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# **Requirement of mesodermal retinoic acid generated by** *Raldh2* **for posterior neural transformation**

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# **Abstract**

Studies in amphibian embryos have suggested that retinoic acid (RA) may function as a signal that stimulates posterior differentiation of the nervous system as postulated by the activationtransformation model for anteroposterior patterning of the nervous system. We have tested this hypothesis in retinaldehyde dehydrogenase-2 (*Raldh2*) null mutant mice lacking RA synthesis in the somitic mesoderm. *Raldh2<sup>-/-</sup>* embryos exhibited neural induction (activation) as evidenced by expression of *Sox1* and *Sox2* along the neural plate, but differentiation of spinal cord neuroectodermal progenitor cells (posterior transformation) did not occur as demonstrated by a loss of *Pax6* and *Olig2* expression along the posterior neural plate. Spinal cord differentiation in  $\text{Radh2}^{-/-}$  embryos was rescued by maternal RA administration, and during the rescue RA was found to act directly in the neuroectoderm but not the somitic mesoderm. RA generated by *Raldh2* in the somitic mesoderm was found to normally travel as a signal throughout the mesoderm and neuroectoderm of the trunk and into tailbud neuroectoderm, but not into tailbud mesoderm. *Raldh2<sup>-/-</sup>* embryos also exhibited increased *Fgf8* expression in the tailbud, and decreased cell proliferation in tailbud neuroectoderm. Our findings demonstrate that RA synthesized in the somitic mesoderm is necessary for posterior neural transformation in the mouse and that *Raldh2* provides the only source of RA for posterior development. An important concept to emerge from our studies is that the somitic mesodermal RA signal acts in the neuroectoderm but not mesoderm to generate a spinal cord fate.

# **Keywords**

Neuroectoderm; Spinal cord; Tailbud; Retinoic acid; *Raldh2*; *Fgf8*; *Cyp26a1*; *Olig2*; Mouse

# **1. Introduction**

A key objective of embryology is to understand the cell–cell signaling pathways regulating generation of differentiated tissues from undifferentiated stem cells. In amniote vertebrate embryos, the primitive ectoderm (epiblast) consists of a pluripotent embryonic stem cell population which differentiates during primitive streak formation (gastrulation) to produce the three primary germ layers (embryonic ectoderm, mesoderm, and endoderm). As development proceeds, the primitive streak stem cell zone regresses posteriorly and cells emerging from the primitive streak become further differentiated and acquire progressively more posterior characteristics. The activation–transformation model for differentiation of neuroectoderm from primitive ectoderm suggests that an activation event occurs (neural induction) followed by a transforming event which posteriorizes later-forming neuroectoderm (Nieuwkoop,

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1952). Neural induction of anterior ectoderm to form prospective brain has been found in amphibian embryos to occur in response to bone morphogenetic protein (BMP) antagonists produced in underlying mesoderm generated in the primitive streak (Sasai and de Robertis, 1997). The involvement of BMP antagonists in amniote neural induction is less clear, but fibroblast growth factor (FGF) produced in the primitive streak and tailbud has been implicated (Stern, 2002). FGF8 signaling is particularly important in the mouse as it is necessary for posterior neural induction as well as generation of mesoderm during gastrulation (Sun et al., 1999). In addition to these signals, paraxial mesoderm emerging from the regressing primitive streak provides another signal needed for differentiation of the posterior nervous system (Muhr et al., 1999). One function of this paraxial mesodermal factor is to attenuate posterior FGF signaling mediated by *Fgf8* expressed in the tailbud (Bertrand et al., 2000; Del Corral et al., 2002). Recent studies in avian embryos indicate that retinoic acid (RA) is the paraxial mesodermal factor, and evidence was provided that opposing actions of RA and FGF signaling pathways control posterior neuronal differentiation (Del Corral et al., 2003; Novitch et al., 2003). Treatment of amphibian embryos with exogenous RA suggests that this nuclear receptor ligand may be a factor that causes a posterior transformation of the nervous system, thus supporting the activation–transformation model for anteroposterior subdivision of the nervous system (Durston et al., 1989; Sive et al., 1990). However, it remains unclear if endogenous RA functions as the postulated posterior transformation signal or whether it also acts upstream at the neural induction step (formation of neuroectoderm from primitive ectoderm). Also, it is unknown whether the RA that opposes tailbud FGF signaling is synthesized exclusively in the paraxial mesoderm, and to what extent RA can travel from paraxial mesoderm (or other potential sources) to target tissues.

The concept of embryonic tissue differentiation occurring in regions of opposing RA and FGF signals was originally demonstrated in studies of proximodistal outgrowth of chick limb buds (Mercader et al., 2000). In mouse embryos, *Fgf8* is required to generate a distal FGF signal needed for limb outgrowth (Lewandoski et al., 2000; Moon and Capecchi, 2000). A gene critical for mouse limb RA synthesis has also been identified. Mouse gene knockout studies revealed that several overlapping alcohol dehydrogenases catalyze the first step of RA synthesis, oxidation of retinol (vitamin A) to retinaldehyde (Molotkov et al., 2002), whereas a single gene encoding retinaldehyde dehydrogenase-2 (*Raldh2*) is essential for the second step of RA synthesis in most embryonic tissues, i.e. oxidation of retinaldehyde to RA (Niederreither et al., 1999; Mic et al., 2002). Further studies revealed that all-*trans*-RA is the endogenous ligand needed to rescue *Raldh2*–/– embryonic development, and that its isomer 9-*cis*-RA is undetectable and unnecessary (Mic et al., 2003). We have further demonstrated that *Raldh2* is required to generate a proximal RA signal in the lateral plate mesoderm that travels distally into the limb bud during outgrowth (Mic et al., 2004b). Thus, *Raldh2* and *Fgf8* function as the primary genetic factors involved in generating opposing RA and FGF signals needed for limb development. As *Raldh2* is also essential for RA synthesis in the paraxial mesoderm, we have now examined *Raldh2*–/– embryos to test the hypothesis that *Raldh2* is required for posterior neural development and to further examine the mechanism of RA action using a genetic lossof-function model. Our studies indicate that *Raldh2* is not required for neural induction of ectoderm emerging from the tailbud, but that it is required for posteriorization of tailbud neuroectoderm to generate cells characteristic of the spinal cord. We demonstrate that RA generated in the somitic mesoderm by RALDH2 is the only source of RA for posterior development and that RA travels throughout the posterior neuroectoderm (trunk and tailbud) and trunk mesoderm, but not into the tailbud mesoderm. Our studies also revealed that the RA which reaches the tailbud neuroectoderm is required to define the anterior border of *Fgf8* expression and to stimulate proliferation of neuroectodermal progenitor cells. In summary, our findings indicate that somitic RA generated by *Raldh2* is required to drive neuroectoderm to a spinal cord fate and that RA acts directly in posterior neuroectoderm but not mesoderm during this process.

# **2. Results**

#### **2.1. Raldh2 is responsible for all RA activity detected in mouse embryos at E8.5**

Genetic studies have revealed that RA signaling activity in mouse embryos is dependent upon *Raldh2* encoding an aldehyde dehydrogenase that synthesizes RA (Niederreither et al., 1999; Mic et al., 2002), and *Cyp26a1* encoding a P450 that degrades RA (Sakai et al., 2001; Abu-Abed et al., 2001). *Raldh2* is initially expressed during mouse development at E7.5 in the paraxial mesoderm, consistent with a role in posterior but not anterior axis development in late primitive streak stage embryos. The sites of RA synthesis and degradation in the posterior region of an E8.5 wild-type mouse embryo are shown by double hybridization with *Raldh2* and *Cyp26a1* probes (Fig. 1A). *Raldh2* mRNA is localized in the somitic paraxial mesoderm anterior to the tailbud while *Cyp26a1* mRNA exists in the tailbud, with the gap between the two domains encompassing the presomitic mesoderm.

RA activity was examined in embryos carrying the *RARE-lacZ* RA-reporter transgene (Rossant et al., 1991). *RARE-lacZ* expression indicated that RA signaling normally occurs within the *Raldh2* expression domain (plus to a certain extent anterior and posterior), but that all RA signaling activity is lost in  $\text{Rad}h2^{-/-}$  embryos ( $N=3$ ; Fig. 1B–C). It can also be observed that RA activity in wild-type embryos is reduced in the tailbud where *Cyp26a1* is expressed, consistent with its role in RA degradation.

#### **2.2. RA is not required for neural induction**

As ectoderm emerges from the tailbud, *Sox1* and *Sox2*, encoding high-mobility group (HMG) transcription factors, are expressed in ectoderm that has undergone neural induction (Pevny et al., 1998). Thus, *Sox1* and *Sox2* play a role in differentiating neuroectoderm from non-neural ectoderm which will form epidermis. In order to examine whether RA is needed for neural induction, we examined E8.5 wild-type and *Raldh2*–/– embryos for expression of *Sox1* and *Sox2* mRNAs. *Raldh2<sup>-/-</sup>* embryos expressed *Sox1* ( $N=2$ ) and *Sox2* ( $N=2$ ) at relatively normal levels in neuroectoderm, indicating that RA is unnecessary for neural induction (Fig. 1D–G). The expression domains of *Sox1* and *Sox2* appear wider in the hindbrain region of *Raldh2*–/– embryos compared to wild-type due to incomplete closure of the neural tube in the mutant.

#### **2.3. Raldh2 is required for posterior neuronal differentiation**

Posteriorization of neuroectoderm during the late stages of gastrulation allows hindbrain and spinal cord to be differentiated from forebrain and midbrain (Muhr et al., 1999). During the process of hindbrain specification, several *Hox* genes including *Hoxb1* are normally induced, and this is lost in *Raldh2<sup>-/-</sup>* embryos (Niederreither et al., 2000). This demonstrates that RA is required for the early stages of neural posteriorization when hindbrain is forming. Here we have examined the role of RA at later stages of development when the spinal cord is developing. During spinal cord development, overlapping expression of the homeobox transcription factors *Pax6* and *Nkx6.1* in the posterior neural tube are required for the basic helix-loop-helix transcription factor *Olig2* to be expressed all along the anteroposterior axis of the spinal cord in a ventral location (Marquardt and Pfaff, 2001). Genetic studies in the mouse have demonstrated that *Olig2* is required for spinal cord motor neuron differentiation (Lu et al., 2002; Takebayashi et al., 2002). We found that a lack of RA synthesis in *Raldh2*–/– mouse embryos resulted in a loss of *Pax6* (*N*=3) and *Olig2* (*N*=3) expression in the posterior neural tube at E8.5 (Fig. 1H–K). A small expression domain of *Olig2* that normally exists in the forebrain was not eliminated in *Raldh2*–/– embryos (Fig. 1J–K). We also previously reported that the forebrain expression domain of *Pax6* (not shown here) is still retained in *Raldh2*–/– embryos (Mic et al., 2004a). We found that *Nkx6.1* expression remained in both anterior and posterior neural tissues of *Raldh2<sup>-/-</sup>* embryos ( $N=3$ ), indicating that RA is not required for its expression (Fig. 1L–M). Our findings indicate that RA signaling is not needed during neural

induction, but that RA is needed for undifferentiated posterior neuroectoderm to differentiate to a spinal cord fate. This requirement for RA during posterior neural transformation is conserved between mammalian and avian embryos as investigations with quail and chick embryos have also shown that RA signaling is necessary to induce *Pax6* and *Olig2* but not *Nkx6.1* in the spinal cord (Del Corral et al., 2003; Novitch et al., 2003).

#### **2.4. Distribution of RA from its site of synthesis in the somitic mesoderm**

Previous studies suggested that *Raldh2* may be responsible for synthesizing the RA needed for spinal cord motor neuron differentiation as *Raldh2* is expressed in somitic mesoderm adjacent to the neural tube (Del Corral et al., 2003; Novitch et al., 2003). Our genetic studies have now shown that *Raldh2* is responsible for all RA detected posteriorly in mouse embryos at E8.5 and that *Raldh2* is necessary for *Olig2* expression in the spinal cord. Thus, it can be concluded that RA is needed for posterior differentiation in mouse embryos and that *Raldh2* expressed in the somitic mesoderm is the only source of RA synthesis for this process in mouse embryos. However, the extent of RA distribution from sites of *Raldh2* expression during posterior development is unclear. This was examined by directly comparing the expression patterns of *Olig2*, *Raldh2*, and *RARE-lacZ* along the anteroposterior and dorsoventral axes of 10-somite stage mouse embryos (Fig. 2). *Raldh2* expression was observed in the somites with a posterior border in the presomitic mesoderm where the next somite was just beginning to condense (Fig. 2E–H). *Olig2* expression was observed ventrally in the developing motor neuron field adjacent to the *Raldh2* expression domain (Fig. 2A–D). *RARE-lacZ* expression was observed at high levels in the trunk mesoderm and neural tube and at lower levels further posterior in the tailbud (Fig. 2I–L). Transverse sections showed that *RARE-lacZ* is expressed throughout the dorsoventral axis of the neural tube adjacent to somite 5 and somite 9, thus demonstrating that RA synthesized by *Raldh2* in the somites can travel to all portions of the posterior neural tube (Fig. 2F,G,J,K). Transverse sections also revealed that *RARE-lacZ* expression occurs at lower levels in the tailbud neuroectoderm and endoderm but not mesoderm (Fig. 2L). These findings provide evidence that RA may function cell non-autonomously during posterior neuronal differentiation, with *Raldh2* generating RA in the somitic mesoderm that travels to the neural tube where it may induce *Pax6* and *Olig2*. As RA was not localized to any particular region of the neural tube, this may explain its ability to induce not only *Olig2*, which is limited to the developing motor neuron domain, but also *Pax6* which is expressed more widely across the dorsoventral axis of the neural tube (Marquardt and Pfaff, 2001). Also, we conclude that the RA generated by *Raldh2* in the somitic mesoderm travels posteriorly to nascent tailbud neuroectoderm, suggesting that RA may also function in this location.

# **2.5. Maternal administration of RA rescues Olig2 expression via a direct effect on the neuroectoderm**

We have previously shown that development of  $Raldh2^{-/-}$  embryos can be substantially rescued by a low dose maternal dietary RA supplement that returns embryonic RA levels to approximately the same level observed in wild-type embryos (Mic et al., 2003). We found here that *Olig2* expression was rescued in E9.5 *Raldh2*–/– embryos following such maternal RA administration starting at E6.75. Progressively more *Olig2* expression was observed with longer RA treatments such that treatment ending at E7.75 resulted in very little *Olig2* expression at E9.5 (*N*=2), treatment ending at E8.5 provided substantial expression of *Olig2* at E9.5 (*N*=2), and continuous treatment up to the point of analysis at E9.5 gave results indistinguishable from untreated wild-type controls (*N*=2) (Fig. 3A–D). It can be concluded that continuous RA signaling is needed to induce *Olig2* expression as new neural tube tissue is generated from the tailbud.

In order to examine the mechanism whereby maternal RA rescues embryonic *Olig2* expression, we examined *RARE-lacZ* expression in E8.5 *Raldh*2<sup>-/-</sup> embryos following low dose maternal

dietary RA administration from E6.75 to the point of analysis at E8.5. A transverse section of such an embryo shows that RA signaling activity is occurring throughout the neural tube and endoderm, but not in the mesoderm (Fig. 3E). Interestingly, *RARE-lacZ* expression was not observed in the somitic mesoderm of *Raldh2*–/– embryos subjected to this dietary RA rescue (*N*=4), indicating that the low dose of exogenous RA entering the embryo was not distributed evenly (Fig. 3E); compare to pattern of endogenous RA in wild-type embryos (Fig. 2K). This phenomenon is clearly dose-dependent as large doses of exogenous RA have previously been shown to induce *RARE-lacZ* in all cells of the embryo at E8.5 (Rossant et al., 1991;Mic et al., 2002). As the normal site of RA synthesis in the somitic mesoderm did not exhibit RA signaling activity in *Raldh2<sup>-/-</sup>* embryos under these rescue conditions, this suggests that RA does not need to act in the mesoderm during posterior neuronal differentiation.

#### **2.6. RA generated by Raldh2 limits the anterior border of Fgf8 expression in the tailbud**

*Fgf8* is normally expressed in the tailbud, but its expression is extinguished anteriorly as cells exit the tailbud (Bertrand et al., 2000; Del Corral et al., 2002). There have been conflicting results concerning the ability of RA to effect posterior *Fgf8* expression, with results in avian embryos suggesting that RA reduces *Fgf8* expression in the tailbud (Del Corral et al., 2003), and results in *Xenopus* embryos suggesting that RA induces *Fgf8* expression in the tailbud (Moreno and Kintner, 2004). Our observation that RA synthesized in the somitic mesoderm of mouse embryos enters the tailbud stimulated us to examine this tissue more closely in *Raldh* $2^{-/-}$  embryos. We performed double hybridizations to examine expression of *Fgf8* and *Uncx4.1* in the same embryo; *Uncx4.1* is a marker for the posterior portion of each somite (Mansouri et al., 1997). Examination of E8.5 *Raldh2*–/– and wild-type embryos matched for somite number demonstrated that the somites were smaller and more densely packed in the absence of RA, that the distance from the posterior end of the embryo to the most recently formed somite was increased, and that *Fgf8* expression extended further anterior in the tailbud  $(N=3; Fig. 4A)$ . When matched for somite number,  $Raldh2^{-/-}$ embryos appeared overall slightly larger than wild-type embryos, with size increases noticed in the tailbud and head, but a size decrease apparent in the trunk where somites are located. This suggests that in the absence of RA the length of the tailbud increases along its anteroposterior axis at the expense of the trunk, with more *Fgf8* expression occurring anteriorly. Our findings thus indicate that RA normally limits the anteroposterior extent of *Fgf8* expression in the mouse tailbud, consistent with previous results in avian embryos (Del Corral et al., 2003). Whether this reflects a direct effect of RA on the *Fgf8* gene remains to be determined, but this possibility is supported by our results with *RARE-lacZ* indicating that RA generated by *Raldh2* in the somitic mesoderm normally travels to the tailbud neuroectoderm which expresses *Fgf8* at relatively high levels.

We examined transverse sections through the tailbud of embryos stained for *Fgf8* expression to determine if a lack of RA effected expression in the ectodermal, mesodermal, or endodermal cell populations. *Raldh2*–/– embryos were found to maintain relatively normal expression of *Fgf8* in all three germ layers (Fig. 4C–D). However, these experiments revealed that the mesodermal compartment was expanded in *Raldh2*–/– embryos (Fig. 4D). These findings suggest that a lack of RA results not only in an expansion of the tailbud along its anteroposterior axis, but also an expansion of its mesodermal component along the dorsoventral axis.

#### **2.7. Expression of Cyp26a1 in the tailbud is not regulated by endogenous RA**

We also examined the effect of a loss of RA on *Cyp26a1* expression in the tailbud. Previous studies have indicated that *Cyp26a1* expression in the mouse tailbud can be repressed by excess RA (Fujii et al., 1997), but *Cyp26a1* expression in the *Xenopus* tailbud was found to be induced by excess RA (Moreno and Kintner, 2004). We found that *Cyp26a1* was expressed at a similar level in the tailbuds of E8.5 *Raldh* $2^{-/-}$  and wild-type embryos matched for somite number, with no significant expansion anteriorly into the larger tailbud that forms in the absence of RA

 $(N=3; Fig. 4B)$ . Also, *Raldh* $2^{-/-}$  embryos were found to maintain relatively normal expression of *Cyp26a1* in all three germ layers of the tailbud (Fig. 4E–F). Our loss-of-function findings thus indicate that endogenous RA is unnecessary for *Cyp26a1* expression in the tailbud, and unnecessary to limit the anterior border of *Cyp26a1* expression. We suggest that the previous observations of RA-mediated alterations in posterior *Cyp26a1* expression are the results of the use of excess exogenous RA and do not reflect endogenous RA action. In support of this suggestion, recent studies show that zebrafish *Cyp26a1* expression is responsive to exogenous RA but not endogenous RA generated by *Raldh2* (Dobbs-McAuliffe et al., 2004).

#### **2.8. RA is required to maintain a high level of cell proliferation in the tailbud neuroectoderm**

In order to determine if a lack of RA altered cell proliferation during posterior development, we examined tailbuds and spinal cords of E8.5 (10-somite) wild-type and *Raldh2<sup>-/-</sup>* embryos with antibodies against phosphohis-tone 3 which is detectable only in cells undergoing mitosis. We found that *Raldh2<sup>-/-</sup>* tailbuds had a significantly lower mitotic index in the neuroectoderm compared to wild-type embryos  $(2.8\pm0.6\% \text{ vs. } 8.1\pm0.5\% \text{; mutant vs. wild-type}; p<0.001)$ , but that the mitotic indices were not significantly different for the underlying mesoderm  $(2.5\pm0.2\%)$ vs.  $3.5\pm0.3\%$ ; mutant vs.wild-type) or endoderm  $(3.5\pm1.2\%$  vs.  $4.5\pm0.4\%$ ; mutant vs.wildtype) (Fig. 5A–B). In contrast, in these same embryos there was no significant difference in neuroectodermal cell proliferation further anterior in the neural tube at the level of somite 9  $(6.2\pm1.1\%$  vs. 7.2 $\pm$ 0.7%; mutant vs.wild-type) (Fig. 5C–D); this provides evidence that the reduction in cell proliferation observed in the tailbud neuroectoderm of E8.5 *Raldh2*–/– embryos is not due to poor health of the embryos. These findings suggest that endogenous RA, which we have shown normally travels from the somitic mesoderm to the tailbud neuroectoderm, is needed to generate a high level of neuroectodermal cell proliferation specifically in the tailbud.

# **3. Discussion**

#### **3.1. RA is required for posterior neural transformation but not for neural induction**

During vertebrate neural development, neuroectoderm becomes differentiated from epidermal ectoderm in the process of neural induction (Sasai and De Robertis, 1997; Stern, 2002). Also, anterior neuroectoderm (forebrain/midbrain) becomes differentiated from posterior neuroectoderm (hindbrain/spinal cord) at an early stage. One mechanism proposed for anteroposterior differentiation is the activation–transformation model in which activation (neural induction) is followed by transformation of the posterior neuroectoderm in response to a posteriorizing signal (Nieuwkoop, 1952). Posterior expansion of the hindbrain upon administration of exogenous RA to amphibian embryos suggested that RA may be a factor normally needed for posterior transformation of the nervous system (Durston et al., 1989; Sive et al., 1990). Exogenous RA also results in an expansion of hindbrain in chick and mouse embryos as previously reviewed (Maden, 2002). Recent studies on avian embryos indicate that spinal cord neuronal differentiation is blocked in the absence of endogenous RA signaling (Del Corral et al., 2003; Novitch et al., 2003). However, these studies did not address whether endogenous RA signaling is needed for neural induction of ectoderm emerging from the tailbud or only for posterior differentiation of neuroectoderm progenitor cells to form neurons of the spinal cord. Also, it is not clear if the role of endogenous RA in neuronal differentiation is conserved in mouse embryos. During mouse development, *Raldh2* is initially expressed at E7.5 in the paraxial mesoderm (Niederreither et al., 1997), consistent with a role in synthesizing RA for patterning the posterior axis in late primitive streak embryos. Our analysis here of *Raldh2*–/– embryos demonstrates that RA is not required for neural induction as expression of *Sox1* and *Sox2* proceeded normally all along the neural plate of embryos lacking RA activity. Thus, although exogenous RA treatment has been reported to increase *Sox1* expression in mouse embryonic stem cells (Guan et al., 2001; Wichterle et al., 2002; Gottlieb, 2002),

endogenous RA is not required for normal induction of *Sox1* in the mouse embryo neural plate. We did find that endogenous RA was required for further differentiation of posterior neuroectoderm as the spinal cord motor neuron marker *Olig2* failed to be expressed all along the posterior neural tube. We found that continuous RA treatment was required to rescue *Olig2* expression in the posterior neural tube of *Raldh2<sup>-/-</sup>* embryos, thus demonstrating that RA continually acts to posteriorize nascent neuroectoderm as it emerges from the tailbud. Our results provide strong support for RA as a posterior transforming factor needed for spinal cord formation as hypothesized in the activation-transformation model of neural development (Nieuwkoop, 1952), and we suggest that this developmental mechanism may be generally applicable to vertebrate embryos.

# **3.2. RA synthesized in the somitic mesoderm acts cell non-autonomously in the posterior neuroectoderm**

The results presented here demonstrate that RA generated by *Raldh2* in the somitic mesoderm performs two cell non-autonomous functions during posterior development: (1) stimulation of cell proliferation in tailbud neuroectoderm and (2) stimulation of neuronal differentiation in neuroectodermal cells exiting the tailbud. The ability of *Raldh2* to perform these two different functions is reinforced by our observation that RA synthesized in the somitic mesoderm travels to the adjacent spinal cord neuroectoderm as well as the more distant tailbud neuroectoderm. These observations were made possible through the use of mouse embryos carrying the *RARElacZ* transgene which allowed us to follow where endogenous RA signaling activity occurs in wild-type embryos, and where RA signaling occurs in *Raldh2<sup>-/-</sup>* embryos rescued by maternal RA treatment. Such experiments have not been possible in avian, amphibian, or fish embryos due to lack of such a transgene. With the tools available for analysis of RA signaling in mouse embryos we have found that *Raldh2* expressed in somitic mesoderm is the only source of endogenous RA for posterior neural development. We also found that maternal dietary administration of a low dose of RA was able to rescue posterior neuronal differentiation in *Raldh2*<sup>-/–</sup> embryos, and interestingly we observed that the RA signaling activity restored to these mutant embryos was limited to the neuroectoderm and endoderm and was not detected in the somitic mesoderm. This could be due to a differential ability of maternal RA to reach different embryonic tissues, or could be due to differing levels of RA degradation or RAbinding proteins in epithelial versus mesenchymal cells. These findings demonstrate a direct requirement for RA in the neuroectoderm during neuronal differentiation and indicate that the RA signal does not need to perform an additional function in the somitic mesoderm in order to initiate *Olig2* expression in the posterior neural tube. A direct role for RA in the neuroectoderm has also been suggested from studies in chick embryos where *Olig2* expression was lost following expression of a dominant-negative RA receptor in the posterior neural tube (Novitch et al., 2003).

#### **3.3. Opposing RA and FGF signals in the tailbud stem cell zone**

A mechanism of tissue differentiation involving opposing RA and FGF signals has been proposed for proximodistal outgrowth of chick limb buds (Mercader et al., 2000) as well as anteroposterior extension of the posterior neural tube and somitogenesis (Del Corral et al., 2003; Moreno and Kintner, 2004). In each case the RA signal was hypothesized to be generated by *Raldh2* expressed in mesoderm and the FGF signal was from *Fgf8* expressed in tailbud ectoderm. Genetic studies in the mouse revealed that the RA signal needed for limb buds is indeed from *Raldh2* and that RA travels from the lateral plate mesoderm to the adjacent limb bud ectoderm where it is needed for correct expression of *Fgf8* in the apical ectodermal ridge (Mic et al., 2004b). As we have now shown that RA normally travels from the somitic mesoderm to the tailbud neuroectoderm, and that its presence there results in a smaller *Fgf8* expression domain compared to  $Raldh2^{-/-}$  embryos, it is now possible to propose that endogenous RA signaling controlled by *Raldh2* performs a direct role in modulating the action

of FGF signaling in the tailbud. In particular, our findings suggest that a balance between RA and FGF signaling in the tailbud stem cell zone may be needed to achieve the appropriate amount of cell proliferation in the developing tailbud neuroectoderm. In particular, we found that endogenous RA generated by *Raldh2* in the somites is released and enters the tailbud neuroectoderm where it is needed to achieve a high level of cell proliferation in this tissue. In contrast, our findings indicate that endogenous RA is not required to regulate neuroectodermal cell proliferation further anterior in the neural tube. Studies on vitamin A deficient quail embryos lacking endogenous RA have also shown that RA is not required to maintain neuroectodermal cell proliferation in the neural tube at the comparable stage examined here (Wilson et al., 2003); the tailbud was not examined in those studies. We speculate that neuroectodermal cells of the neural tube may no longer require RA for cell proliferation as they have moved out of the range of FGF8 signaling which is highest in the tailbud.

We propose that nascent somitic mesoderm that has exited the tailbud sends an RA signal back to the tailbud neuroectoderm that limits *Fgf8* expression and establishes the optimal rate of tailbud neuroectodermal cell proliferation needed to coordinate neural plate and somite formation. This model of mouse embryo posterior development involving opposing RA and FGF signals is consistent with a model proposed for avian posterior neural development (Del Corral et al., 2003), and extends the mechanism by demonstrating that there is a direct effect of somitic RA on tailbud neuroectoderm as well as trunk neuroectoderm. We further suggest that neuroectodermal cells in the tailbud exist in a low RA zone, but upon exiting anteriorly these cells enter a zone of high RA emanating from the directly adjacent somitic mesoderm which stimulates neuronal differentiation (i.e. *Pax6* and *Olig2* expression). Our results provide physiological insight into how RA normally functions in mammalian neural stem cell differentiation which may be useful to develop strategies for cell-based therapies through manipulation of stem cell lines.

# **4. Experimental procedures**

#### **4.1. Mice**

*Raldh2<sup>-/-</sup>* embryos were generated from matings of heterozygous adult *Raldh2<sup>-/+</sup>* mice carrying a targeted deletion of *Raldh2* exons 3 and 4 that we have previously described (Mic et al., 2002). In those studies we demonstrated that *Raldh*2<sup>-/-</sup> embryos develop until E8.5 then undergo developmental arrest and die by E10.5. Mice carrying a *RAREhsplacZ* (*RARE-lacZ*) RA-reporter transgene in which *lacZ* (encoding β-galactosidase) is controlled by a retinoic acid response element (RARE) have been previously described (Rossant et al., 1991). Adult *Raldh2*–/+ mice carrying the *RARE-lacZ* transgene were generated by matings between the two lines, and these mice were used to generate *Raldh2<sup>-/-</sup>* embryos carrying one copy of *RARElacZ*.

# **4.2. Rescue of Raldh2–/– embryos by maternal dietary RA treatment**

Maternal dietary RA rescue of *Raldh2<sup>-/-</sup>* embryos was performed by a slight modification of a previously described method (Mic et al., 2003). Briefly, all-*trans*-RA (Sigma) was dissolved in corn oil and mixed with powdered food at the following concentrations:  $0.1 \text{ mg/g}$  for treatment from E6.75–E7.75; 0.1 mg/g for treatment from E6.75–E8.25; for treatment from E6.75–E9.5 a low dose of 0.1 mg/g was used from E6.75–E8.25 and then the dose was raised to 0.25 mg/g for E8.25–E9.5. Such food was prepared fresh twice each day.

#### **4.3. In situ hybridization and retinoic acid detection**

Whole-mount in situ hybridization was performed as described (Wilkinson, 1992). Detection of RA activity was performed in embryos carrying the *RARE-lacZ* RA-reporter gene by staining

for β-galactosidase activity in situ with X-gal as substrate as described (Rossant et al., 1991). Stained embryos were embedded in 3% agarose and sectioned at 50 μm with a vibratome.

#### **4.4. Cell proliferation assay**

Wild-type and *Raldh2<sup>-/-</sup>* embryos embryos at the 10-somite stage (approximately E8.5) were treated with Dent's fixative (Methanol:dimethylsulfoxide 4:1), endogenous peroxidase activity was quenched by incubation with 5% hydrogen peroxide, and paraffin embedding was performed. Tissue sections  $(7 \mu m)$  were incubated with anti-phosphohistone 3 antibodies as a marker for mitosis (1:200 dilution of 1  $\mu$ g/ $\mu$ l stock antibody giving a final concentration of 0.005 μg/μl; Upstate Cell Signaling Solutions, Lake Placid, NY). Sections were then incubated with peroxidase-linked anti-rabbit secondary antibodies, and finally with diaminobenzidine (DAB) substrate. Total cell numbers were determined by counter-staining with hematoxylin. To obtain mitotic indices (percentage cells undergoing mitosis) for the tailbud and neural tube, the numbers of hematoxylin-stained nuclei and phosphohistone 3-positive nuclei were counted in four sections from two wild-type and *Raldh2<sup>-/-</sup>* embryos. Mitotic indices were determined from the mean cell counts and are presented as mean±SEM; statistical significance was determined for raw data using the unpaired Student's *t* test.

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#### **Fig. 1.**

RA is unnecessary for neural induction but needed for neuronal differentiation. (A) Double hybridization to detect *Raldh2* and *Cyp26a1* mRNAs in an E8.5 wild-type (WT) embryo. (B– C) *RARE-lacZ* expression (RA activity) in E8.5 wild-type and *Raldh2<sup>-/-</sup>* (-/-) embryos; *Raldh2<sup>-/–</sup>* embryos totally lack RA activity. (D–E) *Sox1* and (F–G) *Sox2* mRNAs were examined at E8.5 by whole-mount in situ hybridization; these *Sox* genes are markers of neural induction and were still expressed all along the neural tube of *Raldh2*–/– embryos (*Raldh2*–/– embryos have reduced neural plate folding in the hindbrain region, but neural tube closure occurs in the spinal cord). (H–I) *Pax6* mRNA in wild-type and *Raldh2*–/– embryos at E8.5; the *Raldh* $2^{-/-}$  embryo lacks *Pax6* mRNA in the posterior neural tube, whereas the wild-type embryo exhibits expression throughout the hindbrain and spinal cord. (J–K) *Olig2* mRNA is not detected in the posterior neural tube of an E8.5 *Raldh2*–/– embryo, but expression is still observed in a small domain of the forebrain. (L–M) *Nkx6.1* mRNA is still detected in the brain and posterior neural tube of an E8.5 *Raldh2*–/– embryo with a posterior border of expression at the anterior end of the tailbud (in both wild-type and mutant there is a gap of expression in the hindbrain, with the mutant having a larger gap).

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#### **Fig. 2.**

*Raldh2* functions cell non-autonomously for generation of RA detected in the spinal cord and tailbud. Wild-type embryos at E8.5 (10 somite stage) were compared for expression of the target gene *Olig2* (A–D), for the source of RA synthesis by *Raldh2* (E–H), and for the distribution of RA activity detected with *RARE-lacZ* (I–L). For each, transverse sections are shown at somite 5 (s5), somite 9 (s9), and the tailbud (tb). *Raldh2* expression is limited to the somites, anterior presomitic mesoderm, and anterior lateral plate mesoderm. A high level of RA activity (*RARE-lacZ*) is observed throughout the dorsoventral axis of the neural tube (J,K); the developing motor neuron field (marked by *Olig2* expression) is included within this region of high RA activity (B,C). Weaker RA activity is also detected in the tailbud, but is absent from the mesoderm (I, L). ec, ectoderm; en, endoderm; lpm, lateral plate mesoderm; me, mesoderm; mn, motor neuron field; n, neural tube; psm, presomitic mesoderm; s, somite; tb, tailbud.



#### **Fig. 3.**

Maternal dietary RA treatment rescues spinal cord *Olig2* expression in *Raldh2*–/– embryos. (A) *Olig2* expression in E9.5 wild-type embryo (untreated control). *Olig2* expression is also shown in E9.5 *Raldh2<sup>-/-</sup>* embryos following maternal dietary RA treatment from E6.5–E7.75 (B), E6.75–E8.25 (C), or E6.75–E9.5 (D); progressively more *Olig2* expression along the anteroposterior axis of the spinal cord is observed in *Raldh2*–/– embryos with longer RA treatments. (E) *RARE-lacZ* expression was examined in an E8.5 *Raldh2*–/– embryo treated with RA from E6.75–E8.5; shown is a transverse section through the spinal cord at the level of somite 9 demonstrating that maternally administered RA has reached the neural tube and endoderm, but is absent in the somitic mesoderm. en, endoderm; n, neural tube; s, somite.



#### **Fig. 4.**

*Fgf8* and *Cyp26a1* expression in the tailbud of *Raldh2*–/– embryos. (A) Double hybridization showing *Fgf8* mRNA in the tailbud and *Uncx4.1* mRNA in somites; *Raldh2*–/– embryos exhibit an expansion of the tailbud along its anteroposterior axis characterized by an anterior shift in the border of *Fgf8* expression (wild-type and mutant each have 11 somites with the most posterior somite marked by an asterisk). (B) Double hybridization showing *Cyp26a1* mRNA in the tailbud and *Uncx4.1* mRNA in somites; the *Cyp26a1* tailbud expression domain is not significantly altered in *Raldh2<sup>-/-</sup>* embryos (wild-type has 9 somites and mutant has 8 somites with the most posterior somite marked by an asterisk). (C–F) Transverse sections through the tailbud of embryos stained for *Fgf8* or *Cyp26a1* mRNA (plane of sections depicted by arrows in panels A and B); expression of these genes in the three germ layers is not significantly altered in the mutant; however,  $Raldh2^{-/-}$  tailbuds have an expanded mesodermal zone. ec, ectoderm; en, endoderm; me, mesoderm.



#### **Fig. 5.**

RA stimulates cell proliferation in tailbud neuroectoderm. Shown are transverse sections through the tailbud (A–B) and posterior neural tube at the level of somite 9 (C–D) stained with antibodies against phosphohistone 3 (H3P) to detect cells undergoing mitosis; note the decrease in H3P staining in *Raldh2*–/– tailbud neuroectoderm, but not neural tube neuroectoderm. These sections are representative of those used to determine mitotic indices. ec, ectoderm; en, endoderm; me, mesoderm.