expression of nuclear β -catenin in the MHB, suggesting a role for Wnt signaling in MHB generation and maintenance. Indeed, Wnt1 mutants display prominent abnormalities in the midbrain and the metencephalic regions of the HB (McMahon and Bradley, 1990; Thomas and Capecchi, 1990; Thomas et al., 1991; McMahon et al., 1992). Subsequent studies revealed that Wnt1 expression stimulates cell proliferation and survival, possibly explaining the observed phenotype in Wnt1 mutants (Serbedzija et al., 1996; Megason and McMahon, 2002). In addition, *Fzd3/Fzd6* double mutants exhibit significant defects in midbrain morphogenesis, suggesting that Wnt signals act through these receptors for proper MHB development (Stuebner et al., 2010).

Extensive studies have investigated the transcriptional mechanisms by which Wnt signaling regulates MHB induction (Fig. 6; Wurst and Bally-Cuif, 2001; Raible and Brand, 2004). Early studies with Wnt1 knockout mouse embryos revealed that loss of the midbrain and adjacent metencephalic regions was preceded by a complete loss of *Engrailed 1* and *2* (*En1* and *2*) expression (McMahon et al., 1992). Subsequent studies revealed that expression of *En1* in the developing midbrain of Wnt1 mutant embryos is sufficient to rescue midbrain and anterior HB developmental deficits (Danielian and McMahon, 1996). Likewise, complementary studies in *Xenopus* demonstrated that *En2* is a direct target of the Wnt signaling pathway (McGrew et al., 1999). Collectively, these studies demonstrate that *En1* and *En2* act downstream of Wnt1 signaling activity to regulate MHB emergence and identity. Additional evidence suggests a more complex regulatory network in which Wnt1 signals through *En1* to regulate *FGF8* expression in the MHB (Wurst and Bally-Cuif, 2001; Ye et al., 2001; Raible and Brand, 2004). In turn, *FGF8* acts through *Otx2* and *Gbx2* to establish and maintain MHB cell identity (Joyner et al., 2000).

Wnt Signaling Directs the Specification and Segmentation of the HB

As discussed previously, initial HB development proceeds in two stages—the rhombencephalon first emerges from the developing neural tube and then next divides into eight defined rhombomeres (Guthrie and Lumsden, 1991; Glover, 2001). The Wnt signaling pathway is the primary inducer of the gene network that regulates this initial induction and subsequent segmentation (Fig. 7).

Wnt3a secreted from the neighboring paraxial dorsolateral mesoderm is the initial signaling event that governs the emergence of the rhombencephalon from

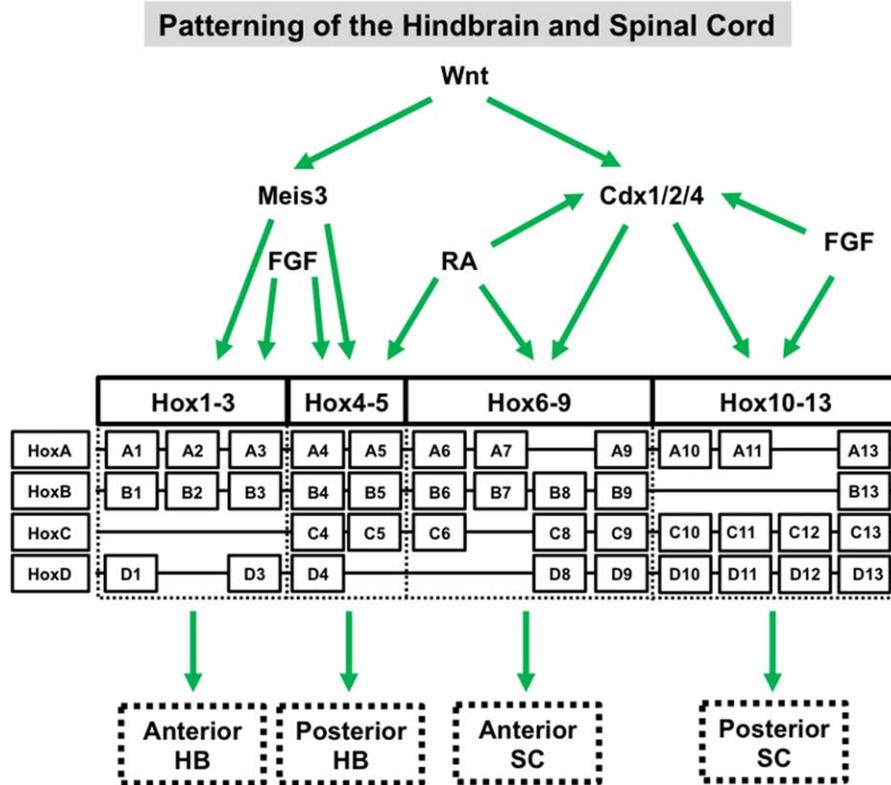


Figure 7 Wnt signaling is refined by FGF and RA signaling to induce and pattern the HB and SC. In the HB, Wnt can directly (not shown) or indirectly (via Meis3) induce the expression of the Hox1–Hox5 paralog group genes. In addition, FGF signaling alone or in concert with RA signaling can further refine the expression pattern of Hox genes in the anterior and posterior HB, respectively. In the SC, Wnt-induced Cdx 1/2/4 expression leads to the induction of the Hox6–Hox13 paralog group genes. Moreover, RA signaling alone or in combination with FGF signaling can further refine the expression, directly or indirectly (via Cdx1), of Hox genes in the anterior and posterior SC, respectively. [Color figure can be viewed at wileyonlinelibrary.com]

the developing neural tube (Elkouby et al., 2010). This initial endogenous signal stimulates the downstream expression of Meis3, which is a direct target of Wnt signaling. Multiple studies have shown that misexpression of Meis3 results in loss of HB, but not SC, structures as well as expansion of anterior cell types (Dibner et al., 2001; Elkouby et al., 2010; Gutkovich et al., 2010). Corresponding studies have also shown that Meis3 expression is sufficient for HB induction in the absence of exogenous Wnt signals (Elkouby et al., 2010). Although other HB makers, such as Gbx2 (Li et al., 2009a) and Hox1–Hox5 paralog group genes (In der Rieden et al., 2010), have been shown to be direct targets of the Wnt pathway, their activation is dependent on the expression of Meis3 (Vlachakis et al., 2001; Dibner et al., 2004; Elkouby et al., 2010, 2012). Together, this suggests a mechanism by which Wnt signaling directly induces Meis3 expression to activate a downstream gene

regulatory network to control HB development. Finally, it should be noted this Wnt signal is further refined by FGF and retinoic acid (RA) signaling to further subdivide the prospective rhombencephalon (Partanen, 2007; Esain et al., 2010; Ishioka et al., 2011; Mazzoni et al., 2013).

Wnt signaling also plays a prominent role in rhombomere boundary formation and maintenance. Specifically, using zebrafish as a model system, Riley et al. found that several Wnt genes (e.g., Wnt1, Wnt3a, Wnt8b, and Wnt10b) were elevated at rhombomere boundaries (Riley et al., 2004). Moreover, knockdown of Tcf-mediated Wnt gene transcription resulted in complete elimination of boundary cell types. Additional work revealed that Wnt signaling, mainly via the action of Wnt1, maintains boundary cell identity by promoting neuronal differentiation in nonboundary regions of the rhombomere (Amoyel et al., 2005).

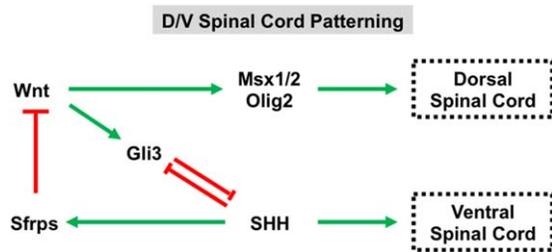


Figure 8 Wnt and SHH signaling act in an opposing manner during dorsal/ventral (D/V) patterning of the SC. Wnt signaling promotes dorsal fate specification in the SC through the activation of target genes such as *Msx1/2* and *Olig2*. Wnt signaling also represses SHH signaling in the dorsal SC by inducing *Gli3* expression. Shh signaling promotes ventral SC cell identities and represses Wnt signaling through induction of Wnt antagonists, such as *Sfrps*. [Color figure can be viewed at wileyonlinelibrary.com]

Wnt Acts in Concert with Multiple Signaling Molecules to Induce and Pattern the SC

In a manner similar to that in the HB, Wnt acts through a complex gene regulatory network to regulate the initial formation of the SC (Melton et al., 2004; Elkouby and Frank, 2010; Fig. 7). Several studies have implicated *Cdx 1, 2, and 4* as the chief transcriptional mediators of the Wnt inducing response in the SC (Lohnes, 2003). These *Cdx* genes have several Tcf binding sites in their regulatory regions and are direct targets of Wnt signaling (Prinos et al., 2001; Lickert and Kemler, 2002; Haremakei et al., 2003; Pilon et al., 2007; Wang and Shashikant, 2007). Moreover, gain- (Prinos et al., 2001; Shimizu et al., 2005; Keenan et al., 2006) and loss-of-function (Prinos et al., 2001; Shimizu et al., 2005) studies confirm the role of these *Cdx* genes in mediating downstream expression of SC-specific *Hox5–13* paralog group genes. It should be noted, although, that there appears to be some functional redundancy between these *Cdx* genes as knockdown of individual *Cdx* genes leads to a range of effects on *Hox* expression patterns whereas only the compound loss-of-function of *Cdx 1, 2, and 4* leads to complete posterior truncation (van den Akker et al., 2002; Shimizu et al., 2005; Faas and Isaacs, 2009; Savory et al., 2009). Finally, both RA and FGF act in concert with Wnt signaling to further refine the *Hox* expression pattern in the developing SC with RA inducing anterior cell fates and FGF promoting posterior identities (Bel-Vialar et al., 2002; Shimizu et al., 2006; Mazzoni et al., 2013; Philipidou and Dasen, 2013).

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As the SC continues to mature, Wnt signals from the roof plate oppose Shh signals from the notochord and ventral plate to pattern cells along the D/V axis (Jessell, 2000; Wilson and Maden, 2005; Ulloa and Marti, 2010). Early studies revealed that both Wnt1 and Wnt3a are highly expressed in the dorsal SC (Parr et al., 1993). Studies in which Wnt1 and Wnt3a were knocked out confirmed the importance of these signaling molecules in dorsal SC specification as Wnt1/Wnt3a double knockout mice showed a marked reduction in neural crest derivatives, dorsolateral neural precursors in the neural tube, and spinal neurons and a concomitant increase in ventral spinal neurons (Ikeya et al., 1997; Muroyama et al., 2002). Mechanistically, Wnt signaling induces the expression of *Gli3* to directly repress Shh signaling (Alvarez-Medina et al., 2008; Yu et al., 2008; Ulloa and Marti, 2010; Fig. 8). In turn, this leads to a gradient of Shh ventral-inducing signaling activity that patterns the SC into various D/V domains.

Wnt and Neurodevelopmental Defects

With its nearly ubiquitous functions in development, it is not surprising that perturbations in Wnt signaling lead to a variety of neurodevelopmental defects in mammals, many of which have been discussed in the previous sections. In addition to the engineered Wnt knockouts described above, classic mouse mutations have also provided evidence for the importance of Wnt signaling in neural development. For example, the recessive mutation *swaying* (*sw*), which is characterized by ataxia, carries a frameshift mutation in the *Wnt1* gene (Thomas et al., 1991), consistent with the phenotype observed in the *Wnt1* knockout model (McMahon and Bradley, 1990; Thomas and Capecchi, 1990). Furthermore, the recessive mutation *vestigial tail* (*vt*) is a hypomorphic allele of *Wnt3a* (Greco et al., 1996), which is required for hippocampal development (Lee et al., 2000), and—together with *Wnt1* deficiency—for proper neural tube development (Ikeya et al., 1997). As the name implies, *vt* mice present with shortened tails due to loss of caudal vertebrae. In contrast, the engineered *Wnt3a* null mutation is embryonic lethal with a severe posterior truncation. As expected, allelic combinations of *vt* and the null mutation produce intermediate phenotypes.

Mutations in WNT genes in humans are extremely rare and the few that have been reported generally did not present with overt neurodevelopmental defects. For example, mutations in WNT1 are associated with a syndrome of severe bone fragility, called osteogenesis imperfecta (OI; Fahiminiya et al., 2013; Keupp et al., 2013; Laine et al., 2013; Pyott et al.,

2013), without overt neurological problems as one may expect given the mouse *Wnt1* knockout phenotype. However, here it is important to note that the neurological defects in mice are quite variable, ranging from ataxia to neonatal death, indicating that loss of function phenotypes are variable in different genetic backgrounds. Furthermore, mice carrying the spontaneous *Wnt1* mutation *sw*, aside from exhibiting the aforementioned neurodevelopmental defects, serve as useful model of OI (Joeng et al., 2014). Mutation of human *WNT3* is embryonic lethal with tetra-amelia, which is characterized by complete loss of all four limbs (Niemann et al., 2004). This severe phenotype has precluded identification of associated neurological defects, as observed in the mouse *Wnt3* knockout. Mutations in *WNT5A* produce Robinow syndrome (Person et al., 2010), which is characterized by multiple morphological defects of stature, skeleton, head, face and external genitalia. Neurodevelopmental defects, as observed in mouse *Wnt5a-Wnt1* double mutants (Andersson et al., 2013), are not common in individuals with Robinow syndrome and intelligence is generally normal.

Mutations in *Porcn* and *Wls*, which encode critical Wnt processing enzymes, are predicted to block secretion of all Wnts and produce “all Wnt mutant” phenotypes. Consistent with this model, global mutations in these genes are embryonic lethal and fail to complete gastrulation, thus precluding analysis of later events, such as neural development. Use of the Cre-Lox recombination system to create conditional alleles of *Porcn* and *Wls* has confirmed the importance of Wnt signaling in various aspects of early neural patterning. For example, heterozygous *Porcn* deleted females are embryonic lethal with open neural tubes and decreased expression of the neuroectodermal markers *Gbx2* and *Sox2* (Liu et al., 2012). Deletion of *Wls* with a neural specific Cre driver (*Wnt1-Cre*) results in deletion in midbrain and HB, a phenotype resembling *Wnt1* mutation (Carpenter et al., 2010; Fu et al., 2011). Interestingly, *PORCN* mutations in humans produce a condition called focal dermal hypoplasia (FDH, also known as Goltz syndrome; Grzeschik et al., 2007; Wang et al., 2007), which is characterized by skin lesions and defects of the gastrointestinal and cardiovascular system. The reason for the extreme pleiotropic phenotypes observed in FDH is due to the fact that *PORCN* is encoded on the X chromosome and hence is subject to random X-inactivation. Neurodevelopmental and cognitive defects are infrequently observed in FDH, which likely indicates that defects due to inactivation of *PORCN* in neural tissues are too severe to permit survival to birth.

Emerging Trends: Human PSC Models to Study Neurodevelopment

Studies in model animal organisms, including frogs, chick and mouse, have provided important insights into the role of Wnt signaling in early neural development and patterning. However, the inherent complexities of these biological systems have made it difficult to dissect the underlying molecular mechanisms associated with complex neurodevelopmental processes. Human PSCs (hPSCs) provide an emerging opportunity to study the mechanisms of human neurodevelopment in an easily controllable and accessible system. In addition, hPSC-based models enable for the resolution of species-specific differences that have been observed in some model systems. For example, studies with model organisms have provided conflicting evidence with regards to whether Wnt induces neural or non-neural ectodermal cell fates. To that end, Lee et al. showed that hPSC lines with amplified *WNT3/WNT9B* revealed that upregulation of Wnt signaling enhanced neural differentiation (Lee et al., 2015).

hPSC-based systems have also been used to model and elucidate the roles of Wnt signaling in early A/P patterning of the neural tube (Kriks et al., 2011; Kirkeby et al., 2012; Moya et al., 2014). For example, Moya et al. developed an *in vitro* hPSC-model that mimics the same effects of the Wnt signaling gradient on the early A/P patterning of the neural tube observed during *in vivo* development (Moya et al., 2014). Moreover, this system allowed for genome-wide expression analysis of cell populations of various A/P regional identities. This analysis provided novel insight into the mechanisms by which Wnt signaling regulates A/P fate in that Wnt signaling appeared to be acting cell autonomously, with Wnt signaling activity restricted to those cells expressing Wnt genes. This suggests that during development WNT proteins may act in an autocrine, rather than paracrine, manner to specify and maintain A/P neural cell identities.

Experiments with hPSC-based models have also allowed for a better understanding of the mechanisms by which Wnt signaling regulates D/V patterning of the forebrain (Li et al., 2009b). Specifically, through the precise activation and inhibition of Wnt and Shh signaling, Li and coworkers were able to generate cells of various telencephalic D/V cell fates. Moreover, the authors identified a Gli3-dependent mechanism by which Wnt signaling represses Shh signaling to promote dorsal telencephalic fates.

More recently, studies with hPSCs have allowed for the unraveling of the temporal mechanisms by

which Wnt patterns the HB and SC (Lippmann et al., 2015). Through the modulation of Wnt, RA, and FGF signaling, Lippmann et al. were able to generate early neural cells with HOX profiles corresponding to specific rhombomeric segments and cervical, thoracic, and lumbosacral identities. In addition, the authors discovered that Wnt, RA, and FGF act in a biphasic manner to induce discrete HB and SC regional identities.

In the future, the use of 3-D culture methods and organoid approaches may allow for the generation of hPSC-based models that more accurately mimic *in vivo* neural development (Lancaster et al., 2013; Meinhardt et al., 2014; Kelava and Lancaster, 2016). A recent study showed that modulation of endogenous Wnt signaling (with IWP2, a Porcn inhibitor) promoted the formation of spheroids that resembled ventral forebrain or the subpallium as monitored by strong induction of NKX2-1 and FOXP1 (Birey et al., 2017). Finally, the use of these systems with next-generation sequencing technologies, such as single-cell RNA sequencing, will allow for a better understanding of how individual cells make fate decisions in early neurodevelopmental processes (Yao et al., 2017).

CONCLUDING REMARKS

Despite the plethora of research to date, there is still much that needs to be unraveled about the role of Wnt signaling in the earliest stages of neural development. For instance, the mechanisms by which the Wnt cell polarity (= noncanonical) pathway acts in concert with the Wnt cell fate (= canonical) pathway in neural fate decisions has not been subject to extensive investigation. In addition, the complex manner in which Wnt interacts with other signaling pathways (e.g., BMP, Shh, FGF, RA, Notch) and components of the cellular microenvironment (e.g., mechanical forces, extracellular matrix proteins) to precisely regulate early neural developmental cell fate has yet to be ascertained. Finally, the downstream epigenetic and genetic gene regulatory networks by which Wnt imparts specific neural cell identities have not been fully identified. In the future, continued advancements with *in vitro* culture systems (e.g., hPSCs), genetic manipulation systems (e.g., CRISPR/Cas9), and molecular analysis techniques (e.g., next-generation sequencing), will allow for a more complete understanding of these multifaceted roles Wnt signaling plays in CNS development.

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