

RESEARCH ARTICLE

Cell delamination in the mesencephalic neural fold and its implication for the origin of ectomesenchyme

Raymond Teck Ho Lee¹, Hiroki Nagai², Yukiko Nakaya², Guojun Sheng², Paul A. Trainor^{3,4}, James A. Weston⁵ and Jean Paul Thiery^{1,6,7,*}

ABSTRACT

The neural crest is a transient structure unique to vertebrate embryos that gives rise to multiple lineages along the rostrocaudal axis. In cranial regions, neural crest cells are thought to differentiate into chondrocytes, osteocytes, pericytes and stromal cells, which are collectively termed ectomesenchyme derivatives, as well as pigment and neuronal derivatives. There is still no consensus as to whether the neural crest can be classified as a homogenous multipotent population of cells. This unresolved controversy has important implications for the formation of ectomesenchyme and for confirmation of whether the neural fold is compartmentalized into distinct domains, each with a different repertoire of derivatives. Here we report in mouse and chicken that cells in the neural fold delaminate over an extended period from different regions of the cranial neural fold to give rise to cells with distinct fates. Importantly, cells that give rise to ectomesenchyme undergo epithelial-mesenchymal transition from a lateral neural fold domain that does not express definitive neural markers, such as Sox1 and N-cadherin. Additionally, the inference that cells originating from the cranial neural ectoderm have a common origin and cell fate with trunk neural crest cells prompted us to revisit the issue of what defines the neural crest and the origin of the ectomesenchyme.

KEY WORDS: Neural crest, Ectomesenchyme, Neural fold

INTRODUCTION

The neural crest of the vertebrate embryo has been defined as the dorsal portion of the neural tube epithelium, which appears when the neural folds elevate and fuse in the midline to form the neural tube (Hörstadius, 1950; Marshall, 1879). During neurulation, cells residing within the neural crest leave the dorsal neural epithelium through an epithelial-mesenchymal transition (EMT) and disperse into the adjacent embryonic interstitial spaces, where they are generally termed migrating neural crest cells. Experiments involving

dorsal neural fold epithelia (Hörstadius, 1950) suggested that the neural crest is the source of multipotent stem cells that give rise to a remarkable diversity of cell types, including pigment cells, neurons and glia of the peripheral and enteric nervous systems. In addition, the neural crest was widely considered to be the source of mesenchymal connective tissues that entered the branchial arches to form components of the craniofacial skeleton and connective tissue of the dorsal fin at the trunk axial levels of fishes and amphibia (Hörstadius, 1950; Le Douarin and Kalcheim, 1999; Raven, 1936).

Grafting studies in avian embryos (Le Douarin and Teillet, 1974; Nakamura and Ayer-le Lievre, 1982) suggested that developmental potential differed among crest-derived cells from different axial levels. Such inferences were based on three assumptions: (1) the entire neural fold epithelial domain was presumed to be neural crest (Dupin et al., 2007; Hörstadius, 1950); (2) these cell populations were initially developmentally homogeneous; and (3) the putative neural crest epithelia involved in these marking and grafting studies at trunk and cranial axial levels were qualitatively the same. Recent publications now question these assumptions (Breau et al., 2008; Weston et al., 2004).

Developmental heterogeneity exists within neural crest-derived populations, at least when they first emerge from the neural epithelium *in vitro* (Henion and Weston, 1997) and *in vivo* (Krispin et al., 2010; McKinney et al., 2013; Nitzan et al., 2013; Shoval and Kalcheim, 2012). Moreover, a population of mesenchyme cells precociously emerges from lateral cranial neural fold epithelium and enters the branchial arches before other cells emerge from the neural tube (Hill and Watson, 1958; Nichols, 1981). This implied early developmental heterogeneity in the cranial neural fold epithelium compared with the trunk, which led to the suggestion that skeletogenic ectomesenchyme might arise from a distinct epithelial domain of the neural fold, designated as ‘metablast’, which, in contrast to trunk neural crest cells, expressed a unique combination of ectodermal and mesodermal markers, such as platelet-derived growth factor receptor alpha (PDGFR α) (Weston et al., 2004). This idea is supported by the finding that these cells were found in established mouse strains that label the ectomesenchyme (Breau et al., 2008).

Studies have yet to directly demonstrate that craniofacial skeletal tissues are formed from the lateral non-neural epithelium of the cranial neural folds (Breau et al., 2008). To test this, we provide a detailed immunohistological and cell fate analysis of the neural fold in the midbrain of both mouse and chicken embryos and show that there are two distinct regions from which cells delaminate. In the midbrain, cells originating from the neural ectoderm labeled through the use of Sox1-Cre give rise predominantly to neuronal derivatives. Direct DiI labeling of corresponding regions within the neural fold in chicken embryos shows that the neural ectoderm gives rise to neuronal derivatives, whereas non-neural ectoderm gives rise to ectomesenchyme. We

¹Institute of Molecular Cell Biology, A*STAR, 61 Biopolis Drive, 138673, Singapore.

²Laboratory for Early Embryogenesis, RIKEN Center for Developmental Biology, 2-2-3 Minatogima-minamimachi, Chuo-Ku, Kobe, Hyogo 650-0047, Japan. ³Stowers Institute for Medical Research, 1000 E 50th Street, Kansas City, MO 64110, USA.

⁴Department of Anatomy and Cell Biology, University of Kansas Medical Center, Kansas City, KS 66160, USA. ⁵Institute of Neuroscience, University of Oregon, Eugene, OR 97403-1254, USA. ⁶Cancer Science Institute, National University of Singapore, 14 Medical Drive, 117599, Singapore. ⁷Department of Biochemistry, Yong Loo Lin School of Medicine, National University of Singapore, 8 Medical Drive, 117596, Singapore.

*Author for correspondence (jpthiery@imcb.a-star.edu.sg)

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conclude that, in both species, the cranial neural fold can be broadly divided into two developmentally distinct domains – the neural and the non-neural ectoderm – that undergo temporally distinct episodes of delamination and give rise to neuronal and ectomesenchymal derivatives, respectively.

RESULTS

Cranial neural fold contains two phenotypically distinct epithelial domains and premigratory cells are initially only found in the non-neural ectoderm

During early development, neural induction results in two epithelial domains that can be distinguished within the neural fold: the neural and the non-neural ectoderm. The neural ectoderm in embryos of both mouse and chicken is characterized by the expression of Sox1

and N-cadherin (cadherin 2), whereas the non-neural ectoderm is characterized by the expression of E-cadherin (cadherin 1) (Dady et al., 2012; Edelman et al., 1983; Hatta and Takeichi, 1986; Nose and Takeichi, 1986; Pevny et al., 1998; Wood and Episkopou, 1999). To characterize the neural fold in mouse embryos, we used E-cadherin antibodies to delineate the non-neural ectoderm and Sox9 as a specific marker for cells that are destined to delaminate. At the onset of neurulation at 2 somites, Sox1 was already expressed in the neural ectoderm (Fig. 1Aa,e) and E-cadherin in the non-neural ectoderm (Fig. 1Ac,g). Some residual E-cadherin is found in the Sox1-expressing neural ectoderm, probably owing to the stability of E-cadherin in the entire ectoderm at earlier stages (Carver et al., 2001). However, at this stage, Sox9 (Fig. 1Ab,f) was co-expressed with E-cadherin in the non-neural ectoderm in a restricted region

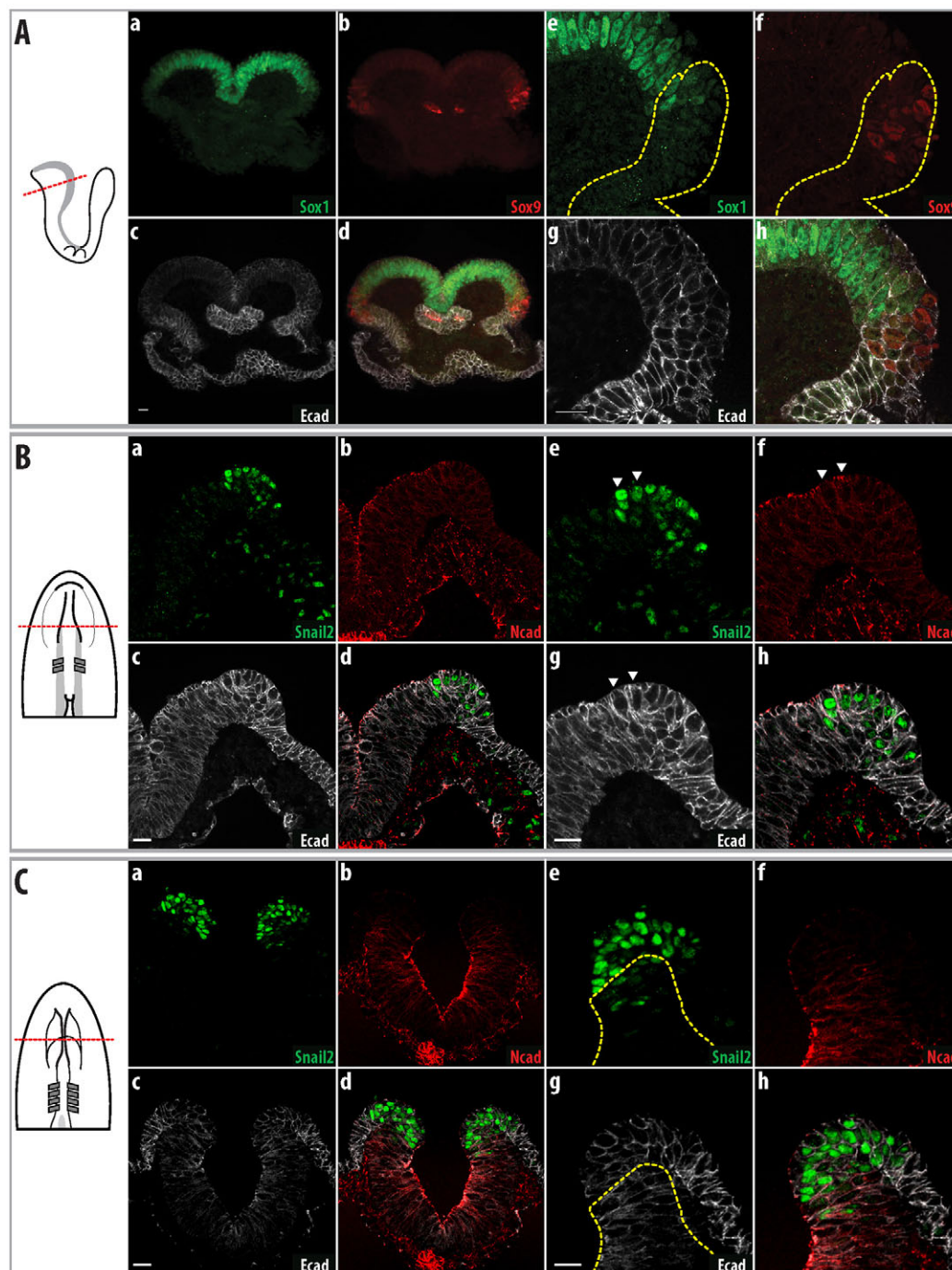


Fig. 1. The cranial neural fold in mouse and chicken embryos contains neural and non-neural ectoderm. At early stages, cells destined to delaminate are only found in the non-neural ectoderm. To the left are schematics of the embryos shown in the images, with the plane of section illustrated. Parts a-d show an overview, whereas e-h show a higher magnification of the neural fold. Neural ectoderm is indicated by the expression of Sox1 (Aa,e) in mouse embryos and N-cadherin in chicken embryos (Bb,f, Cb,f), whereas non-neural ectoderm is indicated by expression of E-cadherin (Ac,g, Bc,g, Cc,g). (A) Two-somite mouse embryo. Sox9 is only expressed in the non-neural ectoderm, which is marked by E-cadherin (Ecad); regions with higher E-cadherin levels are outlined (yellow dotted lines). (B) Two-somite chicken embryo. Snail2 is expressed in the non-neural ectoderm, as marked by E-cadherin. N-cadherin (Ncad) is localized apically in the neural ectoderm. Snail2-expressing cells showing apical staining of N-cadherin (arrowheads). (C) Four-somite chicken embryo. Snail2 is expressed only in the non-neural ectoderm; the region expressing N-cadherin is demarcated by the yellow dotted line. At this stage, N-cadherin is localized apically and laterally in the neural ectoderm, as compared with earlier stages when it was only found apically (B). Scale bars: 20 μ m.

adjacent to, but not overlapping, the Sox1-positive neural epithelium (Fig. 1Ad,h; supplementary material Fig. S1A).

We used E-cadherin (L-CAM) and N-cadherin for the non-neural and neural ectoderm, respectively, and Snail2 to identify cells that were fated to delaminate. N-cadherin was co-expressed with Sox1 in the neural ectoderm in chicken embryos (supplementary material Fig. S2). Thus, the expression pattern of N-cadherin in chicken embryos is comparable to that of Sox1 in mouse embryos, and both factors define the neural ectoderm. Snail2 has been shown to be the earliest indicator of delaminating cells in the epithelium (Nieto et al., 1994). At 2 somites (Fig. 1B), E-cadherin was found in both the neural and non-neural ectoderm (Fig. 1Bc,g); cells found medially in the neural ectoderm showed reduced expression of E-cadherin as compared with cells located more laterally in the non-neural ectoderm (Fig. 1Bc; supplementary material Fig. S1Ba). The neural ectoderm also began to express N-cadherin apically (Fig. 1Bb,f). During this time, the majority of Snail2-expressing cells were found in the non-neural ectoderm lying adjacent to the N-cadherin-expressing neural ectoderm (Fig. 1Ba,e), with a few Snail2-expressing cells found in the neural ectoderm. At 4 somites, the neural ectoderm expressed N-cadherin apically and laterally,

whereas the non-neural ectoderm did not express N-cadherin (Fig. 1Cb,f). E-cadherin expression in the neural ectoderm was also diminished compared with the non-neural ectoderm; its expression was found mainly apically (Fig. 1Cg; supplementary material Fig. S1Ba,b). Snail2 was predominantly expressed in a region that does not express N-cadherin but showed reduced E-cadherin expression compared with lateral areas (Fig. 1Cd,h; supplementary material Fig. S1Ba,b). The *in situ* hybridization pattern of E-cadherin staining in the embryos at 4 somites is similar to that seen by immunostaining (supplementary material Fig. S3A) – that is, E-cadherin is found at higher levels in the non-neural ectoderm as well as in the Snail2 expression region.

In summary, prior to the delamination of cells, the neural fold consists of neural and non-neural ectoderm in both mouse and chicken embryos. The majority of cells that are going to delaminate are found in the lateral non-neural ectoderm.

Precocious EMT of cells within the non-neural ectoderm of mouse embryos

At 4 somites in the midbrain region, most delaminating cells expressing Sox9 (Fig. 2Ab,f) continued to co-express E-cadherin in

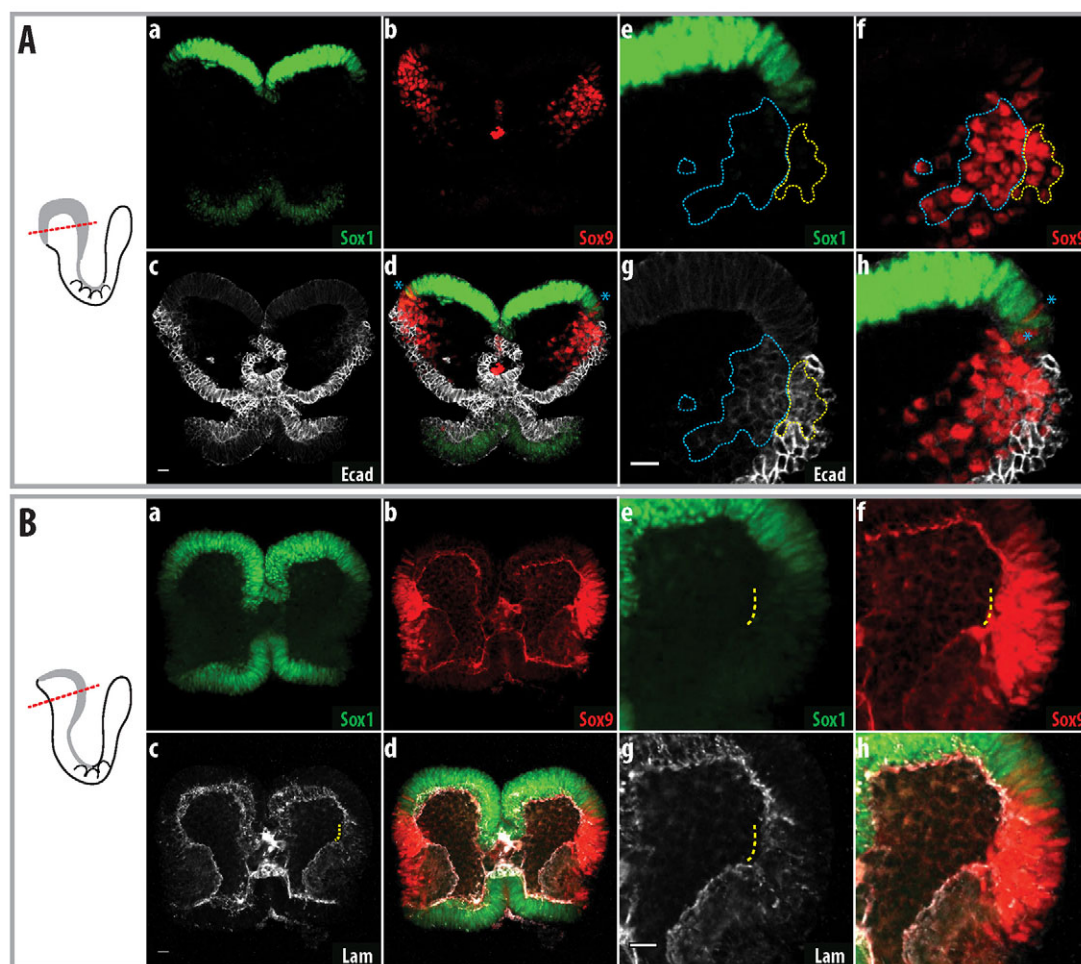


Fig. 2. Cells delaminate from the non-neural ectoderm at ~4 somites in mouse embryos. Delamination of cells in the non-neural ectoderm in mouse embryos. To the left are schematics of the embryos shown in the images, with the plane of section illustrated. Parts a-d of each figure show an overview, whereas e-h show a higher magnification of the neural fold. (A) Four-somite mouse embryo. Sox9 is still expressed in the non-neural ectoderm, as indicated by co-expression with E-cadherin. Sox9-positive cells that have delaminated retain their expression of E-cadherin (at lower levels, blue dotted line). Sox9-expressing cells that co-express low levels of Sox1 are indicated by blue asterisks. Sox9-expressing cells in the non-neural ectoderm are indicated by the yellow dotted line. (B) Three-somite mouse embryo, before cell delamination. Breakage of the basement membrane below the non-neural ectoderm is indicated by the yellow dotted line. Sox9-expressing cells in this region do not express Sox1. Lam, laminin $\alpha 1$. Scale bars: 20 μ m.

the lateral non-neural ectoderm (Fig. 2Ad,h; supplementary material Fig. S1). The dorsal neural ectoderm expressed lower levels of Sox1 than more medial regions (Fig. 2Aa,e; supplementary material Fig. S1) and a few cells appeared with low levels of Sox1 and Sox9 expression (blue asterisk, Fig. 2Ad,h). However, E-cadherin expression in this domain was further reduced and Sox9-positive mesenchymal cells beneath the epithelium showed low levels of E-cadherin within the cytoplasm (blue dotted line, Fig. 2Ac,g); this suggests that these cells have recently delaminated from the E-cadherin-expressing epithelium (see also Breau et al., 2008). To further determine the origin of early delaminating cells, we examined the integrity of the epithelium via the basement membrane component laminin $\alpha 1$. Prior to the delamination of any cells (3–4 somites) there were gaps in laminin $\alpha 1$ immunostaining (yellow dotted line, Fig. 2Bc,g) in the region that was negative for Sox1 (Fig. 2Ba,e). At premigratory stages, Sox9 was expressed only in the non-neural ectoderm (Fig. 1A); this suggests that cells are directly delaminating from the non-neural ectoderm.

Cells in the non-neural ectoderm delaminate first in chicken embryos

In mouse, we have shown that cells are delaminating from the non-neural ectoderm. However, because of the topology of the chicken neural fold, it is not known whether the Snail2-expressing cells in the non-neural ectoderm also delaminate in chicken embryos or how this would occur. In a previous study, it was shown that there is a general lateral-to-medial movement during neural tube closure in the midbrain region (Brouns et al., 2005). To test this idea, we labeled the non-neural ectoderm with Dil and recorded the movement of these cells during neural tube closure. Supplementary material Movie 1 shows the general lateral-to-medial movement of cells during neural tube closure, which is consistent with a previous study (Brouns et al., 2005), but more importantly, also shows that cells in the non-neural ectoderm undergo EMT and disperse into the embryo.

Epithelial cells are polarized, as revealed by the apical and basal localizations of zona occludens 1 (ZO1; also known as Tjp1) and β -dystroglycan (dystroglycan 1), respectively. During EMT, this apical/basal polarization of cells is lost. At 5 somites, when the two neural folds become juxtaposed, we found that Snail2- and E-cadherin-expressing cells (Fig. 3Aa) within the neural fold (Fig. 3Ad) exhibit reduced apical expression of ZO1 compared with cells found more ventrally or laterally (yellow dotted lines, Fig. 3Ae). Similarly, β -dystroglycan (Fig. 3Bb) was found throughout the cortex (yellow dotted area, Fig. 3Be) in these cells. By contrast, Snail2-expressing cells found more ventrally in the neural ectoderm retained their polarized expression of ZO1 and β -dystroglycan (Fig. 3Ae,Be).

Small GTPases control the organization of the cytoskeleton in association with changes in cell morphology and migratory behavior. It was previously shown that RhoB is specifically expressed by neural crest cells but not during the early stages of neural crest formation (Del Barrio and Nieto, 2004; Liu and Jessell, 1998). At 5 somites, RhoB (yellow dotted lines, Fig. 3Cb,e) was only co-expressed in the non-neural ectoderm with Snail2 (Fig. 3Ca) and E-cadherin (Fig. 3Cd), and not in the Snail2-expressing cells situated more medioventrally. The loss of apical/basal polarity and the differential expression of RhoB in Snail2-expressing cells in the non-neural ectoderm indicate that these cells delaminate first. These data, together with the behavioral data of the Dil-labeled cells in the corresponding neural fold epithelium, suggest that RhoB is expressed in the epithelial cells that are the first to delaminate from the neural fold.

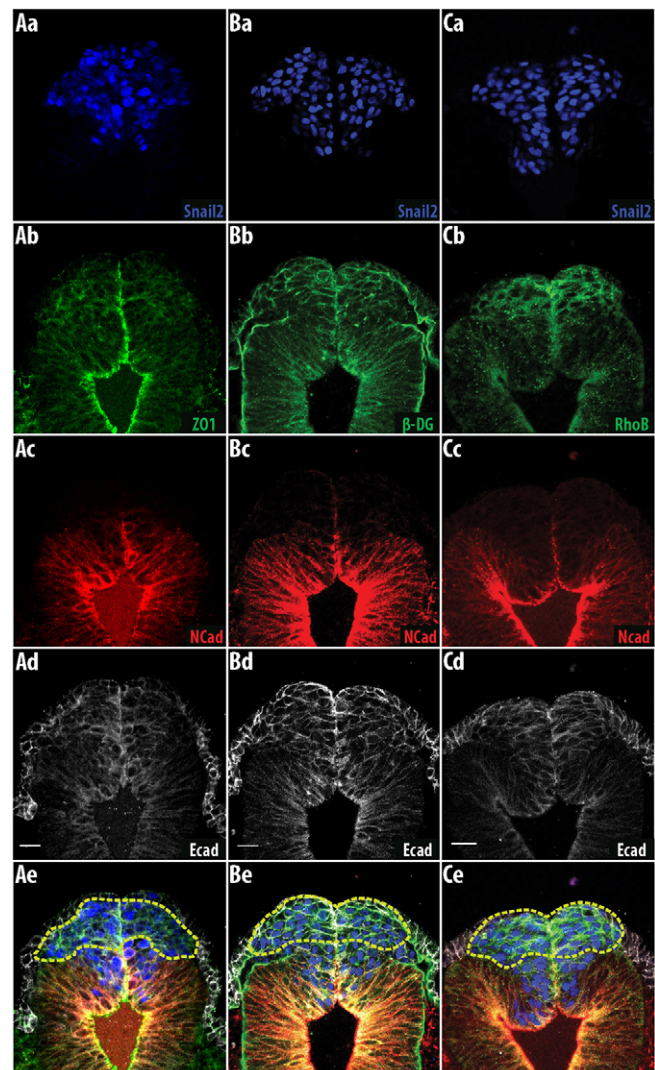


Fig. 3. Snail2-expressing cells in the non-neural ectoderm are depolarized in chicken embryos. Cells in the lateral non-neural ectoderm are depolarized and express RhoB. (A,B) Five-somite chicken embryo. (A) ZO1 is localized apically in Snail2-positive cells within the neural ectoderm, whereas cells in the lateral non-neural ectoderm exhibit reduced apical localization of ZO1. (B) β -dystroglycan (β -DG) is found throughout the cell cortex in the Snail2-expressing cells in the lateral non-neural ectoderm (yellow dotted line), whereas β -dystroglycan in Snail2-expressing cells in the neural ectoderm is basally localized. (C) Six-somite chicken embryo. RhoB is only co-expressed with Snail2 and E-cadherin in the non-neural ectoderm (yellow dotted line). Snail2-expressing cells are found more ventrally and do not express RhoB. Scale bars: 20 μ m.

A second population of delaminating cells originates from the neural ectoderm in cranial regions

Delamination occurs over a prolonged timeframe in cranial regions compared with the trunk region. To determine whether spatiotemporal changes occur during cell delamination, the pattern of expression of Sox9 and Snail2 was established in the neural and non-neural ectoderm, respectively, of mouse and chicken embryos.

The boundary between the neural and non-neural ectoderm was distinct in mouse embryos at 8 somites (Fig. 4A). The dorsal region of the neural ectoderm expressed lower levels of Sox1 than more ventral regions (Fig. 4Aa,e) and Sox9 was expressed in this region of the ectoderm (blue asterisk, Fig. 4Ad,h; supplementary material Fig. S1A) but not in the non-neural ectoderm (Fig. 4Ac,g).

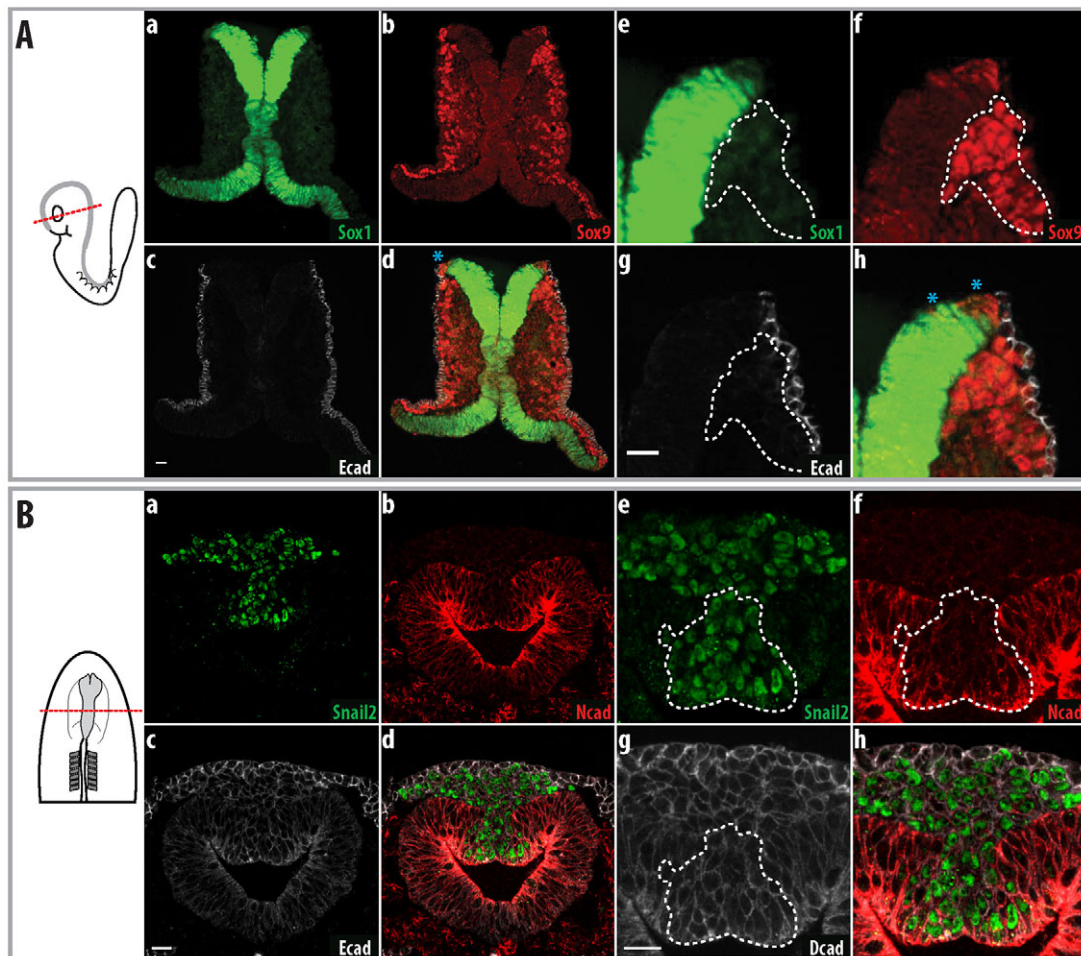


Fig. 4. Cells subsequently delaminate from the neural ectoderm in both mouse and chicken embryos. A second population of cells delaminates from the neural ectoderm. To the left are schematics of the embryos shown in the images, with the plane of section illustrated. Parts a-d show an overview, whereas e-h show a higher magnification of the neural fold. (A) Eight-somite mouse embryo. Sox9 is expressed in the dorsal neural ectoderm that is expressing Sox1; delaminated cells are indicated (white dotted line). (B) Seven-somite chicken embryo. N-cadherin is more strongly expressed in the neural ectoderm, whereas E-cadherin expression in the neural ectoderm is reduced. Snail2 is expressed in the N-cadherin-positive neural epithelia (white dotted line) and in migrating cells juxtaposed between the neural and non-neural ectoderm. Scale bars: 20 μ m.

The neural tube in the midbrain of chicken embryos is closed at 7 somites. N-cadherin was strongly expressed in the neural ectoderm (Fig. 4Bb,f), and lower levels of E-cadherin were found in Snail2-expressing cells and the neural ectoderm (Fig. 4Bc,g; supplementary material Fig. S1Ba,b). At this stage, Snail2-expressing cells were observed between the neural and non-neural ectoderm (Fig. 4Bd,h) and Snail2-expressing cells were found in the neural ectoderm (white dotted line, Fig. 4Bd,h; supplementary material Fig. S1Ba,b). E-cadherin mRNA was only found in the non-neural ectoderm, whereas E-cadherin immunostaining showed that neural ectoderm and delaminated cells are also positive, indicating that E-cadherin protein is stable (supplementary material Fig. S3b). At this stage, Snail2-expressing cells in the neural ectoderm were no longer polarized, as shown by the uniform distribution of ZO1 and β -dystroglycan (yellow dotted line, supplementary material Fig. S4A,B). Furthermore, these cells also began to express RhoB (yellow dotted line, supplementary material Fig. S4C). Similar to the mouse, we noted a second population of delaminating cells in the chicken neural ectoderm.

Orderly delamination of cells in the chicken neural fold

To more definitively show that there are two populations of cells delaminating in the neural fold of chicken embryos and to assess

whether these cells delaminate in an orderly fashion, we labeled different regions of the neural fold and followed their fates shortly after delamination. The non-neural ectoderm at the level of the mesencephalon was labeled by depositing a small drop of DiI onto the non-neural ectoderm using a micromanipulator; care was taken not to pierce the epithelium (supplementary material Fig. S5). When the non-neural ectoderm was labeled (Fig. 5A), these labeled cells were found at the leading edge of migrating cells. By contrast, when the neural ectoderm was labeled, the labeled cells were found at the trailing edge of the migrating cells (Fig. 5B). Fig. 5C shows an embryo in which the cells in the non-neural ectoderm and the neural ectoderm were labeled with DiO and DiI, respectively. These two groups constitute opposite ends of the migrating cells, with little or no cell intercalation. This further shows that cells in the non-neural ectoderm delaminate first, whereas cells in the neural ectoderm delaminate later (Fig. 5D,E).

Cells originating from the neural ectoderm give rise to neurons and glia in mouse embryos

Takashima et al. reported that branchial arch (BA) 1 in Sox1-Cre/YFP embryos was not well labeled when compared with P0-Cre/YFP embryos (Takashima et al., 2007), a transgenic mouse line

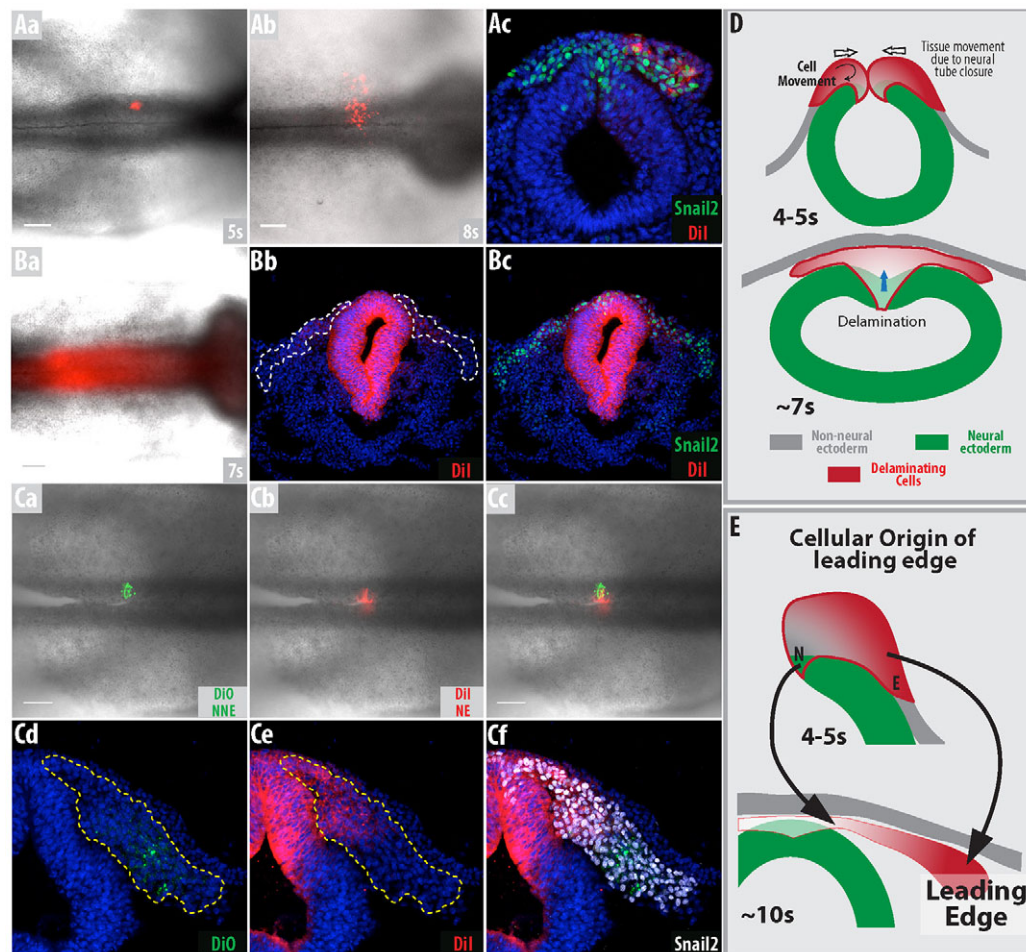


Fig. 5. Cells delaminate from the non-neural ectoderm before the neural ectoderm in chicken embryos. (A-C) Dil and DiO were used to label different regions of the neural fold. (A) Non-neural ectoderm was Dil labeled at 5 somites. The embryos were grown until 8 somites (Ab,c). Labeled cells can be seen at the leading edge of the migrating cells. (B) Neural ectoderm was labeled with Dil at 7 somites. The embryos were grown until 11 somites (Bb,c). Labeled cells are found in the trailing edge of the migrating cells; the white dotted line represents migrating Snail2-positive cells. (C) Labeling was performed at 4 somites and embryos were grown until 10 somites (Cd-f). These two populations of labeled cells do not intermingle. (Cd) DiO-labeled cells form the leading edge of the migrating cells, whereas (Ce) Dil-labeled cells form the trailing edge; yellow dotted line represents migrating Snail2-positive cells. (D,E) Schematics of (D) cell delamination and movement away from the neural fold at 4 and 7 somites based on the results from general tissue movements, loss of basal/apical polarity and the short-term tracing of cells and (E) how they contribute to the migrating mass of cells. The red gradient in the figures represents the two waves of delaminating cells, with red and white representing cells originating from the non-neural and neural ectoderm, respectively. s, somites. Scale bars: 100 μ m.

used to label neural crest derivatives. We showed that Sox1 is expressed in the neural ectoderm (Fig. 1Aa,e) and that the neural ectoderm is YFP-labeled in Sox1-Cre/YFP embryos (supplementary material Fig. S6) prior to cell delamination, suggesting that the neuroectoderm does not give rise to ectomesenchyme.

Cells originating from the midbrain neural folds give rise to cells within BA1 as well as to cells within the trigeminal ganglion (Osumi-Yamashita et al., 1994). In embryonic day (E) 11.5 Sox1-Cre/YFP embryos, there were few YFP-positive cells in BA1 that were also expressing Sox9 (Fig. 6A,C,D), a marker of bone/cartilage precursors (Akiyama et al., 2005; Mori-Akiyama et al., 2003). By contrast, the trigeminal ganglia was relatively well labeled, with the majority of glial cells in the trigeminal ganglia co-expressing YFP and Sox10 (Fig. 6B,E), a glial cell marker (Kuhlbrodt et al., 1998). Furthermore, some neurons within the trigeminal ganglia, which originated from the neural crest, were also labeled and showed co-expression of YFP and HuC/D (Elavl3/4), a neuronal marker (Fig. 6E).

HtPA-Cre/lacZ will label all cells that originate from the neural fold (Pietri et al., 2003). We compared bone/cartilage derivatives in

the skulls of postnatal day 0 mice that had been labeled with HtPA-Cre/lacZ or Sox1-Cre/lacZ. Our results showed that the overall intensity of β -galactosidase was much weaker in Sox1-Cre/lacZ skull preparations than in those labeled with HtPA-Cre/lacZ (Fig. 6F,G). This was especially the case for the mandible, which derives from BA1; the bones and cartilages within this region were not well labeled (Fig. 6F,G).

We further quantified the proportion of bone/cartilage and neuronal precursors in E9.0 Sox1-Cre/YFP embryos (15 somites) after both the trigeminal ganglia (blue dotted lines, Fig. 6H'd) and BA1 (green dotted lines, Fig. 6H'd) had formed. We found far greater numbers of YFP-positive cells in the trigeminal ganglia than in BA1, indicating that Sox1-Cre preferentially labels glial and neuronal derivatives (81.2%, Fig. 6H) compared with ectomesenchyme (14.5%, Fig. 6H). These neural fold-derived cells also expressed Tfp2a (Fig. 6H'b,H'c) as well as Sox9 (Fig. 6H'c,H'c); both of these markers are expressed by migratory cells (Mitchell et al., 1991; Wright et al., 1995). Sox1 was not expressed in the trigeminal ganglion at E9.0 (data not shown), which rules out

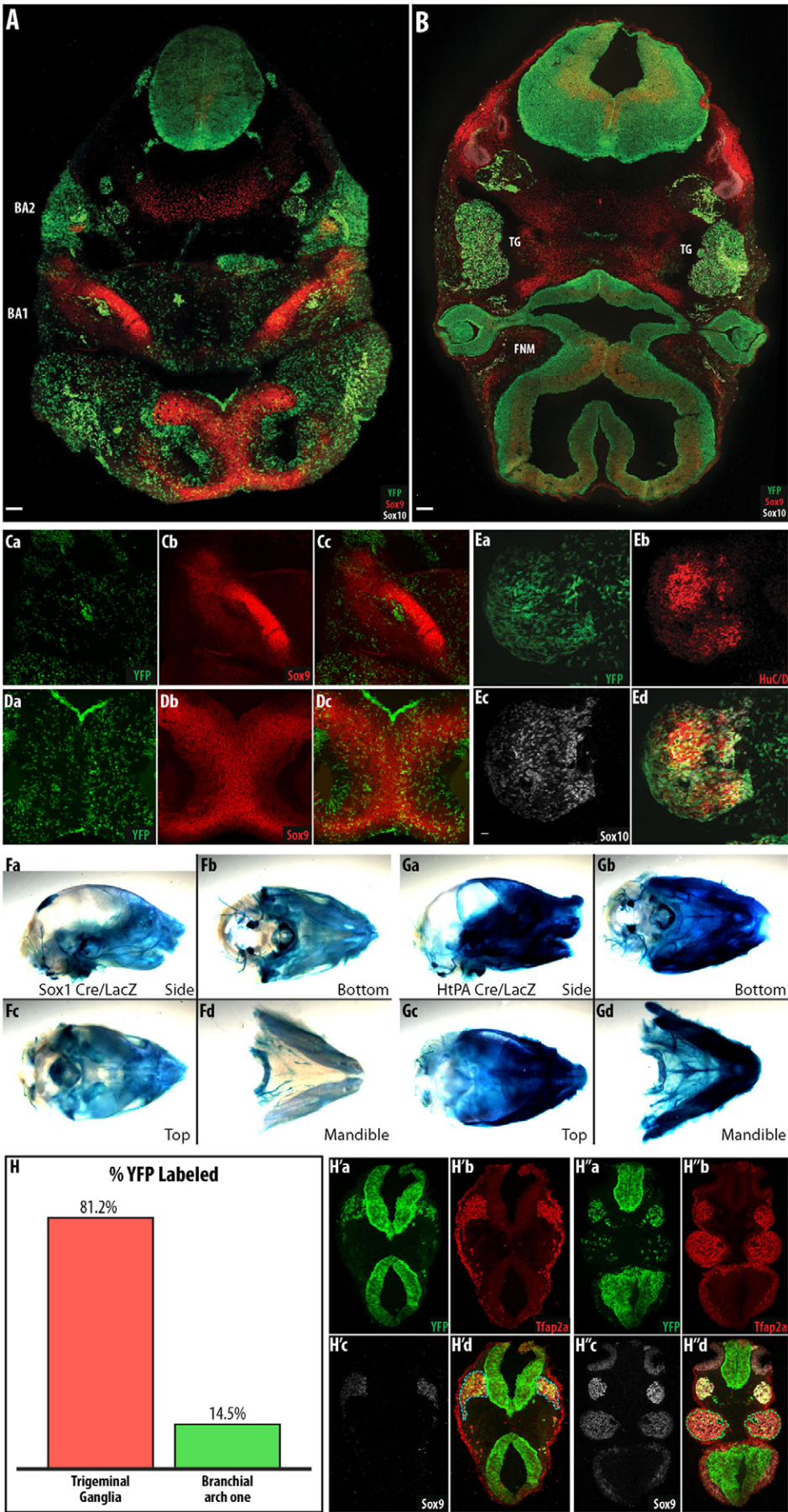


Fig. 6. Neuronal and glial derivatives in the trigeminal ganglia are preferentially labeled in Sox1-Cre. In cranial regions, Sox1-Cre preferentially labels neuronal and glial derivatives compared with the ectomesenchyme. (A-E) E11.5 Sox1-Cre/YFP mouse embryo. Overview of YFP-labeled cells in cranial regions of Sox1-Cre/YFP embryos at the level of BA1 (A) and the trigeminal ganglia (B). (C,D) Cartilage/bone precursors in BA1 (C) and the nasal septum (D) are not well labeled in these regions. (E) More cells in the trigeminal ganglia are labeled with YFP. Sox1-Cre/lacZ (F) and HtPA-Cre/lacZ (G) P0 mouse skull preparations. (H) The percentage of YFP-labeled cells in the anlage of the trigeminal ganglia and BA1 was determined in E9.5 Sox1-Cre/YFP embryos. (H',H'') Representative sections in which the numbers of YFP-labeled and Sox9/Tfap2a-expressing cells were counted in the anlage of the trigeminal ganglia (H') and BA1 (H''). Region counted indicated by blue and green dotted lines for the trigeminal ganglia and branchial arch, respectively. *n*=2. TG, trigeminal ganglia; FNM, frontal nasal process. Scale bars: 40 μ m.

the possibility of labeling because of *de novo* expression of Sox1 in the trigeminal ganglion. Overall, we have shown that the ectomesenchyme in BA1 is not well labeled, whereas the neuronal derivatives are well labeled in Sox1-Cre/YFP embryos.

The incomplete labeling of the trigeminal ganglion prompted us to examine the activation of the YFP reporter at the onset of cell delamination. We found that at 5 somites some YFP-expressing cells are present within the E-cadherin-expressing non-neural ectoderm (blue asterisk, Fig. 7Ad,Bd). Although these cells expressed YFP, they did not express Sox1 protein. As mentioned above, the dorsal region of the neural ectoderm expressed lower levels of Sox1 than the more ventral regions (yellow dotted line, Fig. 7Bd). It is important to note that in Sox1-Cre/YFP embryos the border region between the neural and non-neural ectoderm had a mosaic appearance, i.e. cells were sporadically labeled with YFP. These YFP-labeled cells in the non-neural ectoderm could still be observed at 7 somites.

The formation of BA1 at 8 somites and anlage of the trigeminal ganglia after 10 somites (Chan and Tam, 1988; Theiler, 1989) coincides with the switch in expression of Sox9 from the non-neural ectoderm to the neural ectoderm. We determined the percentage of YFP-labeled migratory cells in Sox1-Cre/YFP embryos as 7.9% at 6 somites and 12.8% at 8 somites (Fig. 7C). At ~10 somites, the cells that had recently delaminated from the neural ectoderm were YFP labeled (Fig. 7D,E), consistent with the percentage of YFP-

labeled cells found in older embryos in this study. The early delaminating cells are most probably derived from the non-neural ectoderm and are ectopically labeled by Sox1-Cre. Thus, although we cannot fully exclude the possibility that some ectomesenchymal derivatives originate from the neural epithelial domain of the cranial neural fold in Sox1-Cre/YFP embryos, these labeled cells most likely originated from cells that ectopically expressed Sox1-Cre in the non-neural epithelial domain.

Non-neural ectoderm-derived and neural epithelial-derived cells give rise to ectomesenchymal cells and neuronal derivatives, respectively, in chicken embryos

We performed DiI labeling of the non-neural and neural ectoderm in chicken embryos to determine the fate of labeled cells 2 days later. In whole-mounts of these embryos, the majority of labeled cells were found in BA1 (Fig. 8A) or in the periocular mesenchyme. Particular cells within the trigeminal ganglia were also labeled with DiI in 2 out of 10 embryos (Fig. 8A,C). When these embryos were sectioned, the DiI-labeled mesenchymal cells within BA1 showed Tfap2a expression (Fig. 8B) (Minarcik and Golden, 2003). It is possible that the DiI-labeled cells within the trigeminal ganglion might be neurons derived from the trigeminal placode, since these two regions are adjacent to each other. However, because of the intermingling of the neurons and glia within the ganglion and the

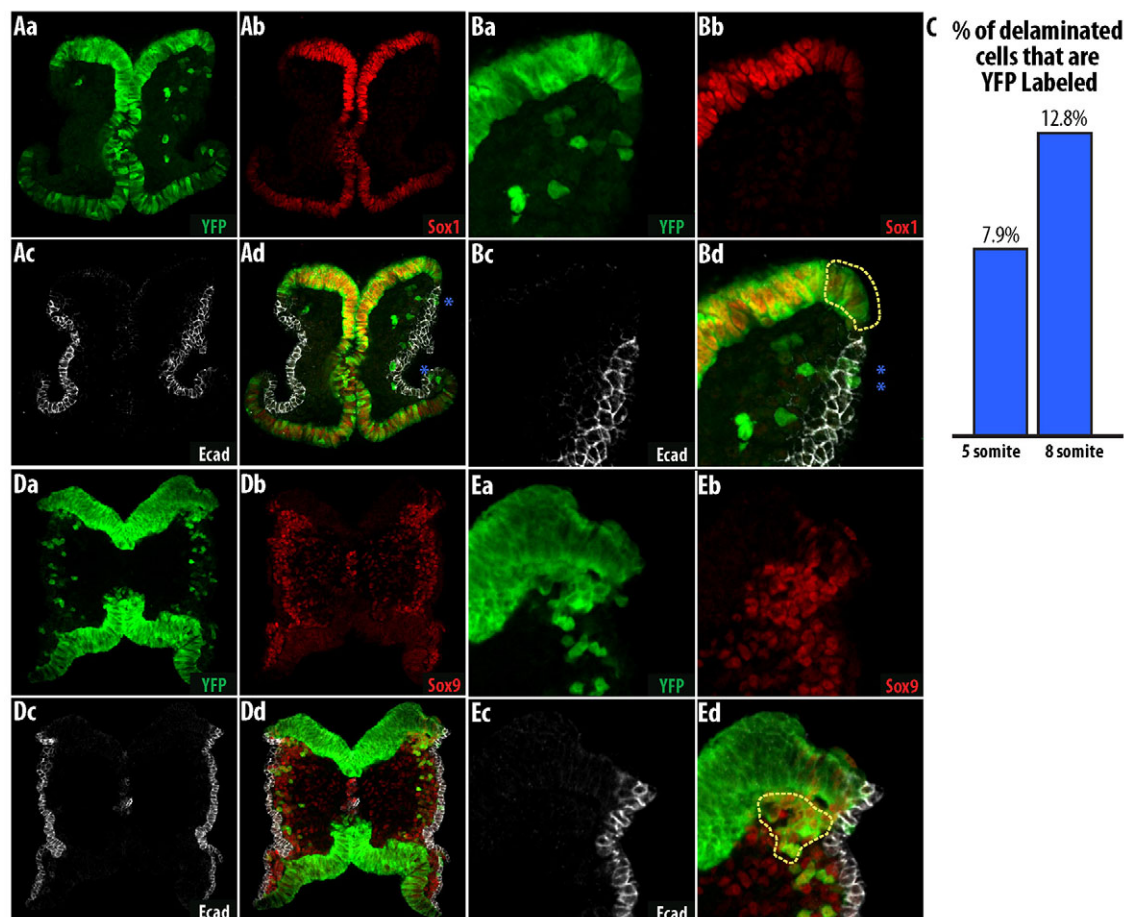


Fig. 7. Sox1-Cre labels some cells in the lateral non-neural ectoderm. The majority of cells labeled in Sox1-Cre originate from the neural ectoderm but there are some ectopically labeled non-neural ectoderm cells. (A,B) YFP-labeled cells in the non-neural ectoderm do not express Sox1 (blue asterisks). YFP-labeled dorsal neural ectoderm expressing lower levels of Sox1 is indicated by the yellow dotted line. (C) There are few YFP-labeled cells in the Sox1-Cre/YFP embryo at the early stages of cell delamination. $n=2$. (D,E) Cells delaminating from the neural ectoderm are better labeled at 10 somites, prior to the formation of the trigeminal ganglia.

punctate DiI signal, we cannot distinguish exactly whether these labeled cells are glia or neurons. To determine whether these cells are derived from the trigeminal placode, we looked at earlier stages when the DiI signal was stronger ($n=14$). Pax3 expression in neural crest cells is lost after 16 somites but Pax3 is expressed by the trigeminal placode and its derivatives (Xu et al., 2008); this provides a means to identify the trigeminal placode as well as its derivatives. In two embryos in which we could observe DiI-labeled cells in the trigeminal ganglia at 20 somites, DiI-labeled cells in the trigeminal ganglia expressed Pax3 (supplementary material Fig. S7A). Furthermore, analysis of non-neural ectoderm-labeled embryos at early stages revealed three embryos with DiI-labeled cells in the Pax3-positive trigeminal placode (supplementary material Fig. S7B,C). Therefore, since we did not consistently observe derivatives in the trigeminal ganglia in all labeled embryos, and as we also demonstrated that the trigeminal placode and its derivatives are labeled in some cases, it is likely that the DiI-positive cells that we observed in the trigeminal ganglion at Hamburger-Hamilton stage (HH) 18 were derived from the trigeminal placode.

Injecting DiI into the lumen of the neural tube at 7 somites only labels cells that originate from the neural ectoderm, as the neural tube is already closed. In whole-mounts of these neural ectoderm-labeled embryos, there were no DiI-labeled cells within BA1 (Fig. 8D). The strong fluorescent signal from the underlying neural tube meant that it was difficult to determine, in whole-mounts, whether labeled cells were present in the trigeminal ganglion. Upon sectioning these embryos, we found numerous DiI-labeled cells within the trigeminal ganglia (Fig. 8E). In this case, because only the neural ectoderm was labeled, these DiI-labeled cells could not have originated from the trigeminal placode.

From these labeling experiments, we conclude that cells within distinct domains of the neural fold give rise to cells with different cell fates. Cells originating from the non-neural ectoderm give rise to mesectodermal derivatives, whereas cells originating from the neural ectoderm give rise to neuronal derivatives.

DISCUSSION

Ectomesenchyme is present in all vertebrates; however, its origin was often debated in early studies because of the considerable variation in the morphogenesis of the head across different classes of vertebrates. In addition, the terms 'dorsal neural tube in the trunk' and 'cranial neural fold' are considered to describe equivalent tissues. Unlike the single episode of delamination in the trunk region, there are two episodes of cell delamination from the neural fold in the midbrain region in both chicken and mouse embryos (summarized in Fig. 9). In cranial regions only the second wave of delaminating cells shares a common origin and cell fate with cells that delaminate in the trunk. Our study provides compelling evidence that these two regions are not equivalent, and that the unique formation of the cranial neural folds may permit the formation of ectomesenchyme.

In the lineage analysis of chicken embryos, we do not address the issue of cell fate potential, as we did not challenge these cells. Furthermore, the small number of Snail2-expressing cells in the neural ectoderm at very early stages could be precursors of cells that delaminate from the neural ectoderm. Below we address a number of issues regarding the definition and pluripotency of these cells at different axial levels.

Origin of ectomesenchyme and topology of the neural fold

The initial study on the origin of the ectomesenchyme showed that it was derived from the lateral non-neural ectoderm (Platt, 1893), a

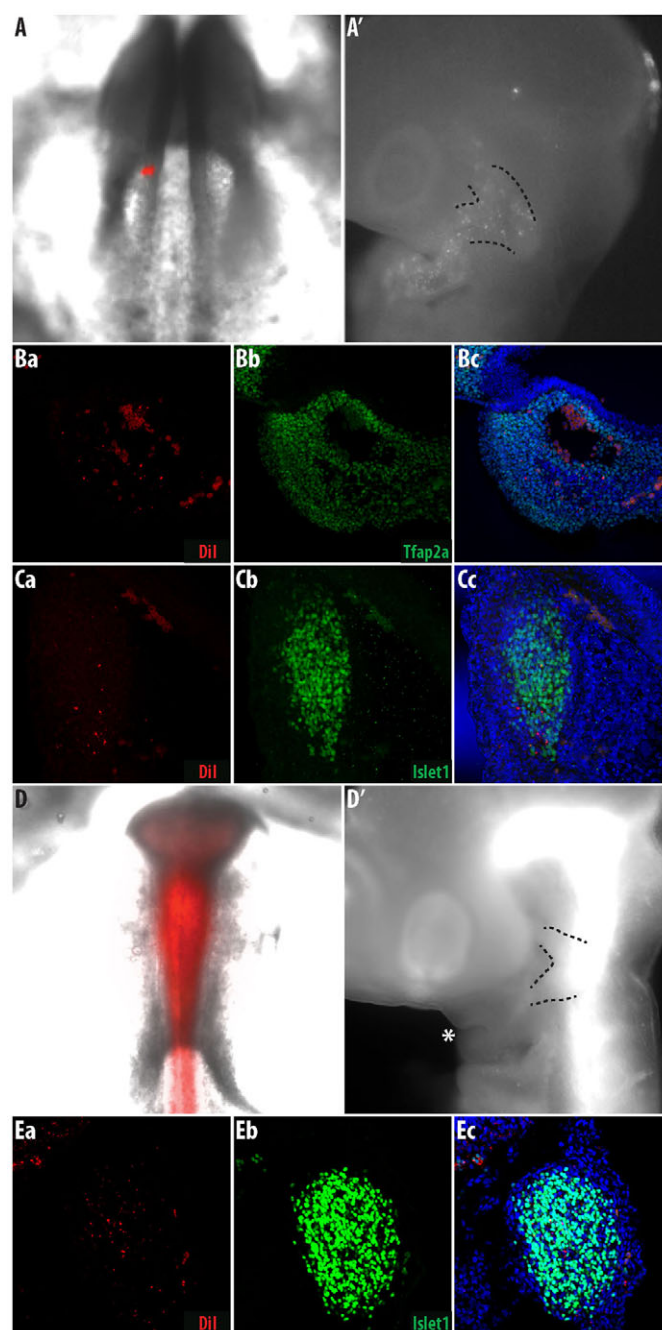


Fig. 8. Two populations of delaminating cells in the neural fold give rise to distinct cell fates. Non-neural ectoderm and neural ectoderm give rise to derivatives in the branchial arches and trigeminal ganglia, respectively. (A-C) Non-neural ectoderm was DiI labeled at 4 somites. All labeled chicken embryos have labeled ectomesenchyme in the pericardial or branchial arch 1 and only two embryos have cells labeled in the trigeminal ganglia; $n=10$. Chicken embryos showing initial DiI labeling of the non-neural ectoderm (A) and labeled derivatives after 2 days of culture (A'). Note that the majority of labeled cells are in the branchial arch (one of two embryos with labeled cells in trigeminal ganglia). Transverse section of (B) branchial arch one (BA1) immunostained for Tfp2a, which is expressed in ectomesenchymal cells and (C) the trigeminal ganglia immunostained for Islet1/2, which is expressed by neuronal cells. (D,E) Neural ectoderm was labeled by DiI at 7 somites; $n=8$. Chicken embryos showing initial DiI labeling of the neural ectoderm (D) and labeled derivatives after 2 days of culture (D'). Note the absence of labeled cells in the branchial arches. (E) Transverse section of the trigeminal ganglia immunostained for Islet1/2. Dotted lines indicated the trigeminal ganglia.

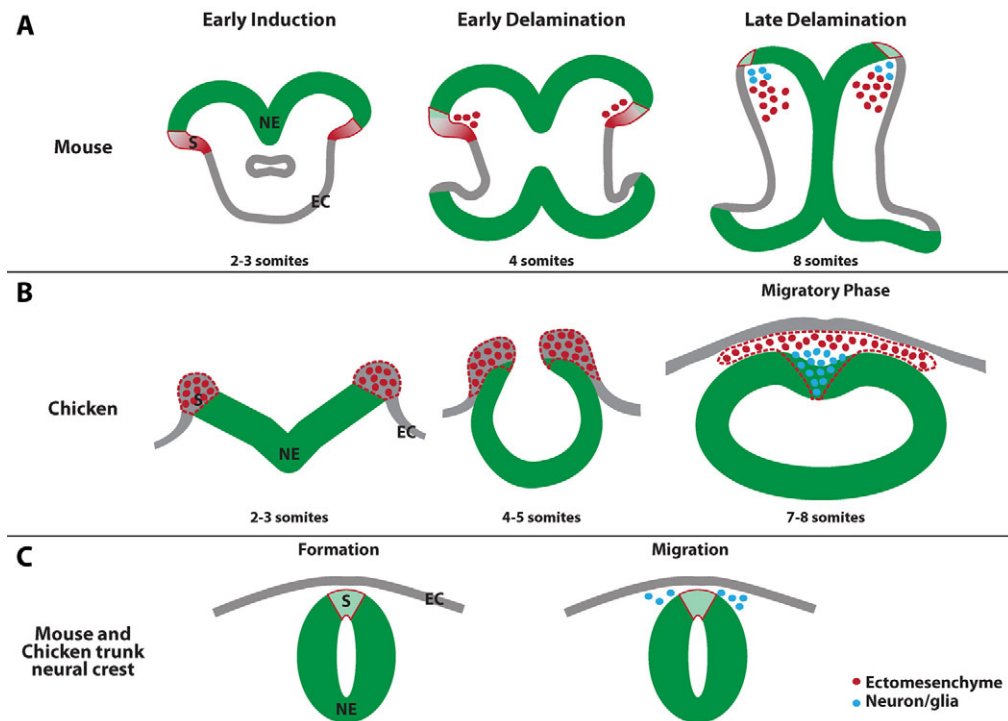


Fig. 9. Model of cell delamination in the midbrain and trunk. There are two waves of cell delamination and these cells adopt distinct cell fates in the midbrain of both mouse (A) and chicken (B) embryos. In the trunk, cells only delaminate when the neural and non-neural ectoderm are already separated (C). EC, non-neural ectoderm; NE, neural ectoderm; S, region where cells delaminate.

region distinct from the neural crest that originated from the dorsal neural ectoderm and gave rise to cranial ganglia. However, subsequent studies dismissed the notion that there were two distinct waves of delaminating cells by conflating the definition of neural crest and neural fold (Hörstadius, 1950; Landacre, 1921). This was probably also complicated by the lateral-to-medial movement of cells during cranial neural tube closure (Brouns et al., 2005). Uncertainties remained as to whether cells were delaminating only from the neural epithelium or also from the overlying non-neural ectoderm, and this was true even in the trunk (Beard, 1888; Hörstadius, 1950). The origin of trunk neural crest from the neural ectoderm was unequivocally shown through grafting of neural tube in avian embryos (Weston, 1963). However, the continuity of the neural and non-neural ectoderm in cranial regions (Johnston, 1966) made it impossible for grafting experiments to determine the origin of cells with sufficient spatial and temporal resolution.

In mammalian embryos, cranial neural tube closure occurs after cell delamination and the neural fold adopts an extended structure with an underlying interstitial space. This creates a scenario in which the origin of delaminating cells can be accurately determined. A number of studies have shown that the lateral non-neural ectoderm gives rise to ectomesenchyme found in the branchial arches in mammals (Breau et al., 2008; Chan and Tam, 1988; Hill and Watson, 1958; Nichols, 1981; Vaglia and Smith, 2003; Weston et al., 2004). We have also observed expression of so-called ‘neural crest specifiers’ such as *Tfap2*, *Pax3* and *Pax7*, in mouse and chicken embryos in both the non-neural and neural ectoderm (data not shown). This would explain why a number of transgenic mouse lines, including *Wnt1-Cre*, *HtPA-Cre*, *Pax3-Cre* and *Pax7-Cre*, label both the neuronal as well as ectomesenchymal derivatives (Danielian et al., 1998; Engleka et al., 2005; Murdoch et al., 2012; Pietri et al., 2003). These observations support the idea that there are two distinct episodes of delamination from cranial neural fold epithelium.

Our results disagree with the findings from a recent report concluding that there is no reduced expression of E-cadherin in the

non-neural ectoderm (Dady et al., 2012). On the contrary, we observed increased expression of E-cadherin in the non-neural ectoderm as compared with the neural ectoderm. Indeed, the E-cadherin *in situ* hybridization results corroborate the results from our immunostaining data and also show that E-cadherin protein is relatively stable. Based on this observation, we define the non-neural ectoderm as the region of ectoderm that expresses higher E-cadherin levels and has an absence of N-cadherin. The pattern of cell delamination in chicken embryos is comparable to that of mouse embryos, as *Snail2* is initially found in a region devoid of *Sox1* or N-cadherin expression, whereas at later stages *Snail2* expression is found in the neural ectoderm (compare Fig. 9A with 9B).

The ventral-to-dorsal formation of derivatives – early fate restriction

Numerous studies have shown that neural crest derivatives are formed in a ventral-to-dorsal sequence (Baker et al., 1997; Krispin et al., 2010; Lumsden et al., 1991; McKinney et al., 2013; Nitzan et al., 2013; Osumi-Yamashita et al., 1994; Shoval and Kalcheim, 2012). It is interesting to note that, in the study by Lumsden et al. (Lumsden et al., 1991), although no attempt was made to label distinct regions of the neural fold, the authors found that, when the neural fold in the midbrain was labeled at early stages, only labeling of the ectomesenchyme was observed in the majority of cases. This is also consistent with our current findings. By contrast, the study by Baker et al. (Baker et al., 1997) shows that early migrating cells can give rise to all of the different derivatives. The discrepancy could be because of the different methodology employed, as the grafts used in the study by Baker et al. would comprise significantly more cells than the number of cells that we labeled with *Dil*.

Recent studies disagree as to whether trunk neural crest cells are fate restricted prior to delamination (Krispin et al., 2010; McKinney et al., 2013; Nitzan et al., 2013). Krispin et al. and McKinney et al. looked at whether neural crest cells delaminate from the neural tube in an orderly fashion and whether fate restriction occurs prior to

delamination. The only consensus from these studies was that precursors of the sympathetic ganglia were the first cells to delaminate and they were fate restricted. However, Nitzan et al. showed that *Foxd3* expression in the dorsal neural tube is heterogeneous and that melanoblast precursors downregulate *Foxd3* before delamination. Perturbing the expression of *Foxd3* in the neural tube caused these cells to adopt neuronal or melanocytic cell fates. This is corroborated by the expression of melanoblast markers prior to delamination (Erickson and Goins, 1995; Wehrle-Haller and Weston, 1995; Wilson et al., 2004). In addition to melanoblasts, in the dorsal neural tube of mice there is a population of neurogenin 2-expressing trunk neural crest cells that preferentially give rise to cells within the dorsal root ganglia (Zirlinger et al., 2002). Similarly, in zebrafish, it has been shown that both the cranial and trunk neural crest cells are already fate restricted prior to delamination (Raible and Eisen, 1994; Schilling and Kimmel, 1994). This shows that it is highly likely that fate restriction occurs prior to delamination.

Previously, it was shown that downregulation of *Sox2* was required for the induction of neural crest cells (Remboutsika et al., 2011; Wakamatsu et al., 2004). *Sox1*, like *Sox2*, is expressed at lower levels in the dorsal region of the neural epithelia, indicating that these factors might be involved in the positioning of the neural plate border (Wakamatsu et al., 2004). The expression level of *Sox1* might be important for inducing cells to adopt a neuronal fate and for generating heterogeneity within the neural fold, since *Sox1* is differentially expressed by the two populations of delaminating cells. In mice, the removal of *Sox1* causes only microphthalmia (Nishiguchi et al., 1998); however, this could result from compensation by other members of *SoxB1* gene family.

Is ectomesenchyme a unique derivative of the cranial neural fold?

When transplanted into the trunk, the cranial neural fold readily forms ectopic cartilage and bone (Le Douarin and Teillet, 1974); however, the trunk neural crest does not seem to give rise to cartilage or bone when transplanted into cranial regions (Nakamura and Ayer-le Lievre, 1982). This has led to questioning as to whether ectomesenchyme is a unique derivative of the cranial neural fold. It has been suggested that both cranial and trunk neural crest cells have the same developmental potential and that formation of the ectomesenchyme is due to inductive environmental factors within the branchial arches (Abzhanov et al., 2003; Baker et al., 1997; Blentic et al., 2008; Lumsden, 1988; McGonnell and Graham, 2002). However, a study has shown that trunk and cranial ‘neural crest’ respond differently to the same signal, and that the formation of ectomesenchyme by trunk neural crest cells involves their conversion to a more cranial-like state (Abzhanov et al., 2003). This might indicate that trunk neural crest only forms ectomesenchyme via extensive cellular reprogramming. Although *Sox1-Cre* was used to show that, in the trunk, mesenchymal stem cells (MSCs) were derived from the neural epithelium, the significance of this is unclear, as MSCs were found only in embryos, and trunk neural crest does not normally give rise to mesenchymal lineages. In some vertebrates, such as the turtle, the trunk neural crest appears to give rise to ectomesenchymal derivatives in the plastron and fins (Clark et al., 2001; Raven, 1936; Smith and Hall, 1990). However, at least in zebrafish, fin rays and scales are not derived from the neural crest (Lee et al., 2013; Mongera et al., 2013). By contrast, the cranial neural fold, when transplanted into the trunk, readily forms ectopic cartilage and bone, even in a foreign environment (Le Douarin and Teillet, 1974). This implies that there could be intrinsic differences between the cranial and trunk ectodermal regions.

In 1950, Hörstadius stated that it was unclear whether the neural crest should be considered an outgrowth from the central nervous system or a separate rudiment (Hörstadius, 1950). Other than the ability of neural crest cells to undergo EMT, it has remained challenging to define these cells in terms of their origin or differentiation potential. Based on the presence of two separate epithelial domains in the cranial neural fold – one giving rise to ectomesenchyme and the other sharing an origin and cell fate with cells that delaminate in the trunk – it might be time to reconsider the nature of the neural crest and the origin of ectomesenchyme. The term ‘metablast’ was proposed by Weston et al. to explain the unique potential of the cranial neural fold to give rise to ectomesenchyme (Weston et al., 2004). This term was coined to take into account the unique population of cells that delaminates in the cranial region, which has a different origin and fate from the neural crest. Alternatively, we can adopt a broader definition of the neural crest: neural crest cells originate from the border of the neural and non-neural ectoderm and subpopulations of such broadly defined neural crest cells can be further refined.

Given the close proximity of the two epithelial domains within the cranial neural fold, we are faced with the dilemma of how to define the cells delaminating from the non-neural ectoderm. Further work is needed to adequately assess how these two regions segregate and to determine their relatedness. This might also serve to deepen our understanding of the emergence of ectomesenchyme in vertebrates and of how different derivatives are formed.

MATERIALS AND METHODS

Mouse strains

All experiments involving animals were reviewed and approved by the Institutional Animal Care and Use Committee of the Agency for Science, Technology and Research (A*STAR) of Biomedical Science Institutes. *Sox1-Cre* (Takashima et al., 2007) and *HtPA-Cre* (Pietri et al., 2003) were bred with reporter strains *Rosa26-YFP* (Srinivas et al., 2001) and *Rosa26-lacZ* (Soriano, 1999) for lineage analysis.

Chicken embryo manipulation

Chicken embryos (Chew's Agriculture, Singapore; Shiroyama Farm, Kanagawa, Japan) used for lineage analysis were cultured by a modification of the cornish pasty method (Nagai et al., 2011). Two embryos were joined together, forming a flat disc, but only one embryo was *Dil* or *DiO* labeled. The *Dil*-labeled embryo was immobilized by the addition of 0.1% Bacto Agar and cultured at the air/medium interface to increase the survival of the labeled embryos. After 2 days of culture, the embryos reached HH16–18, similar to *in ovo* development. *Dil* labeling was performed using a PLI-100 microinjector (Harvard Apparatus). *Dil* was used at 1 mg/ml in dimethylformamide for lineage tracing.

Chicken embryos were cultured dorsal side upward on agarose dishes using the paper culture method (Chapman et al., 2001). A small hole was made in the vitelline membrane and the non-neural ectoderm of the neural fold was labeled with *Dil*. Time-lapse video microscopy was carried out using an Olympus BX51W1 microscope fitted with an incubator box.

Histology and immunohistochemistry

Embryos were fixed in 4% paraformaldehyde, washed with PBS, equilibrated in 30% sucrose and embedded in Tissue-Tek OCT compound (Sakura Finetek). Sectioning was performed on a Leica CM1900 cryostat. Immunohistochemistry was performed as described (Breau et al., 2008). Images were acquired using a Zeiss LSM700 or an Olympus FV1000 confocal microscope.

Immunohistochemistry was performed using: goat anti-*Sox1* (1:100, AF3369, R&D Systems) and rat anti-N-cadherin (1:100, 205605, NCD2, Calbiochem); rabbit anti-*Sox9* (1:1000, AB5535, Millipore); rat anti-E-cadherin (1:100, U3254, Decma, Sigma); goat anti-L-CAM [1:400, chicken E-cadherin (Thiery et al., 1984)]; rabbit anti-Snai2 (1:100, #9585, Cell

Signaling Technology); mouse anti-RhoB (56.4H7), mouse anti-Islet1/2 (39.4D5), mouse anti-Pax3 (Pax3) and mouse anti-Tfap2a (3B5) (1:100, DSHB); chicken anti-GFP (1:500, ab13970, Abcam); mouse anti- β -dystroglycan (1:100, B-DG-CE, Novocastro Laboratories); mouse anti-HuC/D (A21271) and mouse anti-ZO-1 (339100) (1:100, Invitrogen); and rat anti-laminin α 1 (1:200, MAB1905, Chemicon). RhoB immunostaining was performed as described (Liu and Jessell, 1998). Quantification of images was carried out using Fiji (Schindelin et al., 2012) (supplementary material Fig. S1).

For *lacZ* staining of skull preparations, X-gal staining was performed as described (Pietri et al., 2004).

Fluorescent *in situ* hybridization (FISH)

E-cadherin FISH in chicken embryos was performed using the Fluorescein Tyramide Signal Amplification Kit (Perkin Elmer). Chicken E-cadherin probe (provided by J. L. Duband, University Pierre and Marie Curie-CNRS, Paris, France) was used at 75 ng/ml, with hybridization performed at 70°C. After FISH, the chicken embryos were sectioned and immunostained for Snail2.

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Competing interests

The authors declare no competing financial interests.

Author contributions

R.T.H.L., H.N. and Y.N. performed the experiments. R.T.H.L., G.S. and P.A.T. analyzed the data. R.T.H.L., J.A.W. and J.P.T. conceived the study and wrote the manuscript.

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Supplementary material

Supplementary material available online at <http://dev.biologists.org/lookup/suppl/doi:10.1242/dev.094680/-/DC1>

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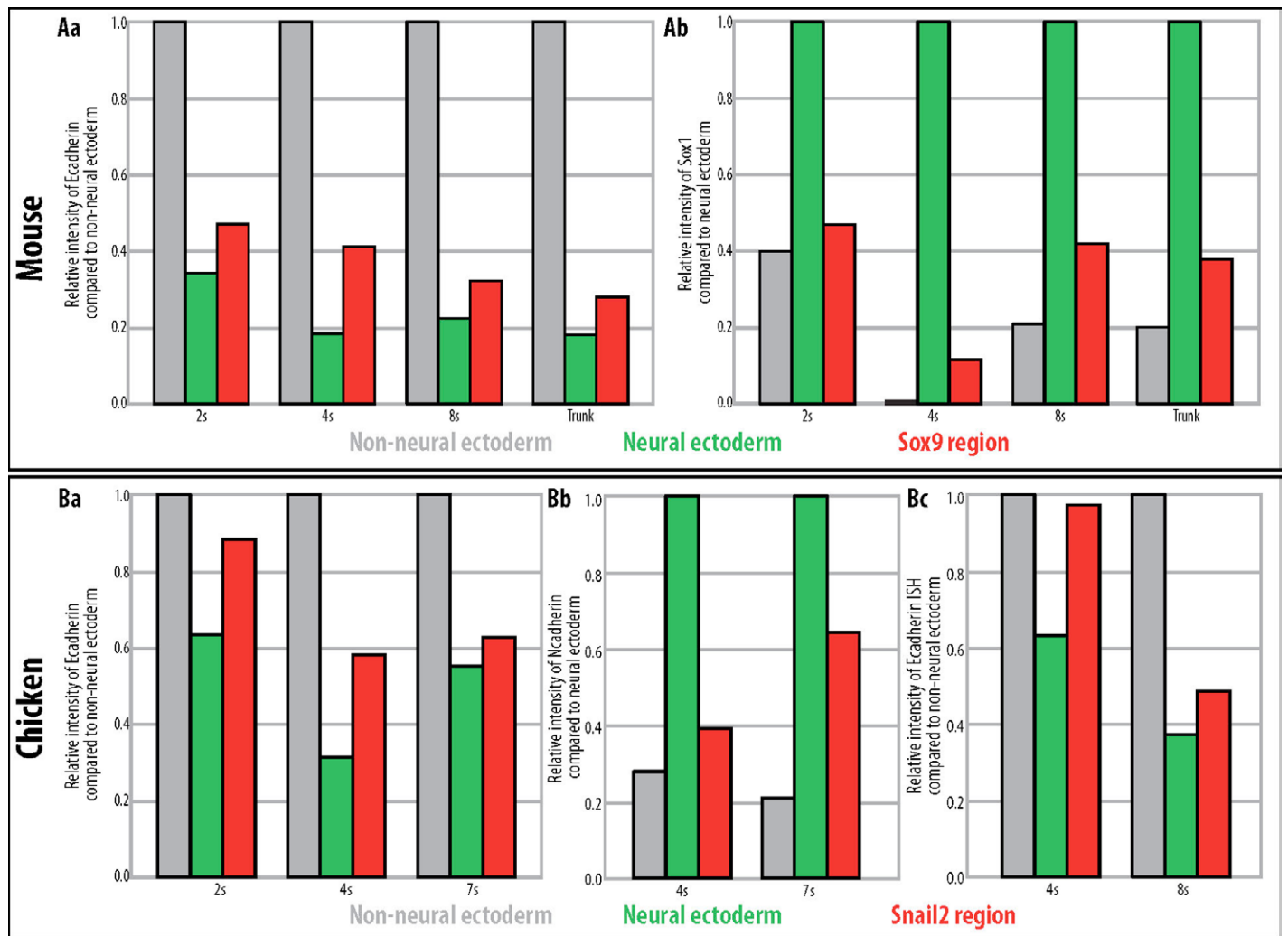


Fig. S1. Quantification of E-cadherin and Sox1/N-cadherin signals in the non-neural ectoderm, neural ectoderm, and in premigratory cells in embryos. The Sox9/Snail2 region is defined as cells remaining in the ectoderm that expresses the respective markers. We compared the expression level of E-cadherin and Sox1/N-cadherin in the region of interest by normalizing it with E-cadherin found in non-neural ectoderm and Sox1/N-cadherin found in neural ectoderm. This allowed us to determine whether the region of interest was more similar to non-neural or neural ectoderm. Quantification of the initial N-cadherin signal in chicken embryos was not performed as it would be misleading because it is expressed mainly apically. (A) Quantification of E-cadherin and Sox1 in different regions of the mouse embryo. At early stages (2 and 4 somites), the region expressing Sox9 shows higher and lower expression levels of E-cadherin (Aa) and Sox1 (Ab) as compared with those expressions in the neural ectoderm. By comparison, at 8 somites, the region expressing Sox9 shows lower levels of E-cadherin expression (Aa) and higher levels of Sox1 expression (Ab). This later population of Sox9-expressing cells in the cranial region is similar to Sox9-expressing cells in the trunk. In the cranial region, the expression level of Sox1 in the Sox9-expressing region is consistently lower as compared with the neural ectoderm at all stages, whereas the expression level of E-cadherin is initially higher but falls to levels similar to the neural ectoderm at later stages. Quantification of immunostaining for E-cadherin (Ba) and N-cadherin (Bb) in chicken embryos. Quantification of *in situ* hybridization of E-cadherin in chicken embryos (Bc). At early stages, between 2 and 4 somites, the Snail2-expressing region shows a similar pattern of E-cadherin and N-cadherin expression as that seen in non-neural ectoderm (high expression of E-cadherin and low expression of N-cadherin) as compared with that observed in the neural ectoderm. The pattern of E-cadherin mRNA expression in Snail2-expressing cells at 4 somites is also similar to the non-neural ectoderm (Bc). However, at 7 somites, the Snail2-expressing region shows a more similar pattern of E-cadherin and N-cadherin expression to that of the neural ectoderm (low expression of E-cadherin (Ba) and higher expression of N-cadherin (Bb) as compared with that of the non-neural ectoderm. At 8 somites, the pattern of E-cadherin mRNA expression in Snail2-expressing cells is no longer similar to the non-neural ectoderm (Bc). There is a clear difference in the mRNA expression of E-cadherin between the non-neural and neural ectoderm (Bc).

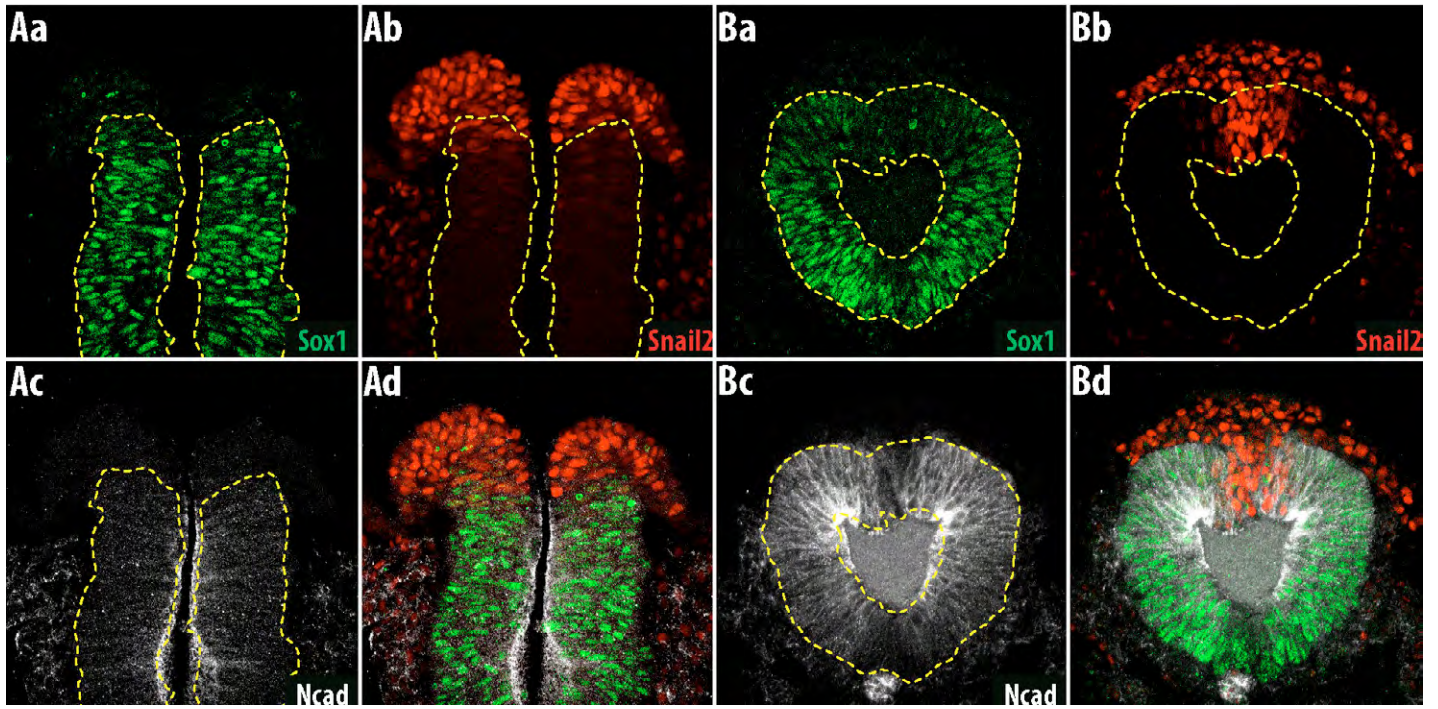


Fig. S2. Sox1 and N-cadherin are similarly expressed in the neural epithelia. Sox1 and N-cadherin expression in the neural epithelia. (A,B) Sox1-expressing cells in the neural epithelia also co-express N-cadherin (yellow dotted line represent Sox1 expressing region). (A) Chicken embryo at 4 somites. At this stage the majority of Snail2 expressing cells do not express Ncad or Sox1. At these early stages, we observed some Snail2 expressing cells in the neural ectoderm. (B) Chicken embryo at 7 somites. Premigratory Snail2-expressing cells are found in the neural ectoderm.

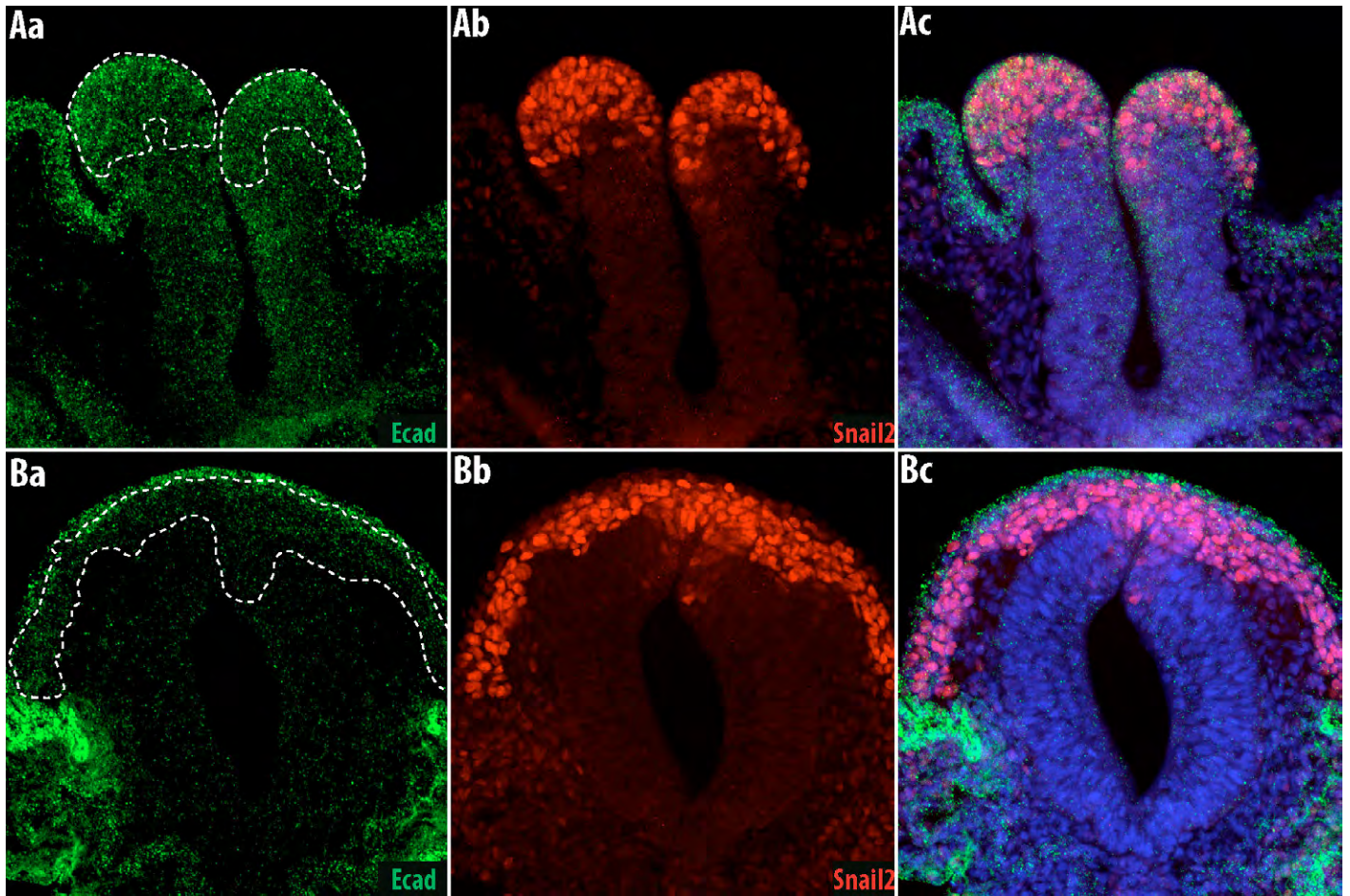


Fig. S3. E-cadherin in-situ shows a similar pattern to E-cadherin immunostaining. Fluorescent in situ hybridization of E-cadherin and Snail2 immunostaining in chicken embryos (A,B). (A) At 4 somites, Snail2 expressing cells still expressed relatively high levels of E-cadherin as compared with those in the neural ectoderm (white dotted area represent Snail2 expressing region). (B) At 8 somites this changes, with only the overlying non-neural ectoderm expressing E-cadherin (white dotted area represents the Snail2 expressing region). E-cadherin expression in the neural ectoderm and Snail2-expressing cells is different from what we observed for E-cadherin immunostaining (Fig. 4B).

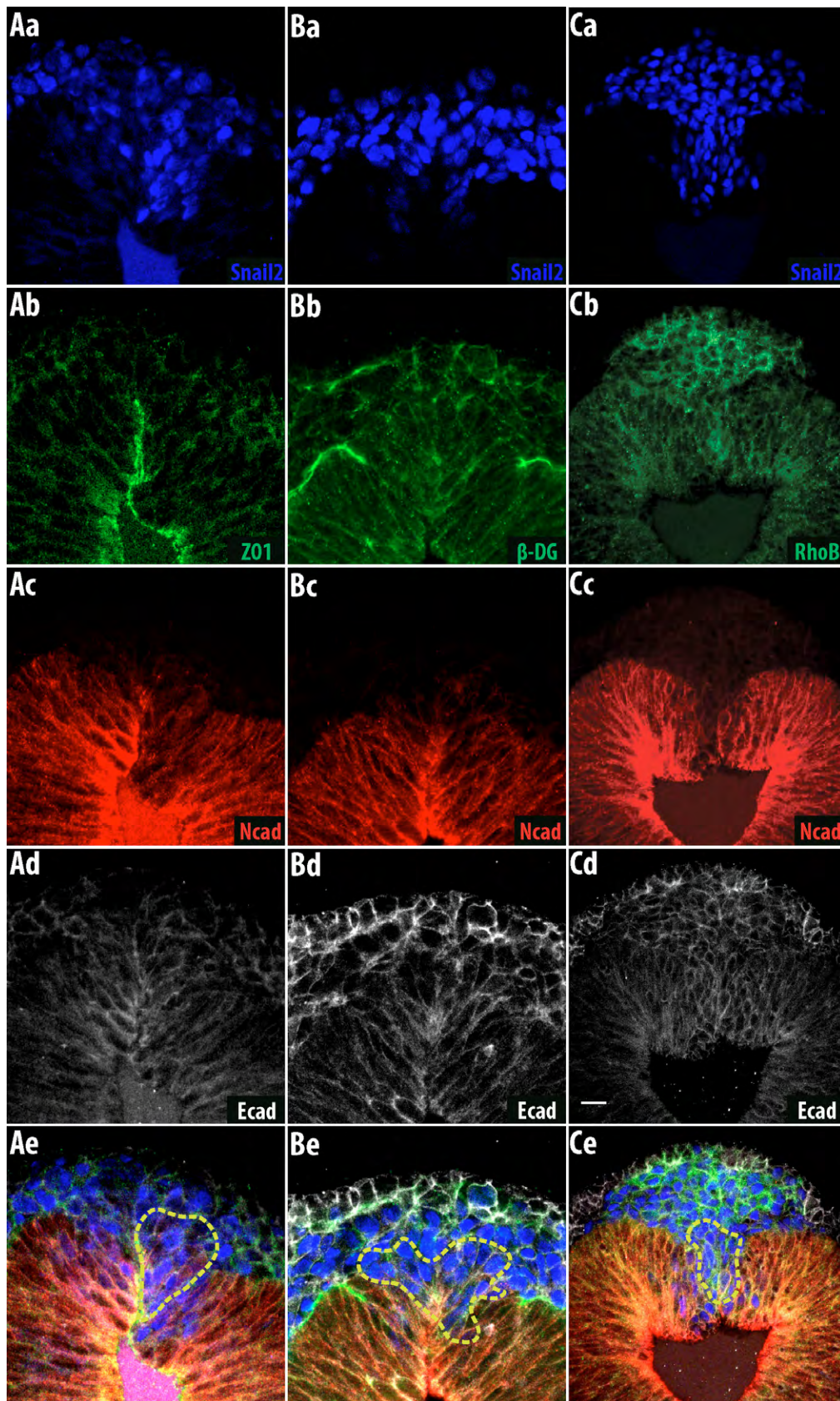


Fig. S4. In chicken embryos cells delaminate from the neural ectoderm at 7 somites. Snail2-expressing cells in the neural epithelia delaminate at about 7 somites in chicken embryos. (A-C) Chicken embryos (7 somites). Snail2-expressing cells in the neural epithelia are undergoing EMT, as they have lost apical and basal polarity as shown by the cortical localization of ZO1 (A) and β -Dystroglycan (B) (yellow dotted line). RhoB is expressed in Snail2-positive cells found in the neural ectoderm (yellow dotted line), further indicating that these cells are undergoing EMT (C).

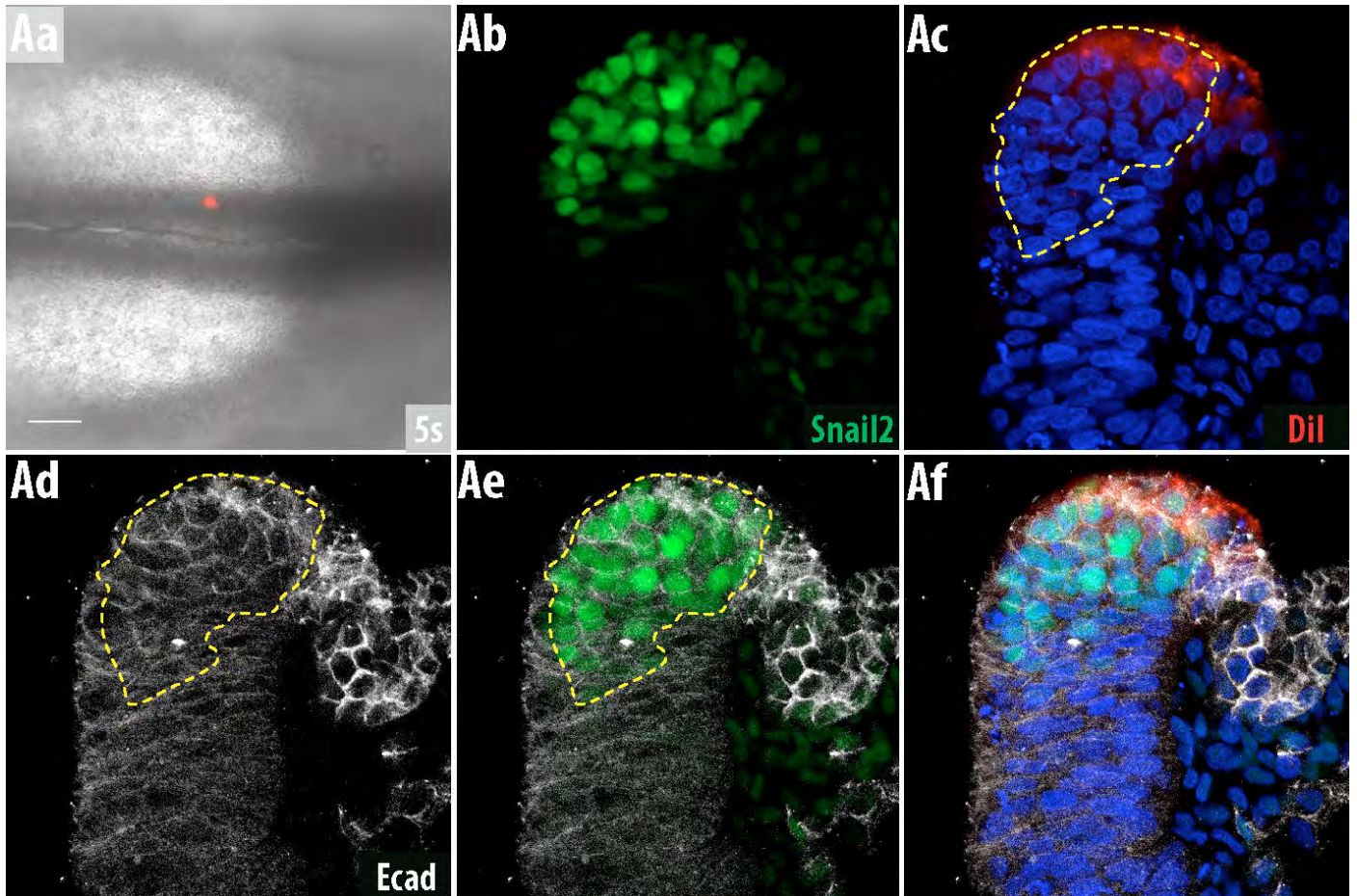


Fig. S5. DiI labeling of the non-neural ectoderm. A chicken embryo used for lineage analysis (between 4 and 5 somites and 6 embryos were analyzed) with the non-neural ectoderm labeled with DiI. All DiI-labeled cells were found in the non-neural ectoderm expressing Snail2 and E-cadherin (Ad,Ae), yellow dotted lines represent the region where Snail2 is expressed.

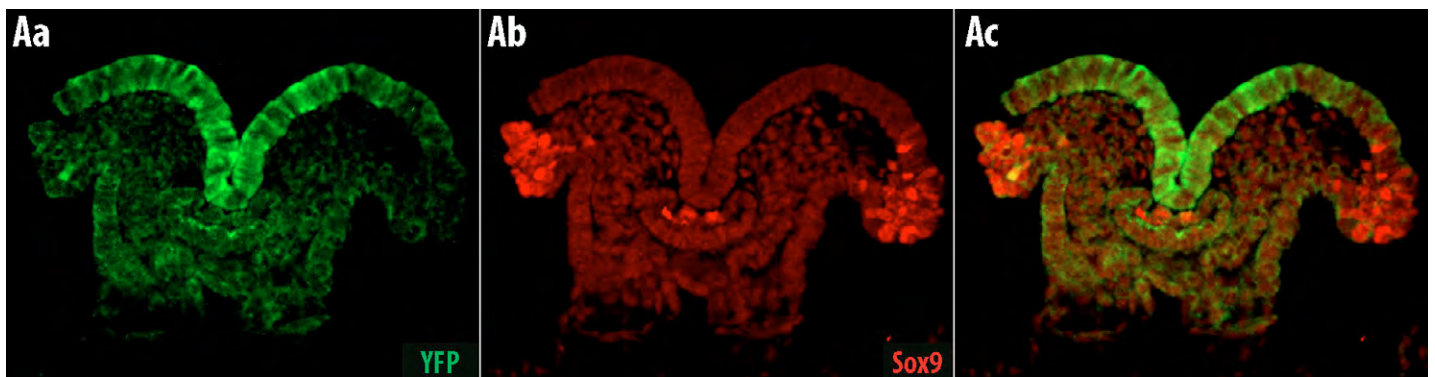


Fig. S6. YFP reporter is expressed in the neural ectoderm prior to the delamination of cells in the neural fold. Mouse embryo (3 somites). Activation of YFP reporter by Sox1 Cre occurs prior to the delamination of cells. YFP (Aa) is expressed in the neural ectoderm prior to the delamination of cells (Ab).

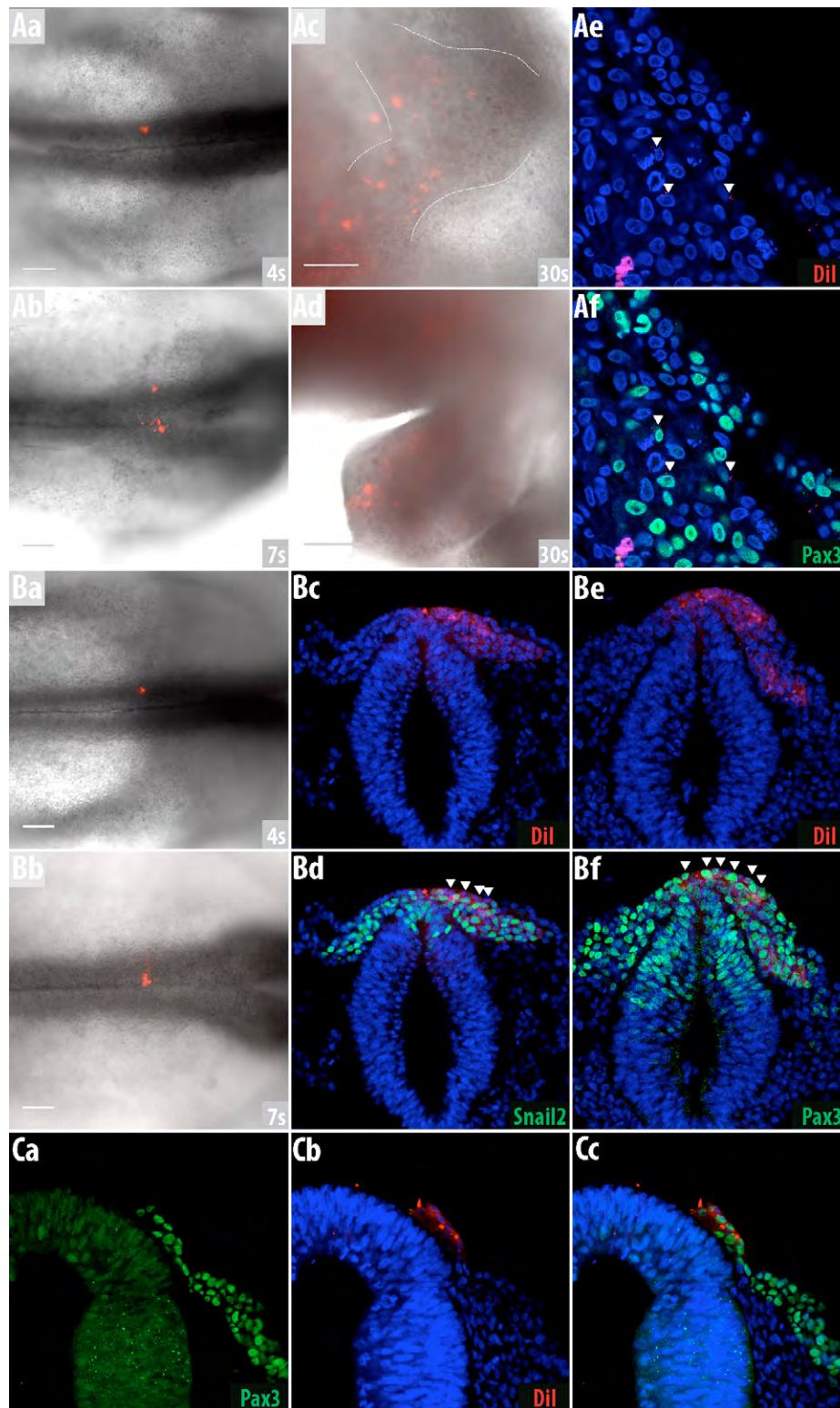


Fig. S7. The trigeminal placode and its derivatives are also labeled in some non-neural DiI-labeled embryos. DiI labeling of the lateral non-neural ectoderm also labels the trigeminal placode. We analyzed 2 embryos at ~30 somites where there was labeled cells in the trigeminal ganglia (A), and analyzed 3 embryos which show DiI labeling on the dorsal midline at 7 somites (B,C). DiI was used to label the non-neural ectoderm at 4 somites (A) and some DiI-labeled cells have migrated close to the midline of the neural tube at 7 somites (Ab). After growing to ~30 somites, DiI-labeled cells can be clearly seen in the trigeminal ganglia (Ac) as well as in the branchial arch (Ad). DiI-labeled cells in the trigeminal ganglia are Pax3-positive, indicating that they are derived from the trigeminal placode (Ae,Af). DiI-labeled cells can be seen in the trigeminal placode when the non-neural ectoderm is labeled at 4 somites (B,C). Chicken embryo at 4 somites (Ba) and 7 somites (Bb-f). DiI can be seen in cells that are Snail2-negative (arrowheads in Bd). These cells on the surface ectoderm are Pax3-positive (arrowheads in Bf). (C) DiI-labeled chicken embryo (13 somites), showing that the Pax3-expressing trigeminal placode (Ca) is labeled with DiI (Cb).



Movie 1. Time-lapse movie of delaminating cells in the mesencephalon labeled with DiI. Chicken embryos were labeled with DiI at 4somites and imaged every 3 minutes. Cells move medially before undergoing EMT and migrating away. Anterior is to the right. The dotted line represents the midline and the line represents regions of the neural tube that have not fused.