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Wnt stabilization of β -catenin reveals principles for morphogen receptor-scaffold assemblies

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Abstract

Wnt signaling stabilizes -catenin through the LRP6 receptor signaling complex, which antagonizes the -catenin destruction complex. The Axin scaffold and associated glycogen synthase kinase-3 (GSK3) have central roles in both assemblies, but the transduction mechanism from the receptor to the destruction complex is contentious. We report that Wnt signaling is governed by phosphorylation regulation of Axin scaffolding function. Phosphorylation by GSK3 kept Axin activated ("open") for -catenin interaction and poised for engagement of LRP6. Formation of the Wnt-induced LRP6-Axin signaling complex promoted Axin dephosphorylation by protein phosphatase-1, and inactivated ("closed") Axin through an intra-molecular interaction. Inactivation of Axin diminished its association with -catenin and LRP6, thereby inhibiting - catenin phosphorylation and enabling activated LRP6 to selectively recruit active Axin for inactivation reiteratively. Our findings reveal mechanisms for scaffold regulation and morphogen signaling.

Signaling by secreted Wnt morphogens governs developmental, homeostatic, and pathological processes by regulating -catenin stability, and represents a critical target for cancer and disease therapeutics (1, 2). Without Wnt stimulation, cytosolic -catenin concentrations are kept low because a "destruction complex" assembled by the Axin scaffold binds to -catenin, *Adenomatosis polyposis coli* (APC), casein kinase-1 (CK1), and glycogen synthase kinase-3 (GSK3), and promotes phosphorylation of -catenin by CK1 and GSK3 thus ensuring -catenin ubiquitination and degradation (1–3). Upon Wnt stimulation, a receptor complex on the cell surface is formed between Frizzled (Fz) and LDL receptor-related protein 6 (LRP6), resulting in phosphorylation and activation of LRP6 and its recruitment of Axin (4–7). Assembly of the Fz-LRP6 complex and associated

Supplementary Materials

Materials and Methods Figs. S1 to S18

References (31-39)

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Dishevelled (Dvl) and the Axin destruction complex, referred to collectively as the "LRP6 signaling complex (signalosome)", inhibits phosphorylation of -catenin thereby causing its stabilization (6–10). The mechanism by which LRP6 activation leads to -catenin stabilization remains enigmatic (1, 2, 11).

Axin is a phospho-protein and central to assemblies of both destruction (12–15) and signaling complexes (4–10), and becomes dephosphorylated upon Wnt stimulation (16, 17). We generated an antibody, Ab-pS497/500 (fig. S1A to S1C), for Axin phosphorylated at serines 497 and 500, which are GSK3 phosphorylation sites in vitro (18). Axin phosphorylation at S497/S500 was decreased within 15–30 min of Wnt3a treatment of mouse L fibroblasts (Fig. 1A), embryonic fibroblasts (fig. S1D), and human embryonic kidney (HEK) 293T cells (Fig. 1C and 1D). Wnt-induced dephosphorylation of Axin likely reflects the counterbalance between GSK3 and a protein phosphatase (PP) such as PP1, whose catalytic subunit, PP1c, was identified in an RNAi screen in Drosophila cells as a requirement for Wnt/ -catenin signaling (19). Through a functional cDNA overexpression screen in HEK293T cells, we identified PP1c , one of the three PP1c genes in the human genome (20), as an activator of the Wnt/ -catenin signaling reporter TOP-Flash (fig. S2A). PP1c overexpression decreased phosphorylation of Axin but not of LRP6 (Fig. 1B); a pharmacological PP1 inhibitor, Tautomycin (TM), prevented Wnt-induced dephosphorylation of Axin without affecting LRP6 phosphorylation (Fig. 1C and fig. S3).

PP1 has pleiotropic roles and its specificity is conferred by hundreds of PP1c-binding proteins (20). Inhibitor-2 (I2, or PPP1R2) is a specific inhibitor of PP1c (20). Overexpression of I2 countered Wnt3a-induced Axin dephosphorylation (without affecting LRP6 phosphorylation) and -catenin stabilization (Fig. 1D and fig. S2B), inhibited Wnt3a-or PP1c -activated TOP-Flash (fig. S2C and S2D), and antagonized -catenin stabilization by an activated LRP6 (fig. S2E). Depletion of the endogenous I2 with shRNAs resulted in accumulation of -catenin and increased TOP-Flash (Fig. 1E and fig. S2F). A morpholino antisense-oligonucleotide (MO) that targets Xenopus I2 mRNA and blocked I2 protein synthesis caused deficiency in Xenopus head development and reduced anterior marker expression, which were restored by human I2 mRNA injection or knockdown of -catenin (Fig. 1F and fig. S4). Thus I2 antagonizes Wnt/ -catenin signaling and participates in vertebrate anteriorization, which requires Wnt pathway inhibition (21).

Recent models of Wnt signaling (22, 23) have overlooked Axin phosphorylation, and one argued that Wnt/LRP6 signaling maintains an intact Axin destruction complex without inhibiting -catenin phosphorylation (22). We reevaluated this critical issue. We found that in multiple Wnt-responsive cells, the rate of -catenin phosphorylation was suppressed under Wnt stimulation (fig. S5) as reported (3, 24), and this correlated with Axin dephosphorylation.

Association of Axin and -catenin is the cornerstone binary interaction within the destruction complex. Co-immunoprecipitation (co-IP) of endogenous proteins showed that the amount of -catenin associated with Axin appeared to be unchanged or decrease after 0.5 hours of Wnt treatment but to increase after 2 hours (fig. S6A and S6B). However normalization to the amount of cytosolic -catenin made it clear that Wnt stimulation weakened the interaction between -catenin and Axin (Fig. 2A and 2B, left panels). We estimated that the dissociation constant (Kd) of Axin- -catenin association was increased by Wnt3a (fig. S6C), signifying a weaker interaction. Indeed Axin from extracts of Wnt3a-treated cells exhibited reduced capability to associate with -catenin in an in vitro binding assay (Fig. 3A) (17). Such diminished Axin- -catenin association observed in vitro and in vivo correlated with, and appeared to be attributable to Axin dephosphorylation by PP1, because it could be partially restored by I2 overexpression or TM treatment (Fig. 3A to 3C).

For -catenin co-immunoprecipitated with Axin from Wnt3a-treated cells, the amount of phosphorylation relative to that of -catenin was reduced (Fig. 2A and 2B, middle panels; fig. S6A and S6B), implying a diminished rate of -catenin phosphorylation in the Axin complex in Wnt-treated cells. Association of Axin with GSK3 remained constant regardless of Wnt treatment (fig. S6A and S6B) (22, 25). Thus Wnt signaling inhibits Axin- -catenin association and -catenin phosphorylation, consistent with earlier findings (3, 17), kinetic modeling (24), and the prevailing model (1, 11, 14, 15) (see Fig. S6C).

Wnt-induced phosphorylation of LRP6 recruits Axin to assemble the signaling complex (5-7). Intriguingly, co-IP of endogenous proteins showed that Wnt-induced association of LRP6 and Axin was prominent at 0.5 hours but diminished after 2 hours (Fig. 2A and 2B, right panels; and fig. S6A and S6B), despite of persistence of phospho-LRP6 (fig. S5). The diminished binding between phospho-LRP6 and Axin appeared to result from Axin dephosphorylation by PP1, because the binding was enhanced by I2 overexpression or TM treatment (Fig. 3B and 3C) and reduced by I2 depletion (Fig. 3D). Axin from extracts of Wnt3a-treated cells showed reduced capability to associate with phospho-LRP6 in an in vitro binding assay (fig. S7A), and this reduction appeared to result from Axin dephosphorylation by PP1 because it was prevented by I2 overexpression (Fig. 3A). Complementarily, Axin from extracts of cells treated with a pharmacological GSK3 inhibitor, BIO or SB216763, or of Gsk3 -/-;Gsk3 -/- cells (26), had minimal association with phospho-LRP6 (Fig. 3E and fig. S7B). Thus phosphorylation of Axin by GSK3 enhanced, whereas dephosphorylation of Axin by PP1 diminished, Axin's ability to associate with phospho-LRP6 (and -catenin), implying that activated LRP6 selectively recruits the phosphorylated form of Axin that is active in -catenin association and degradation.

Above results imply that Axin may undergo a phosphorylation-dependent conformational change. Axin is an intrinsically disordered protein with individual partner-binding domains (fig. S8A) (14, 15). Axin's -catenin binding domain (Axin-BCD) associated, in vitro and when overexpressed in cells, with Axin DIX domain (Axin-DIX) but not homologous Dvl2 DIX domain (Dvl-DIX) (Fig. 4A, fig. S8B and S9), reflecting a specific and direct interaction. Phosphorylation of Axin-BCD by GSK3 in vitro inhibited BCD-DIX binding (Fig. 4A and fig. S8C). Therefore there may be an intra-molecular BCD-DIX interaction that is prevented upon Axin phosphorylation by GSK3, providing an explanation for how phosphorylation of Axin enhances its association with -catenin and phospho-LRP6. Indeed Axin intra- and inter-molecular interactions appeared to be mutually exclusive because catenin and Axin-DIX competed for binding to Axin-BCD (Fig. 4B and fig. S10A). A positively charged histidine-rich region of BCD (fig. S9, S10B, and S11) and a negatively charged loop of DIX (fig. S12A and S13) participated in BCD-DIX interaction, which appeared to be disrupted by negative charges generated in BCD through phosphorylation by GSK3 (figs. S10C, S10D, and S11). Axin(SD4), which contains phosphomimetic aspartic acid substitutions of 4 serines (including S497 and S500) in BCD (fig. S11), and Axin(DA), which contains alanine substitutions of acidic residues in DIX (fig. S13), were each expected to have a weaker intra-molecular interaction (fig. S10D and S12A) and were indeed more effective in inhibiting Wnt/ -catenin signaling than the WT Axin (Fig. 4C). Axin(SA4), which contains alanine substitutions of the 4 serines in BCD (fig. S11) and was predicted to have a stronger intra-molecular interaction (fig. S10C), was less effective in inhibiting Wnt/ -catenin signaling (Fig. 4C). These results support a model that Axin "autoinhibits" through the BCD-DIX intra-molecular interaction. Live cell FRET (fluorescence resonance energy transfer) imaging corroborated this model by demonstrating a Wnt3a-induced proximity of Axin's carboxyl DIX to its amino terminus (fig. S14 to S17), likely through Axin dephosphorylation.

We propose a Wnt signaling model (Fig. 4D) that unifies findings on the two Axin complexes mediating LRP6 signaling and -catenin destruction. Without Wnt, Axin is associated with and phosphorylated by GSK3 and is in an activated ("open") conformation for -catenin binding and phosphorylation and poised for engagement of LRP6 (Fig. 4D). With Wnt, LRP6 undergoes Fz/Dvl-dependent phosphorylation and recruits the active Axin destruction complex to form the signaling complex, in which GSK3 bound to Axin is inhibited by phospho-LRP6 (27-29), leading to initial inhibition of -catenin phosphorylation and tipping the balance towards Axin dephosphorylation by PP1. Dephosphorylated Axin adopts an inactivated ("closed") conformation through intramolecular autoinhibition, and becomes incompetent for association with -catenin or phospho-LRP6, leading to disassembly of destruction and signaling complexes (Fig. 4D). Phospho-LRP6 is thus freed for another round of recruitment of phosphorylated-activated Axin for inactivation while ignoring dephosphorylated-inactivated Axin, and the steps likely reiterate to keep -catenin phosphorylation suppressed. Wnt-induced biphasic assembly and disassembly of the LRP6 signaling complex appear to enable phospho-LRP6 to inactivate Axin in a "catalytic" manner, underlying Wnt stabilization of -catenin in broad component stoichiometries. GSK3 acts as an "assembler" of destruction (16-18) and signaling complexes (4, 6, 8-10) through phosphorylation of Axin and LRP6, whereas PP1 dephosphorylates Axin (19) to disassemble both complexes while leaving phospho-LRP6 unperturbed for continuous signaling. Our model further explains the -catenin stabilization kinetics. Elevating levels of -catenin, by competing against Axin intra-molecular autoinhibition, could promote reassembly of the Axin-GSK3- -catenin complex and counter its disassembly by Wnt (Fig. 4D), thereby plateauing when equilibrium is achieved. This implies a safeguard mechanism by which rising concentrations of -catenin could trigger its own degradation to avoid excessive accumulation. Axin represents a scaffold with an on/off switch controlled through a ligand- and phosphorylation-dependent intra-molecular interaction, which additionally could serve as a feedback sensor of target (-catenin) concentrations. Similar scaffold functions could occur in other pathways as highlighted by yeast Ste5 (30).

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

Acknowledgments

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Fig. 1.

Wnt-induced Axin dephosphorylation by PP1 and effects of I2 on Wnt signaling and Xenopus anteriorization. (A) Wnt3a-induced Axin dephosphorylation, LRP6 phosphorylation and -catenin stabilization in L cells. Protein detections were performed by immunoblotting throughout the paper unless specified otherwise. actin: a loading control.
(B) Effect of PP1c overexpression on phosphorylation of Axin but not LRP6 in HEK293T cells. (C and D) Effects of TM (C) or I2 overexpression (D) on Wnt3a-induced Axin dephosphorylation and -catenin stabilization in HEK293T cells. (E) Effects of I2 depletion with shRNAs (sh-I2) on -catenin stabilization (top) and TOP-Flash (bottom) in HEK293T cells. shGFP: an shRNA against GFP. Error bars represent SD of triplicates. (F) Effect of I2 depletion with an I2MO in embryos and its rescue by human I2 mRNA. CoMo/Uninj: control MO-injected/uninjected.



Fig. 2.

Wnt regulation of Axin- -catenin and Axin-LRP6 association, and of -catenin phosphorylation in the Axin complex. (**A** and **B**) Quantifications of Wnt3a effects on ratios of Axin-associated -catenin versus input -catenin (left), Axin-associated phospho- - catenin versus Axin-associated -catenin (middle), and Axin-associated LRP6 versus total LRP6 (right) in L (A) and HEK293T (B) cells. Error bars represent SEM of triplicates.



Fig. 3.

Effects of Wnt-regulated Axin phosphorylation on its binding to -catenin and LRP6. (**A**) HEK293T cells overexpressing I2 (or control) were treated with Wnt3a. Lysates were incubated with GST- -catenin or GST-phospho-LRP6C in vitro. Bound or input Axin, and phospho-Axin were examined. (**B** and **C**) L cells treated with TM (B) and HEK293T cells overexpressing I2 (or control) (C) were stimulated with Wnt3a for 0.5 hours. Lysates were immunoprecipitated with an Axin antibody, and Axin-associated -catenin and LRP6, and input proteins were examined. Quantifications show Axin-associated -catenin versus cytosolic -catenin (top), and Axin-associated LRP6 versus total LRP6 (bottom). Note that TM or I2 reduced Wnt3a-induced -catenin levels (Input). Error bars represent SEM of triplicates. *P<0.05 and **P<0.01 with student-Newman-Keuls test. (**D**) HEK293T cells expressing an I2 (or control) shRNA were treated with Wnt3a for 0.5 hours. Lysates were immunoprecipitated with an Axin antibody, and Axin-associated LRP6 and input Jysates were examined. (**E**) In vitro binding to GST-phospho-LRP6C by Axin from Jysates of WT or Gsk3 -/-;Gsk3 -/- mouse ES cells. Bound or input Axin was examined.



Fig. 4.

A phosphorylation-regulated Axin intra-molecular interaction and a Wnt signaling model. (A) Association of Axin-DIX with GST-Axin-BCD and its inhibition by GSK3 phosphorylation of Axin-BCD. (B) Competition of -catenin association with GST-Axin-BCD by Axin-DIX (lanes 1–6) and vice versa (lanes 7–12). Purified recombinant proteins were used in these in vitro assays. Axin-DIX or -catenin was detected by immunoblotting and GST or GST-Axin-BCD by Ponceau staining (A and B). The lower band of -catenin (B) was a proteolytic fragment. (C) Comparisons of Axin(SD4), Axin(SA4), and Axin(DA) with Axin in antagonizing Wnt-induced TOP-Flash in HEK293T cells. X-axes represent DNA doses transfected. Note larger TOP-Flash differences at lower overexpression doses. Insets show levels of overexpressed Axin at the 1ng dose and that of endogenous Axin (con). (D) An "Axin inactivation" model for Wnt stabilization of -catenin. APC and CK1 were omitted for clarity. See text and fig. S18.