

Reelin Signaling and Cdk5 in the Control of Neuronal Positioning

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Abstract

Neuronal positioning is important for higher brain function because it is the architectural basis of the formation of precise synaptic circuits. Analysis of neurological mutant mice has led to dramatic progress in the identification and characterization of molecules important for neuronal positioning in the developing mammalian brain. Among these molecules, identification of signal pathways mediated by Reelin and Cdk5 kinase has provided a conceptual framework for exploring the molecular mechanisms underlying proper neuronal positioning in the developing mammalian brain. Recent evidence has implicated synergism between Reelin signaling and Cdk5 in contributing to the proper positioning of selective neuronal populations.

Index Entries: Reelin; Dab1; CR-50; VLDLR; ApoER2; CNR; Cajal-Retzius cell; Cdk5; p35; p39.

Introduction

Most neurons migrate from the site of their birth near the ventricle toward the outer surface of the central nervous system (CNS), where they are integrated into specific brain

circuits. Proper positioning of neurons is particularly critical for the formation of cyto-architecturally distinct brain regions, such as the cerebral cortex, hippocampus, and cerebellum. During the development of laminated brain structures, a series of coordinated migrations takes neurons from their site of origin to their final destinations, where they adopt definitive morphological phenotypes by elaborating dendritic and axonal processes (1). In the mammalian brain, neurons settle into six neuronal

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laminae within the cerebral cortex in an inside-out manner, with the earliest-generated neurons positioned in the deepest layers, and later-generated neurons occupying the more superficial layers (2).

Remarkable progress in the identification and characterization of genes important to neuronal positioning in the developing brain has been made by analysis of neurological mutant mice (3). In addition to this, genetic studies of human brain malformations have also proven a valuable means of identifying the molecules that regulate neuronal migration (4). Such studies also serve to relate the significance of genes initially identified in mutant mice, such as *reelin*, to more complex human brain development (5,6). The mutations of *reelin* were first identified in several known alleles of *reeler* (5), and mutations have also been recently reported in humans with autosomal recessive lissencephaly associated with cerebellar hypoplasia (6). Lissencephaly is a condition caused by several neuronal migration disorders that result in a smooth brain. Although mice are normally lissencephalic, neuronal migration defects result in similar misplacements of neurons in laminated brain structures in both humans and mice.

Reelin Signaling

The mouse mutant *reeler* exhibits an ataxic gait and has served as a prototype for the investigation of neuronal migration (7). The *reeler* mouse is characterized by abnormalities of neuronal alignment in several neural systems, including the cerebral cortex, cerebellar cortex, hippocampus, and neuronal ectopia in several brainstem nuclei and the spinal cord (8–14). The reason for the intense interest in a single ataxic mouse is that the *reeler* phenotype highlights most of the major events in normal cortical development. Neocortical development begins with the appearance of the preplate above the ventricular zone; the preplate is composed of subcortical projections, Cajal–Retzius cells, and subplate

neurons (15). In the next phase, the neurons destined to form the future projection neurons of the neocortex leave the ventricular zone, migrate into preplate, and split it in two layers: 1) a superficial layer, known as the marginal zone, and 2) a deeper layer, known as the subplate (15). Ogawa et al. generated CR-50 monoclonal antibody by immunizing homogenates of wild-type embryonic brain to *reeler* mutant mouse (16), and it recognized Cajal–Retzius neurons in the marginal zone of the cerebral cortex and external granular layer of the cerebellum. The cerebral cortex of *reeler* did not react with this antibody, even though Cajal–Retzius cells are present (16,17), and CR-50 antigen was therefore considered a candidate for the *reeler*-gene product. Independently, the *reeler*-gene named *reelin* (*Reln*) has been cloned by insertional mutation of a transgene, and molecular cloning has revealed that Reelin is a large (approx 385 kDa) secreted protein (5). Subsequent studies have shown that CR-50 antibody recognizes a tertiary structure of a region near the N-terminal of Reelin, which is defective in Jackson-type *reeler* (18). In the Orleans *reeler*, Reelin is truncated by a mutation near the C-terminal, and the truncated Reelin protein is not secreted (19). It is important to note that several experimental approaches have revealed that CR-50 functions as a blocking antibody both in vivo and in vitro. CR-50 antibody disrupts the normal alignment of neurons and converts to *reeler* phenotype in reaggregation-culture of cerebral cortex (16), and in hippocampal structure when injected in the embryonic stage (20). Miyata et al. showed that normal alignment of Purkinje cells is blocked by CR-50 antibody, and that abnormal positioning of Purkinje cells in *reeler* cerebellum is rescued by transplanted granule cells isolated from normal cerebellum (21). Biochemistry and structural biology methods have shown that Reelin forms a homomultimer driven by the CR-50 epitope through electrostatic interaction (22). Mutant Reelin that lacks the CR-50 epitope region does not form homomultimers, and Reelin

homomultimers are disrupted in the presence of CR-50, suggesting that multimerization of Reelin is required for it to function (22). Another biochemical study demonstrated a potential role for Reelin as a serine protease of extracellular matrix (23).

Shortly after Reelin was identified, Davisson et al. at Jackson Laboratory described a *reeler* phenotype in a new spontaneous mutant named *scrambler* (24). Another *reeler* phenotype, spontaneous mutant mouse named *yotari*, was found in a colony of IP₃R1 mutant mice, and was shown to have normal Reelin protein expression (25). The *scrambler* locus was mapped to near the *disabled-1* (*Dab1*) gene on chromosome 4 (24), and mice with targeted disruption of *Dab1* were found to exhibit behavioral and anatomical abnormalities indistinguishable from those of *reeler* and *scrambler/yotari* (26). It was then shown that *scrambler/yotari* mice express a mutated form of *Dab1* mRNA; a drastic decrease in *Dab1* protein was found in *scrambler* brain, while an absence of *Dab1* protein was found in *yotari* brain (27). Later, Kojima et al. found that the *yotari* mutation in *Dab1* is caused by a replacement of gene sequence with a long interspersed nuclear element (L1) fragment (28). Characterization of the gene mutation of *Dab1* in *yotari* made it easy to identify *yotari* embryos by PCR-based genotyping (28). *Dab1* is an intracellular adaptor protein containing a motif known as a protein interaction/phosphotyrosine binding (PI/PTB) domain (29). Tyrosine phosphorylation of *Dab1* promotes its interaction with several nonreceptor tyrosine kinases, including Src, Fyn, and Ab1 through their SH2 domains, implicating *Dab1* function in kinase-signaling cascades during brain development (29).

In 1999, behavioral and neuroanatomical defects identical to those in *reeler* were reported in mice lacking both very-low-density lipoprotein receptor (VLDLR) and apolipoprotein E receptor-2 (ApoER2) (30). These members of the low-density lipoprotein receptor (LDLR) family have been shown to bind to the PI/PTB domain of *Dab1* (31). VLDLR and ApoER2 were originally identi-

fied as the receptors for lipoproteins, such as ApoE, and both receptors were enriched in brain. The patterns of expression of their mRNA considerably overlap that of *Dab1* (30), indicating that VLDLR and ApoER2 act in concert as components of the Reelin receptors. Two groups using different methods have independently demonstrated their ability to bind Reelin (32,33). Several lines of evidence have now placed Reelin, VLDLR/ ApoER2, and *Dab1* in a common signal pathway that controls neuronal positioning in the developing CNS.

1. Mutations in these genes result in identical neuroanatomical defects of neuronal positioning (5,25,26,27,30).
2. Reelin binds extracellularly to the ectodomain of VLDLR/ ApoER2, and *Dab1* interacts intracellularly with the cytoplasmic tails of these lipoprotein receptors (31–33).
3. The levels of *Dab1* protein increase several-fold and tyrosine-phosphorylation of *Dab1* protein decreases in the brains of *reeler* and VLDLR/ ApoER2-deficient mice (34).
4. Reelin is expressed in the cells located adjacent to the migrating neurons that express *Dab1* and VLDLR/ ApoER2 (26,27,30).
5. Addition of Reelin to neuronal culture induces tyrosine-phosphorylation of *Dab1* (35).

Tyr¹⁹⁸ and Tyr²²⁰ are considered to represent the major sites of Reelin-induced *Dab1*-phosphorylation in embryonic neurons (36). Mutation of five tyrosines including these two residues in *Dab1* results in *reeler* phenotype, suggesting that tyrosine-phosphorylation of *Dab1* is essential for Reelin signaling (37).

So far, the kinases responsible for in vivo tyrosyl-phosphorylation of *Dab1* are not known. Reelin has also been shown to bind to cadherin-related neuronal receptors (CNRs) (38) and $\alpha\beta 1$ integrins (39). A number of genes have been suggested to play a role in the Reelin signaling (see Table 1), and while many of them have been directly implicated in this signaling, others (40–46), including the genes coding neurotrophic factors (41,42) and transcription factors, appear to be involved in the generation or survival of Cajal–Retzius cells (43–46).

Table 1
Genes Implicated in Reelin Signaling and Cdk5 for the Control of Neuronal Positioning

Gene	Name	Protein type	Chromosome location			Refs.
			Human	Mouse		
<i>Bdnf</i>	Brain-derived neurotrophic factor	Neurotrophin	11p13	1		(41)
<i>NGFβ</i>	Nerve growth factor β subunit	Neurotrophin	1p13	3		(101)
<i>Ntf-4</i>	Neurotrophin 4	Neurotrophin	19	7		(42)
<i>Reln</i>	Reelin	Extracellular protein	7q22	5		(5)
<i>ApoER2</i>	Apolipoprotein E receptor 2	Transmembrane	1p34	ND ^a		(30)
<i>CNR</i>	Cadherin-related neuronal receptor	Transmembrane	5q31	18		(38)
<i>itga-3</i>	α3 integrin	Integrin receptor subunit	17	11		(39)
<i>PS-1</i>	Presenilin-1	Transmembrane	14q24	ND		(40)
<i>Vldlr</i>	Very low density lipoprotein receptor	Transmembrane	9p24	19		(30)
<i>Abl</i>	Abelson proto-oncogene	Nonreceptor type tyrosine kinase	9q34	2		(102)
<i>CASK</i>	Membrane-associated guanylate kinase	Cytoplasmic	Xp11	X		(43)
<i>Cdk5</i>	Cyclin-dependent kinase 5	Serine/threonine kinase	7q36	5		(47)
<i>Cdk5r</i>	p35	Cdk5 regulatory subunit	17q11	11		(48)
<i>Cdk5r2</i>	p39	Cdk5 regulatory subunit	2q35	ND		(55)
<i>Dab1</i>	Disabled 1	Cytoplasmic adaptor protein	1q	4		(26,27)
<i>Src</i>	Src proto-oncogene	Non-receptor type tyrosine kinase	20q13	2		(29)
<i>Brn-1</i>	Brn-1	POU domain transcription factor	3p14	1		(103)
<i>Brn-2</i>	Brn-2	POU domain transcription factor	6q16	4		(103)
<i>Emx2</i>	Emx2	Transcription factor related empty spiracles	10q26	19		(45)
<i>p73</i>	p73 (Transformation-related protein)	p53 related protein	1p36	ND		(44)
<i>Tbr-1</i>	T-brain 1	Transcription factor related Brachyury gene	2q23-27	2		(46)

^a ND, Not determined.

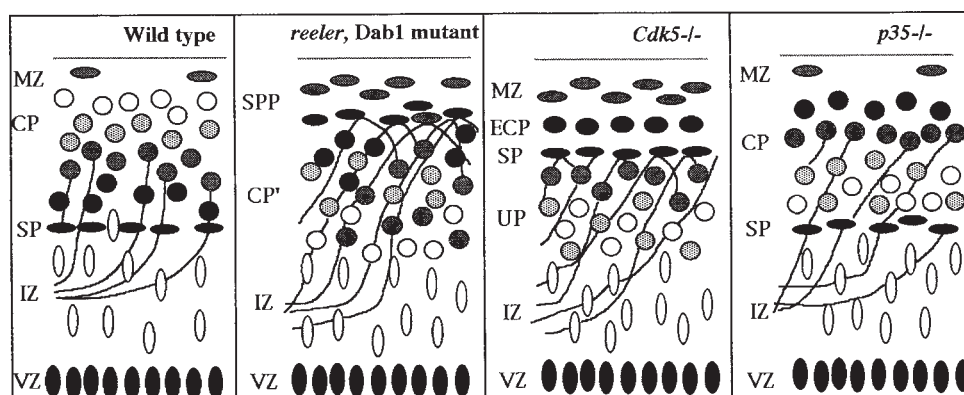


Fig. 1. Schematic representation of the development of the cerebral cortex in wild-type and mutant mice. In wild-type mice, the initial wave of migrating neurons splits the preplate into a marginal zone (MZ) and subplate (SP), and successive waves of migrating neurons move past their predecessors and form the cortical plate (CP) in an inside-out fashion. Marginal zone contains Cajal-Retzius and other cells. The migrating neurons are represented as dark ovals. At the sites where they stopped to form the cortical plate, the neurons are indicated by circles in gray-scale with respect to their final cell division. In *reeler* and *Dab1* mutant mice, migrating cortical plate neurons are incapable of splitting the preplate, which remains as a superplate (SPP) with mutant neurons stacked up in inverted order. In *Cdk5*^{-/-} mice, the initial wave of migrating neurons, destined to become layer VI (indicated by black circles), from the ventricular zone (VZ) cross the intermediate zone (IZ) and split the preplate forming a narrow CP. However, later-born cortical plate neurons (light-gray and white circles) stack up *reeler*-like under the subplate in an inverted fashion termed the 'underplate' (UP). In *p35*^{-/-} mice, initial defects result in an inverted CP similar to that seen in *Cdk5*^{-/-} mice, but later compensatory effects by Cdk5/p39 result in the normal positioning of the subplate neurons. In the adult of *p35*^{-/-} mice there is distinguishable layer I, while in *reeler* no cell-free layer I can be observed in the cerebral cortex.

Cdk5 Kinase

Based on the phenotypes of knockout mice, Cdk5 and its activating subunit p35 are considered to be other critical molecules for neuronal migration (47,48). Cdk5 is a serine/threonine kinase with close homology to other Cdks (49,50) and is a unique Cdk because its kinase activity can be detected mainly in postmitotic neurons (51). Association of Cdk5 with a neuron-specific regulatory subunit, either p35 or its isoform p39, is critical for kinase activity (52,53). *Cdk5*^{-/-} mice exhibit embryonic lethality associated with disruption of the cortical laminar structures in the cerebral cortex, olfactory bulb, hippocampus, and cerebellar cortex (47), whereas *p35*^{-/-} mice showed milder phenotypes than *Cdk5*^{-/-} mice owing to the redundancy of Cdk5 activator isoforms (48,54).

Moreover, *p35*^{-/-}*p39*^{-/-} mice display a phenotype identical to *Cdk5*^{-/-} mice, confirming redundancy in these subunits (55). Neuronal birthdate labeling by BrdU revealed an inverted pattern of cell layers in the cerebral cortex in *Cdk5*^{-/-} mice (56). While *Cdk5*^{-/-} and *p35*^{-/-} mice demonstrate some similarities with *reeler* and *scrambler/yotari* mice, the development of the embryonic cerebral cortex in *Cdk5*^{-/-} and *p35*^{-/-} mice also shows significant differences including the splitting of the preplate (56,57) (see Fig. 1). In *reeler* and *scrambler/yotari* mutants, the migrating cortical neurons appear incapable of splitting the preplate, with cortical plate neurons stacking up in an inverted order beneath the preplate (8,16). In *Cdk5*^{-/-} and *p35*^{-/-} mice, earlier-born neurons split the preplate successfully, and later-born neurons stack up in an inverted

layer under the subplate (56,57). The observations of Morest and others, along with the observations in the analysis of *Cdk5*^{-/-} mice, suggest that the earliest cortical plate neurons (destined for layer VI) may translocate their nucleus within their leading process (58,59), and this migration mode is Cdk5-independent (56). Recent analyses using in vitro slice cultures have supported the view that nuclear (somal) translocation is prominent within the early precursor, while migration guided by radial glia is used by the later-born cortical neurons (59,60). Because the migration of later-born neurons is selectively impaired in *Cdk5*^{-/-} mice (56), it has been suggested that Cdk5 may be involved in the locomotion of migrating neurons along radial fibers (59). The precise role of Cdk5 kinase in neuronal positioning, however, remains to be expounded upon.

Similarities and Differences Between *Reeler* Phenotype and Cdk5-Deficient Phenotype

In addition to the cerebral cortex, *reeler* phenotype and Cdk5/p35 deficiency share similarities and differences in abnormalities of neuronal positioning in other areas of the CNS. With the exception of a few cell types, Reelin signaling and Cdk5 seem to be involved in the neuronal positioning of the same neuronal types, including Purkinje cells in the cerebellum. The cerebellum is another typical region of the CNS where both Reelin/Dab1 deficiency and Cdk5/p35 deficiency result in a similar phenotype at first glance, but detailed analysis has revealed significant differences. *Cdk5*^{-/-} mice exhibit a cerebellar abnormality that resembles *reeler* and Dab1 mutants (47,61). *Cdk5*^{-/-} ↔ *Cdk5*^{+/+} chimeric mice were used to analyze the role of Cdk5 in the neuronal positioning in the cerebellum because most cerebellar morphogenesis occurs after birth, and the *Cdk5*^{-/-} mice die in the perinatal period (61). Analysis of the cerebellum of these chimeric mice demonstrated that Cdk5 defi-

ciency arrests the migration of Purkinje cells in a cell-autonomous manner (61). This is in contrast to the case of normal ↔ *reeler* chimeric mice in which migration of *reeler*-derived Purkinje cells are rescued (62). Cdk5 deficiency also results in a failure of granule cells to complete migration from the external to the internal granule cell layer of the cerebellum (61). A small but significant number of mutant Purkinje cells successfully complete migration to the Purkinje cell layer in *reeler* and *scrambler/yotari* (10,54,63), whereas none of the *Cdk5*^{-/-} Purkinje cells in the chimeras migrate successfully to the cerebellar cortex (61). This finding suggests that the Cdk5 mutation causes a more severe arrest of Purkinje cell migration than the *reeler* or Dab1 mutation. Cdk5-deficiency also results in significant defects in the migration of granule cells (61), whereas defective positioning of granule cells is considered to be a secondary phenomenon in *reeler* and Dab1 mutants.

Relation Between Reelin Signaling and Cdk5/p35

Because of the phenotypic similarities and differences between Cdk5/p35 and Reelin/Dab1 mutants, several models have been proposed regarding the relation between Reelin/Dab1 signaling and Cdk5/p35 (64,65). There is however, no evidence that Cdk5/p35 is a downstream effector of Reelin/Dab1 signaling. The close chromosomal proximity of loci for *cdk5* (66) and *reeler* on chromosome 5 (67) raises concerns that the *reelin* gene might be down-regulated owing to an insertion-induced alteration in a long-distance enhancer located in the vicinity of *cdk5*. However, no differences in the levels of expression of *reelin* and Dab1 mRNA were found between wild-type and *Cdk5*^{-/-} mouse brains at E16.5 (56), confirming that the migration defect of *Cdk5*^{-/-} neurons is not mediated by changes in the expression of either Reelin or Dab1. To investigate the relation between Reelin signal-

ing and Cdk5/p35, p35 and Reelin/Dab1 double-mutant mouse lines were established (54). The deteriorations of the migration defects in these double mutants are typically seen in the Purkinje cells in the cerebellum and the pyramidal neurons in the hippocampus (54); these synergistic effects between two pathways have also been observed in neuronal positioning in the hindbrain. Migration arrest of facial branchiomotor (fbm) neurons and lack of typical alignment of the inferior olive (IO) have been observed in *Cdk5*^{-/-} mice (68). Interestingly, abnormalities in the positioning of fbm and IO neurons have been observed in the *reeler* and Dab1 mutant, and are more severe in the double-null mice for p35 and Dab1 (68). These findings indicate that Cdk5/p35 is not a downstream effector of Reelin signaling, but that Cdk5/p35 and Reelin/Dab1 synergistically contribute to the positioning of the neurons in the developing brain. However, we also find that the addition of heterozygosity of the Dab1 mutation to the *p35*^{-/-} genotype results in extensive migration defects of the granule and Purkinje cells in the cerebellum (54). Therefore, Reelin signaling can influence Cdk5 kinase activity even though Reelin signaling and Cdk5 do not lie in a single cascade. Alternatively, this can also be explained by a regulatory function of Cdk5 in Reelin signaling. Interestingly, Cdk5 phosphorylates serine/threonine sites in the C-terminal region of Dab1 in vitro (cited in ref. 33 as unpublished data). Most recently, mutant mice that express truncated Dab1 lacking the C-terminal region of Dab1 have been generated by a knock-in strategy (69), and analysis of these mutant mice has revealed that a single copy of the truncated gene does not support normal development of the neocortex and hippocampus—suggesting that the C-terminal region of Dab1 p80 is involved in signaling to downstream effector molecules (69). Thus, it will be important to investigate the significance of serine/threonine phosphorylation of Dab1 by Cdk5 in Reelin signaling in vivo.

Possible Function of Reelin Signaling in the Control of Neuronal Positioning

Despite the remarkable progress in the identification of the components of Reelin signaling, the exact function of Reelin in the control of neuronal positioning still remains to be determined. Several hypotheses have been proposed to explain the *reeler* phenotype and Reelin function in neuronal positioning. There is no doubt that Reelin plays an important role in the final phases of neuronal positioning. Reelin is expressed by Cajal–Retzius cells in the cerebral cortex, and is secreted at sites where migrating neurons terminate their migration and detach from radial fibers. Since granule cells in the external granule cell layer of the cerebellum secrete Reelin, and migrating Purkinje cells respond to it, it has been suggested that Reelin functions as a stop signal (or a repellent signal) for migrating neurons in the cerebral cortex (39,70), and similarly as a stop signal or chemoattractant for migrating Purkinje cells in the cerebellum (71,72). A recent attempt to resolve this issue has been made by using transgenic mice, in which Reelin is ectopically expressed in the ventricular zone (73). No effect on neuronal positioning in the cerebrum and cerebellum was observed in wild-type background, but ectopic Reelin has actually been shown to rescue some parts of the neuroanatomical abnormalities in *reeler* background (73). Based on this observation, the authors suspect that Reelin might participate in multiple events critical for neuronal positioning, instead of being a simple positional signal as previously thought (73). However, the fact that the transgene produces only small amounts of Reelin (estimated to be 10–20% of endogenous Reelin protein levels) it is difficult to interpret the effects of ectopic-Reelin from the transgene. Another possible function of Reelin has been proposed based on the study of autonomic neurons in the spinal cord (14). It was shown to regulate the migration of sympathetic preganglionic neurons in

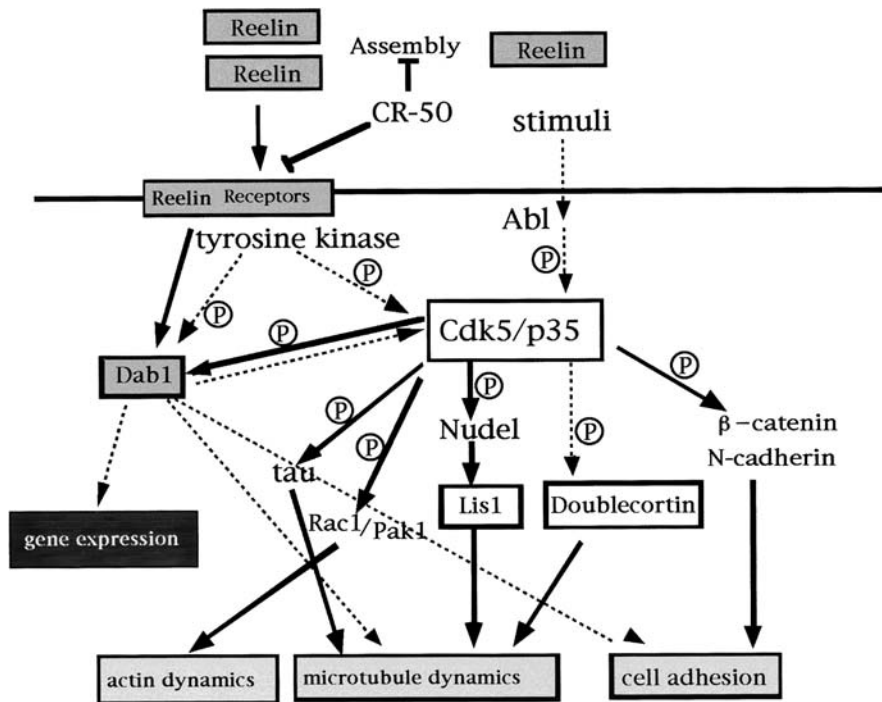


Fig. 2. Molecular model of the signaling pathway of Reelin and Cdk5 in the control of neuronal positioning. Reelin forms homomultimers and binds to its receptors, VLDLR/ApoERs and others. CR-50—a functional blocking antibody to Reelin—is thought to inhibit multimerization of Reelin and binding to receptors. Binding of Reelin to its receptors induces Dab1 tyrosine-phosphorylation, and tyrosine-phosphorylated Dab1 may act as an adapter for the recruitment of various SH2-domain containing proteins and activate downstream signaling cascades (broken lines). Cdk5 phosphorylates Dab1 on serine/threonine residues and may modulate Reelin-induced tyrosine-phosphorylation, or affect Dab1-localization within neurons. Cdk5 phosphorylates cytoskeletal proteins and their associating proteins, which affects the organization of the cytoskeleton. Interaction and phosphorylation of Pak1 by Cdk5 may modulate actin cytoskeleton dynamics. Cdk5/p35 also interacts with β -catenin, resulting in decreased association between β -catenin and N-cadherin, and loss of cell adhesion.

the spinal cord, and based on the observation of a relationship between Reelin and migrating preganglionic neurons, it has been proposed that Reelin may function as a barrier to migrating preganglionic neurons (14).

Possible Function of Cdk5/p35 in the Control of Neuronal Positioning

The results of several studies have suggested that Cdk5/p35 kinase regulates actin dynamics and/or microtubules. The localization of Cdk5 and p35 in the growth cone and interac-

tions with Rac1 and Pak1 suggest that Cdk5/p35 kinase may be involved in the modulation of actin cytoskeletal dynamics (74). Cdk5/p35 may also regulate microtubules, because microtubule-associated protein tau and MAP1b are good substrates of Cdk5/p35 kinase (75,76). Interestingly, the causative genes of human lissencephaly, *Lis1* (77) and *doublecortin* (78,79) are implicated in microtubule reorganization. Possible involvement of Cdk5 in *Lis1* function in the neuronal migration has been discussed, because *Nudel* is one of the *Aspergillus* NUDE gene homologs and interacts with *Lis1*, and is also an in vivo sub-

strate of Cdk5/p35 (80,81). Another possible role of Cdk5/p35 kinase is regulation of cell adhesion mediated by N-cadherin during cortical development. Cdk5 has been shown to negatively regulate N-cadherin-mediated adhesion via an association between p35 and β -catenin (82). This has led to the proposal that Cdk5 kinase activity functions to suppress N-cadherin-mediated adhesion in migrating neurons during corticogenesis. An influence of Reelin signaling on Cdk5 kinase activity, resulting in abrogation of N-cadherin-mediated adhesion, has also been proposed as a possible function of Reelin (65,82). This hypothesis is consistent with the observation that early generated cohorts of neurons in *reeler* mice are more adhesive than their normal counterparts (83,84).

Other Functions of Reelin Signaling and Cdk5 Besides Neuronal Positioning in Developing CNS

Soriano and colleagues have shown that Reelin is important for correct formation of synaptic circuits in the hippocampus (85,86). *Reeler* mice exhibit several alterations in entorhinohippocampal projections, including reduced axonal branching, a decreased number of synapses, and abnormal topography of synapses (85,86); exposure of entorhinohippocampal slices to CR-50 leads to the formation of a *reeler*-type abnormality. Rice et al. demonstrated involvement of Reelin signaling in the organization of synaptic connections in the retina, where Reelin and Dab1 are expressed in adjacent cell layers in the adult (87). After postnatal day 6, Dab1 was found to accumulate in type AII amacrine cells, and defective synaptic layerings of type AII amacrine cells within the inner plexiform layer was observed in *reeler* and *scrambler* retina (87). These findings suggest that, in addition to its essential role during neuronal migration, Reelin signaling contributes to the proper patterning of synaptic connectivity.

Cdk5^{-/-} and *p35*^{-/-} mice have defects in the fasciculation of several axonal tracts including callosal and thalamocortical axons (56,57) (T. Ohshima, unpublished observation). In *Drosophila*, an increase or decrease in Cdk5 activity results in errors in axon pathfinding and target recognition of motor nerves (88). These observations suggest a role of Cdk5 in axon guidance and targeting. This is another example for the comparison with the *reeler* phenotype. In *reeler*, afferent projections to the visual, olfactory, somatosensory, and motor cortices find their correct targets despite being located in ectopic position (89–91).

Reelin and Cdk5 in Neuronal Degeneration

A recent study detected increased levels of tau phosphorylation in *reeler* and VLDLR/ApoER2 null mice in adulthood (32). Hyperphosphorylation of tau, which leads to microtubule dissociation, has been associated with Alzheimer's disease (AD) (92); and we have found an increased level of phosphorylated-tau in *yotari* brain (T. Ohshima et al., unpublished observation), suggesting that Reelin signaling participates in regulating cytoskeletal protein dynamics in the adult brain. Since it is well-known that the inheritance of the ApoE4 allele is a risk factor in the late onset of AD (93), and binding of Reelin to VLDLR/ApoER2 is inhibited by ApoE3 and ApoE4, but not by ApoE2 (32), it can be hypothesized that Reelin signaling acts antagonistically to a pathway involving ApoE, ApoE receptors, and tau phosphorylation (3). Dab1 has been shown to be capable of binding to their NPxY motif of the cytoplasmic domains of the APP family proteins: APP, APLP1, and APLP2 via the PTB domain (31,94,95). It will be interesting to see whether Reelin signaling affects the generation of amyloid β -protein, which is a proteolytic product of APP.

In addition to its possible physiological function in adult brains as well as in synaptic plas-

tivity, dopamine signaling, and drug addiction [reviewed in ref. 96]), Cdk5 is also considered to be involved in the pathology of neurodegenerative disorders, such as AD (96,97) and amyotrophic lateral sclerosis (ALS) (98). Accumulation of p25—which is a cleavage product of p35 and a stable protein—and elevated Cdk5 activity have been reported in the brains of AD patients (97). Although these observations have not been consistent among investigators (99,100), a proposed model, in which elevated Cdk5 activity causes hyperphosphorylation of tau and neuronal apoptosis, is an appealing explanation of the neuropathology of neurodegenerative disorders with hyperphosphorylation of neurocytoskeletal proteins, such as tau and neurofilament proteins, and neuronal loss.

Conclusion

Further studies are necessary to understand the molecular mechanism of Reelin signaling and Cdk5 in neuronal positioning by analysis of mutant mice, and investigating the signal cascades and molecular interactions within migrating neurons in vivo and in vitro. Identification of the downstream cascade of Reelin signaling and in vivo substrates of Cdk5 involved in neuronal migration will provide valuable insights into the molecular and cellular mechanism of the neuronal migration underlying the precise positioning of the neurons in brain development.

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