

# Induction of tooth and eye by transplantation of activin A-treated, undifferentiated presumptive ectodermal *Xenopus* cells into the abdomen

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**ABSTRACT** Activin A can induce the *Xenopus* presumptive ectoderm (animal cap) to form different types of mesoderm and endoderm at different concentrations and the animal cap treated with activin can function as an organizer during early development. The dissociated *Xenopus* animal cap cells treated with activin form an aggregate and it develops into various tissues *in vitro*. In this study, to induce jaw cartilage from undifferentiated cells effectively, we developed a culture method to manipulate body patterning *in vitro*, using activin A and dissociated animal cap cells. An aggregate consisting only of activin A-treated dissociated cells developed into endodermal tissues. However, when activin A-treated cells were mixed with untreated cells at a ratio of 1:5, the aggregate developed cartilage with the maxillofacial regional marker genes, *goosecoid*, *Xenopus Distal-less 4* and *X-Hoxa2*. When this aggregate was transplanted into the abdominal region of host embryos, maxillofacial structures containing cartilage and eye developed. We raised these embryos to adulthood and found that tooth germ had developed in the transplanted tissue. Here, we show the induction of jaw cartilage, tooth germ and eye structures from animal caps using activin A in the aggregation culture method. This differentiation system will help to promote a better understanding of the regulating mechanisms of body patterning and tooth induction in vertebrates.

**KEY WORDS:** jaw, tooth, activin, amelogenin, animal cap

## Introduction

Activin A, a member of the TGF- $\beta$  family, has been shown to be a crucial morphogen in the vertebrate body plan, as demonstrated by its potent mesoderm-inducing activity on *Xenopus* undifferentiated presumptive ectoderm cells (animal cap) (Asashima *et al.*, 2000, Green 2002). Activin A induces mesoderm and endoderm from *Xenopus* undifferentiated presumptive ectoderm in a dose-dependent manner, whereas the untreated ectoderm forms an irregular-shaped epidermis (Asashima *et al.*, 1990, Green and Smith 1990, Jones *et al.*, 1993, Smith *et al.*, 1993). At low concentrations of activin A, ventral mesoderm, such as blood-like cells, coelomic epithelium and mesenchyme are induced in the explants (Miyanao *et al.*, 1998). At intermediate concentrations, muscle and neural tissue are induced (Tamai *et al.*, 1999). At high

concentrations, notochord, the most dorsal mesoderm is induced (Ninomiya *et al.*, 1999). These tissues induced by activin A are indistinguishable at the histological level from those found in normal embryos (Okabayashi and Asashima 2003). Furthermore, activin A can elicit gene cascades in the animal caps with a time course that mimics the sequence in normal embryonal development. When the dissociated animal cap cells are exposed to activin A, the aggregate of the dissociated cells distinctly expresses a range of genes in a dose-dependent manner (Green *et al.*, 1992). An activin A-loaded bead can cause waves of gene expression to spread out through a static population of undissociated blastula

*Abbreviations used in this paper:* AA, activin-A; FDA, fluorescein-dextran-amine; ODC, ornithine decarboxylase; TGF, transforming growth factor; TRDA, texas-red-dextran-amine.

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cells (Gurdon *et al.*, 1994). The animal cap cells can make a low to high gene response to activin concentration.

The most characteristic property of activin A is the induction of organizer activity. When an activin A-treated animal cap is transplanted into the ventral side of early gastrulae, a well-organized, secondary embryo is induced (Ninomiya *et al.*, 1998). By combining these systems, we have demonstrated a sandwiched culture method that reproduces the fundamental embryonic body *in vitro* (Ariizumi and Asashima 1994). Animal caps treated with activin A, were incubated (preculture) and then sandwiched between untreated two animal caps. By this sandwiched culture method, a head or trunk-tail structure was induced depending on activin A concentration and the preculture period following the activin A-treatment. It is therefore theoretically possible to reproduce embryonic induction using activin A and the sandwiched culture method and to generate a developing embryo from undifferentiated presumptive ectoderm *in vitro*.

During craniofacial development, neural crest-derived mesenchymal cells migrate to the branchial arches and contribute extensively to formation of maxillofacial structures. Tooth formation requires a serial interaction between epithelium and neural crest-derived mesenchyme in jaw (Chai *et al.*, 2000, Wilde 1955, Zeichner-David *et al.*, 1995). We hypothesized that tooth reproduction requires jaw induction. Studies in *Xenopus* development have given rise to an instructive model that cell fate can be manipulated by morphogens or inducing factors and *Xenopus* has been used

extensively to probe the events in early development of vertebrates (De Robertis *et al.*, 2000, Gurdon *et al.*, 2003, Melton 1987). Recently, we succeeded in inducing jaw cartilage from *Xenopus* undifferentiated presumptive ectoderm using sandwiched culture method and activin A (Furue *et al.*, 2002). The animal cap treated with the high concentration of activin A, was sandwiched with untreated animal caps developed jaw cartilage with the expression of the maxillofacial regional marker genes. We hypothesized that we could induce tooth germ from animal caps using activin A. Green *et al.*, demonstrated that the activin A-treated dissociated animal cap cells interact with untreated animal caps cells, resulting in induction of neural tissues (Green *et al.*, 1997). We demonstrated that, when cells treated with activin A were mixed with untreated cells, the activin A-treated (1 ng/ml) cells concentrated in a central mass of the aggregates and they effectively formed notochord (Kuroda *et al.*, 2002, Kuroda *et al.*, 1999). Therefore, we further hypothesized that it would be possible to reproduce jaw effectively *in vitro* using activin A and dissociated animal cap cells and jaw induction would lead to tooth formation. In this study, we have developed an aggregation culture method to manipulate the body patterning and we could induce the maxillofacial region effectively in the aggregate of the animal cap cells. When these aggregates were transplanted into the abdominal region of host embryos, maxillofacial structures containing cartilage and eye developed in the transplanted tissues. Furthermore, we raised these embryos to adulthood and found that tooth germ had developed in transplanted tissue. This aggregation culture method will promote understanding of the reproduction of body patterning and tooth induction in vertebrates.

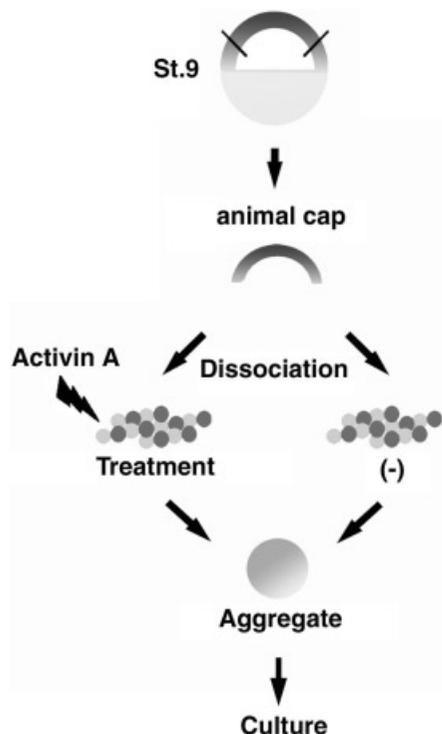
## Results

### *The effect of the mixture ratio of activin A-treated and untreated animal cap cells on tissue differentiation in aggregates*

We dissected 12 animal caps, dissociated the animal cap cells, treated these cells with activin A and made an aggregate (Fig. 1). After 7 days in culture, we performed histological analysis with PAS/Alcian blue staining. When all of the dissociated cells of 12 animal caps were treated with 25 ng/ml of activin A for 1 h and then formed an aggregate (AA-aggregate), the aggregates developed into endodermal tissues (Fig. 2A). When dissociated cells of 6 animal caps treated with 25 ng/ml of activin A for 1 h were mixed with untreated dissociated cells of 6 animal caps (1:1 aggregate), the aggregates developed into mesodermal tissues, such as notochord and muscle (Fig. 2B). Finally, cartilage tissue was observed in the aggregate (Fig. 2C) when dissociated cells of 2 animal caps were treated with 25 ng/ml of activin A for 1 h and then mixed with untreated cells of 10 animal caps (1:5 aggregate). The frequency of the aggregate that formed cartilage tissue was 67% (20/30) of the 1:5 aggregates, 11% (5/42) of the 1:1 aggregates and none (0/20) of the AA-aggregates. These results indicated that the ratio of mixture with activin A-treated and untreated animal cap cells reflected the induced-tissues in the aggregates and suggested that head-region was induced in 1:5 aggregate.

### *Gene expression in the aggregates*

In order to confirm the induced-region in the aggregates, we examined gene expression of *Xenopus Hoxa2* (*X-Hoxa2*), *Xnot*,

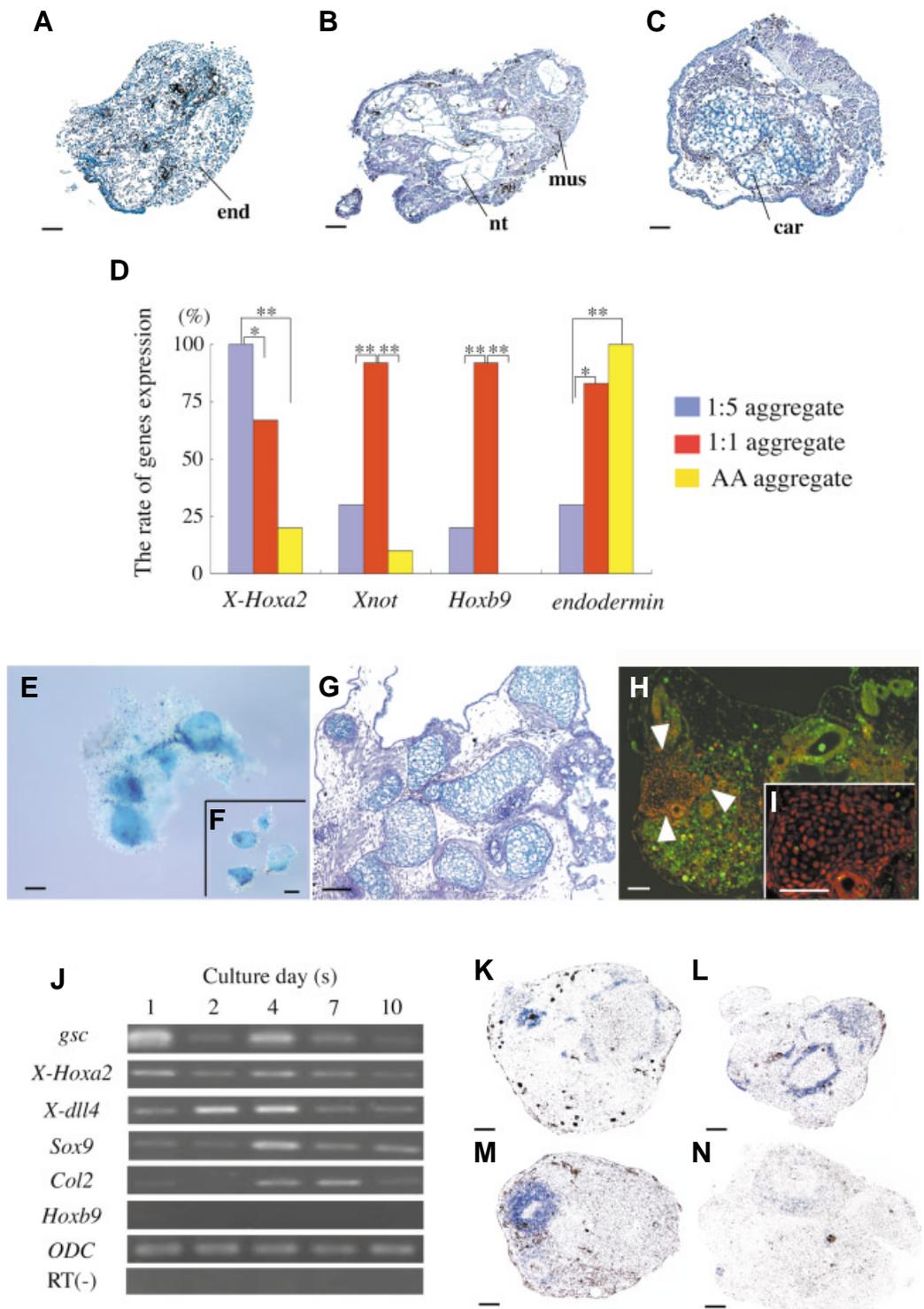


**Fig. 1. Schematic diagram of the procedures involved in the aggregation culture method.** Animal caps were dissected from *Xenopus* embryos (stage 9) and dissociated in calcium- and magnesium-free medium. The dissociated cells were incubated with or without 25 ng/ml activin A for 1 h. After washing away free activin A five times, the activin A-treated cells were mixed with untreated cells and aggregated in calcium- and magnesium-containing medium.

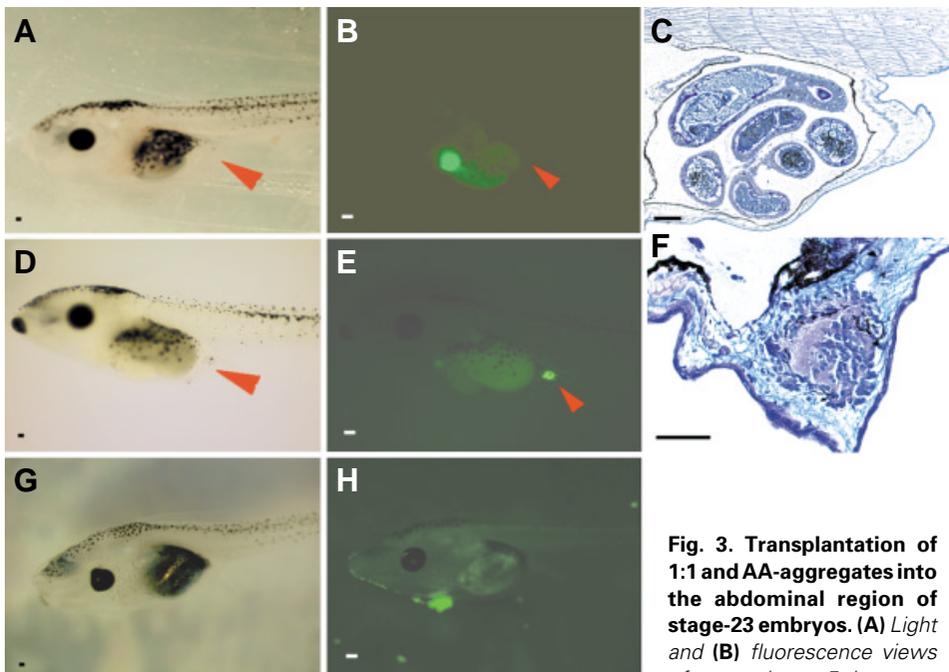
*Hoxb9*, *endodermin* 1d-cultured explants using RT-PCR (Fig. 2D). *X-Hoxa2* is expressed in the 2nd pharyngeal arch and associated with the evolution of jaws (Pasqualetti *et al.*, 2000). The expression of *Xnot*, a homeobox gene is restricted to the organizer region and presumptive notochord (von Dassow *et al.*, 1993). *Hoxb9*, a posterior neural marker, is expressed in the posterior part of the spinal cord (Wright *et al.*, 1990). *endodermin*, a pan-endodermal marker, is localized almost exclusively to the endoderm (Sasai *et al.*, 1996). In replicate experiments, AA-aggregates expressed *endodermin* at the frequency of 100% (10/10), but *X-Hoxa2* at 20% (2/10), *Xnot* at 10% (1/10), *Hoxb9* at 0% (0/10). The 1:1 aggregates expressed *endodermin* at 83% (10/12), *Hoxb9* at 92% (11/12), *X-Hoxa2* at 58% (7/12) and *Xnot* at 92% (11/12). The 1:5 aggregates expressed *X-Hoxa2* at 100% (10/10), but *endodermin* at 30%, (3/10), *Hoxb9* at 20% (2/10) and *Xnot* at 30% (3/10). These results revealed that endodermal tissues were induced in the AA-aggregates, trunk-tail region was induced in the 1:1 aggregates and maxillofacial region was induced in the 1:5 aggregates.

**Cell lineage of activin A-treated animal cap cells in the aggregates**

We further cultured 1:5 aggregates for 7 days (Fig. 2 E,F), or 14 days (Fig. 2G) and observed the induction of a large amount of cartilage tissue which was detected by Alcian blue staining and morphology. To follow cell lineage of activin A-treated cells in the aggregation induced cartilage, we analyzed 1:5 aggregates made of the activin A-treated cells of animal caps derived from embryos labelled with texas-red-dextran-amine (TRDA) and untreated cells of animal caps derived from embryo labeled with fluorescein-dextran-amine (FDA). The Alcian blue-stained cartilage was composed of TRDA-labeled-activin A-treated



**Fig. 2. The phenotypes of AA-aggregate, 1:1 aggregate and 1:5 aggregate.** PAS/Alcian blue stained-sections of (A) AA-aggregate, (B) 1:1 aggregate and (C) 1:5 aggregates on culture day 7. (D) RT-PCR analysis of the genes expressed in aggregates. (\* $p < 0.05$ , \*\* $p < 0.01$ , AA-aggregate, 1:5 aggregate,  $n = 10$ ; 1:1 aggregate,  $n = 12$ ). (E) Whole mount Alcian blue staining of the 1:5 aggregate on culture day 7. (F) Extracted cartilage tissue from (E). (G) PAS/Alcian blue stained-section of the 1:5 aggregate on culture day 14. (H) Cell-lineage analysis of the 1:5 aggregate on culture day 7. White arrowheads point to cartilage tissue. (I) Higher magnification of (H). (J) RT-PCR analysis of the 1:5 aggregates following 10-day-culture. (K) *X-dll4*, (L) *col2*, (M) *gsc* and (N) *Sox9* expression, in the 1:5 aggregates. end, endodermal tissue; car, cartilage; mus, muscle; nt, notochord. Bars, 100  $\mu\text{m}$ .



**Fig. 3. Transplantation of 1:1 and AA-aggregates into the abdominal region of stage-23 embryos.** (A) Light and (B) fluorescence views of an embryo, 5-day-transplanted with AA aggregate. (C) A section of 10-day-transplanted tissues. (D) Light and (E) fluorescence views of an embryo, 5-day-transplanted with 1:1 aggregate. (F) A section of 10-day-transplanted tissues. (G) Light and (H) fluorescence views of a 7-day normal embryo. The fluorescence in the anterior region is the fed baits in the abdomen. Bars, 100  $\mu$ m.

planted with AA aggregate. (C) A section of 10-day-transplanted tissues. (D) Light and (E) fluorescence views of an embryo, 5-day-transplanted with 1:1 aggregate. (F) A section of 10-day-transplanted tissues. (G) Light and (H) fluorescence views of a 7-day normal embryo. The fluorescence in the anterior region is the fed baits in the abdomen. Bars, 100  $\mu$ m.

cells, but not FDA-labeled-untreated cells (Fig. 2 H,I). These results indicate that cartilage was derived from activin A-treated cells.

#### Maxillofacial regional marker gene expression in 1:5 aggregates

To confirm whether the maxillofacial region was really induced in the 1:5 aggregates, we further examined the gene expression of *gooseoid* (*gsc*), *Xenopus Distal-less 4* (*X-dll4*), *X-Hoxa2*, *Sox9* and Collagen type 2 (*Col2*). *gsc* is a homeobox-containing gene that is expressed in the gastrula dorsal lip (Cho *et al.*, 1991, McKendry *et al.*, 1998), in the ventral head region (Newman *et al.*, 1997) and then in the lower jaw cartilages (Furue *et al.*, 2002). *X-dll4*, a homeobox gene, is expressed in anterior ectodermal derivatives (ventral forebrain, cranial neural crest, cement gland) (Papalopulu and Kintner 1993), then in ventral head region including maxillofacial cartilages (Furue *et al.*, 2002). *Sox9* is expressed in cranial neural crest cells and pharyngeal arches (Spokony *et al.*, 2002). *Col2* is expressed in immature chondrocytes (Bieker and Yazdani-Buicky 1992, Seufert *et al.*, 1994) which was localized in craniofacial skeleton primordium and anterior notochord (Furue *et al.*, 2002). Analysis using RT-PCR showed that all these gene expression were detected in the 1:5 aggregates (Fig. 2J). The sections of the 1:5 aggregates were hybridized with DIG-labeled RNA probes of *X-dll4*, *col2*, *gsc* and *Sox9*. The expression of all these genes were localized in the 1:5 aggregates, but were not uniformly expressed (Fig. 2 K-N). These results indicated that 1:5 aggregates gave rise to maxillofacial derivatives.

#### Implantation of the aggregates into the abdominal region

When we transplanted AA-aggregates into the abdominal region, the transplanted tissues were assimilated with the host intestine (Fig.

3 A-C). When we transplanted 1:1 aggregates into the abdominal region, the transplanted tissues did not develop into cartilage (Fig. 3D-F). We transplanted 60 cases of 1:5 aggregates into the abdominal region. 44 cases of them were developed in the host tissues and GFP fluorescence was restricted under the abdomen region (Fig. 4A-C). PAS/Alucian blue staining of sections of the transplanted tissues showed the transplanted tissue developed into muscle, cartilage, neural and eye tissues, which were arranged into a maxillofacial-like-structure (Fig. 4 D,E) at 59% (26/44) although the aggregates *in vitro* developed into cartilage and mesenchymal tissues. Analysis using *in situ* hybridization revealed that the expressions of *Col2* (Fig. 4F), *gsc* (Fig. 4G), *X-dll4* (Fig. 4H) and *Sox9* (Fig. 4I) were detected in the section of transplanted tissues. The expressions of *gsc*, *X-Hoxa2*, *X-dll4*, *Sox9* and *Col2*, were detected, but *Hoxb9* expression was not detected in the dissected transplanted tissues by RT-PCR (Fig. 4L). Analysis using *in situ* hybridization showed that *Pax6*, a universal master control gene for eye morphogenesis (Onuma *et al.*, 2002) was expressed in the

transplanted tissue (Fig. 4J) like normal eye (Fig. 4K). RT-PCR analysis showed that *Pax6* was expressed in aggregates *in vitro* and in 1-day- and 3-day-transplanted into the abdomen, but it was not detected in normal abdomen (Fig. 4M). These results indicated that the transplanted 1:5 aggregates could develop into maxillofacial structures including cartilage and eye with gene expression in abdominal region.

#### Tooth induction from the aggregates in transplanted abdomen

*Xenopus laevis* develops teeth in upper jaw in adulthood. We hypothesized that tooth germ could be developed in aggregates transplanted into the abdominal region. We transplanted 1:5 aggregates into the abdominal region of embryos and raise six of the host embryos until stage 66 when tooth germ differentiates in the upper jaw in normal *Xenopus*. The protruding transplanted tissue was seen in the proximal area between both legs (Fig. 5 A-C). The section of the transplanted tissues showed that cartilage differentiated into membranous bone and surrounding the thin membranous bone, tooth germ-like structures were observed in three of six hosts (Fig. 5 D-F). We immunostained the tooth germ-like tissue with an anti-amelogenin antibody. We detected amelogenin protein in tooth germ-like structures in the transplanted tissue (Fig. 5G), which resembled normal tooth germ (Fig. 5H). These results indicated that tooth germ was developed from 1:5 aggregate.

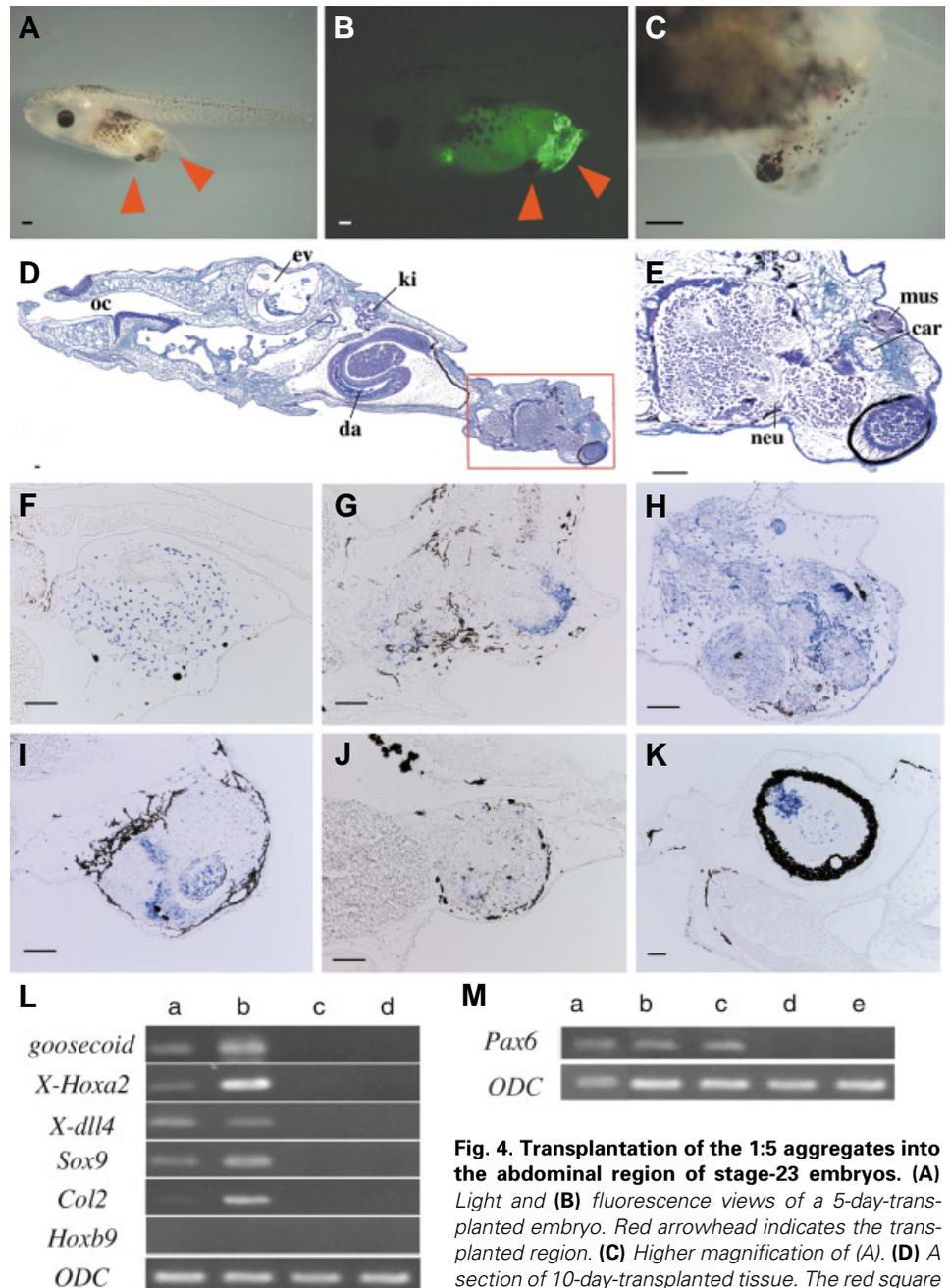
#### Discussion

The experiment described here showed that the mixture ratio of activin A-treated and untreated animal cap cells reflects the body patterning when we reproduce the tissues using animal caps cells. As previously described (Ninomiya *et al.*, 1999), endodermal

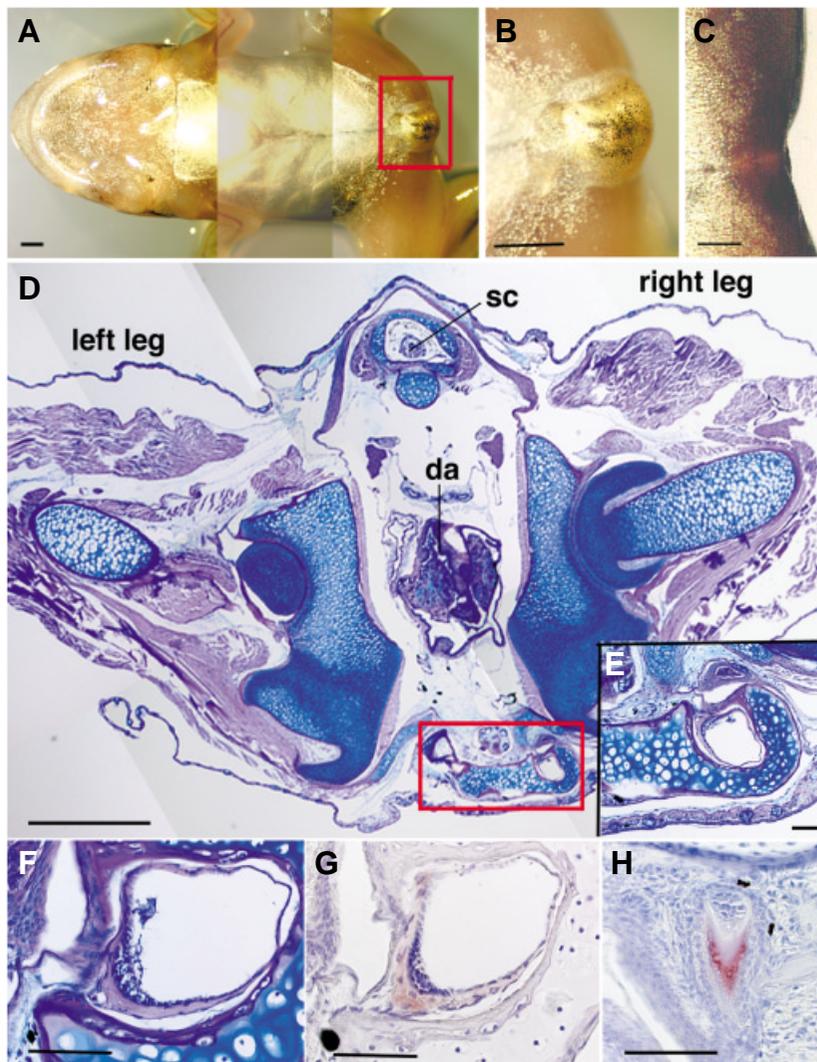
tissues with endoderm gene expression were induced in the AA-aggregates composing of all of the cells treated with high concentration of activin A (25 ng/ml). In the 1:1 aggregate of mixture of activin-treated and untreated animal cap cells at the ratio of 1:1, the trunk-tail region consisting of mesodermal tissues, such as notochord and muscle were induced with gene expression of *endodermin*, *Hoxb9* and *Xnot*. In the 1:5 aggregate of mixture of activin A-treated and untreated animal cap cells at the ratio of 1:5, the maxillofacial region including cartilage tissue was induced with gene expression of *X-Hoxa2*. We previously showed the induction of jaw cartilages with gene expressions of *gsc*, *X-dll4* and *Col2* *in vitro* from the *Xenopus* animal cap sheets using sandwiched culture method and activin A (Furue *et al.*, 2002). The 1:5 aggregates also expressed *gsc*, *X-dll4*, *Sox9* and *Col2*, suggesting that the maxillofacial region was effectively induced in the 1:5 aggregate. Gurdon demonstrated that activin A-loaded bead causes waves of gene expression to spread out through a static population of animal cap cells (Gurdon *et al.*, 1999). Furthermore, Green *et al.*, demonstrated that the activin A-induced mesoderm affects on non-induced animal caps cells resulting in inducing neural tissues and the dissociated cells treated with high concentration of activin A function as anterior positional value (Green *et al.*, 1997). These findings proposed that the mixtures at different ratio of activin A-treated and untreated dissociated animal cap cells develop into different positional values depending on their ratio even though the aggregates contained the cells treated with the same range of activin A. Analysis of cell lineage in 1:5 aggregate revealed that the induced cartilage was derived from activin A-treated cells, but not untreated cells. When we previously tried inducing jaw cartilage *in vitro* from *Xenopus* animal cap sheet (Furue *et al.*, 2002), cartilage was poorly induced from an animal cap sheet treated with activin A, but it was induced from an activin A-treated animal cap sheet sandwiched between two animal cap sheets. These findings suggested that induction of the maxillofacial region required the interaction between activin A-treated and untreated animal cap cells. It is assumed that mixing of cells with anterior positional values and untreated cells induces the wide range of genes and

further that the untreated cells might be affected by these gene products resulting in the induction of the maxillofacial structures.

To investigate whether the animal cap cell aggregates could develop into the maxillofacial structures *in vivo*, we made aggregates from eggs derived from GFP-transgenic *Xenopus* and transplanted the aggregates into the abdominal region of embryos. The



**Fig. 4. Transplantation of the 1:5 aggregates into the abdominal region of stage-23 embryos. (A)** Light and **(B)** fluorescence views of a 5-day-transplanted embryo. Red arrowhead indicates the transplanted region. **(C)** Higher magnification of **(A)**. **(D)** A section of 10-day-transplanted tissue. The red square indicates transplanted tissue. **(E)** Higher magnification of **(D)**. **(F)** *Col2*, **(G)** *gsc*, **(H)** *X-dll4*, **(I)** *Sox9*, **(J)** *Pax6* expression in the transplanted tissue. **(K)** *Pax6* expression in the normal eye of a stage-41 embryo. **(L)** RT-PCR analysis of **(a)** 1-day and **(b)** 3-day-transplanted tissue in the abdominal region and **(c)** 1-day- and **(d)** 3-day- normal abdominal tissue. **(M)** RT-PCR analysis of *Pax6* expression in **(a)** the aggregates on culture day 1; **(b)** 1-day- and **(c)** 3-day-transplanted tissue in the abdominal region and **(d)** 1-day- and **(e)** 3-day- normal abdominal tissue. Abbreviations: car, cartilage; ev, ear vesicle; da, digestive apparatus; ki, kidney; mus, muscle; neu, neural tissue; nt, notochord; oc, oral cavity. Bars, 100  $\mu$ m.



**Fig. 5. Transplanted tissue in the abdominal region of grown host.** (A) Ventral view of the grown host (stage 66). (B) Higher magnification of (A). (C) Normal view. (D) A section of the transplanted tissue. (E) Higher magnification and (F) tooth germ-like tissue of (D). (G) Immunolocalization of amelogenin in tooth germ-like structure in the serial section of (F) and (H) normal tooth germ in a stage-66 frog. Red squares outline the transplanted regions. Abbreviations: da, digestive apparatus; sc, spinal cord. Bars, 1 mm (A-D); 100  $\mu$ m (E-H).

maxillofacial structure consisting of muscle, cartilage, neural and eye tissues was developed in the 1:5 aggregates transplanted into the abdominal region. It is well-known that *Pax6* is a universal master control gene for eye morphogenesis (Onuma *et al.*, 2002) because the ectopic expression of mouse *Pax6* in *Drosophila* induced compound eye (Gehring 2002, Halder *et al.*, 1995). Actually, *Pax6* was expressed in the aggregates transplanted into abdomen. *Pax6* was expressed also in aggregates *in vitro* although eye induction was never found in the aggregate *in vitro*. The abdomen is a good region to culture the aggregates. These findings demonstrated that jaw cartilage induced *in vitro* could be developed *in vivo*.

We expected that the aggregates could be further cultured in the abdomen of host and develop tooth germ. We transplanted 1:5 aggregates into the abdominal region of embryos and raise the

host embryos until stage 66. We found that cartilage differentiated into membranous bone and surrounding the membranous bone, tooth germ-like structures was developed in the transplanted tissues. Amelogenin, a main component of tooth enamel matrix proteins (Fukae and Shimizu 1974, Satchell *et al.*, 2002), is expressed in ameloblasts at highest concentration during tooth formation. Amelogenin is present in *Xenopus* (Toyosawa *et al.*, 1998). We detected amelogenin protein in tooth germ-like structures in the transplanted tissue, which resembled normal tooth germ. Thus, tooth germ was developed from the 1:5 aggregates transplanted into tadpoles. Currently no master gene such as *Pax6* in eye development is known for tooth development. The experimental differentiation system described here may allow the discovery of such regulatory genes in tooth development. Actually, we have detected several genes related to neural crests in the 1:5 aggregates compared with the 1:1 aggregates. We will further investigate gene expression in the 1:5 aggregates to understand a gene cascade to tooth induction.

Deciphering how to manipulate cell differentiation during early development is essential for functional applications. The findings presented here indicated that activin A regulates body patterning and cell fate and it induces a developmental cascade for maxillofacial structures including jaw and tooth. Activin A, or activin-like signal such as nodal, has been considered an important morphogen during mammalian development (Vincent *et al.*, 2003) and in activin  $\beta$ A mutant mouse embryos incisor and mandibular molar teeth fail to develop beyond the bud stage (Ferguson *et al.*, 1998). Matzuk *et al.*, (Matzuk *et al.*, 1995) also reported that activin receptor II-deficient mice showed skeletal and facial abnormalities reminiscent of the Pierre-Robin syndrome in humans, which is characterized by brachygnathia/micrognathia. Mouse ES embryoid bodies treated with activin A expressed genes found at the organizer reported in *Xenopus* (Johansson B.M. and M.V. 1995, Perea-Gomez *et al.*, 2001). We believe that the differentiation system described here can be applied to mammalian ES cells. This methodology will promote an understanding of the regulating mechanisms of body patterning and tooth induction and it may aid in the development of engineered tissues for tissue replacement therapies in humans.

## Materials and Methods

### Aggregation culture method

Fertilized eggs were obtained from both male and female *Xenopus laevis* injected with 600 IU of gonadotropin (Gestron; Denka Seiyaku Co., Kawasaki, Japan) and their jelly coat were removed with Steinberg's solution containing 4.5% cysteine hydrochloride (pH 7.8) (Ariizumi *et al.*, 1991). Animal cap sheets cut from stage-9 *Xenopus* embryos were dissociated in CMF-Steinberg's solution containing 0.1 % fatty acid-free bovine serum albumin (Sigma Chemical Co.). We treated the dissociated cells with 25 ng/ml activin

A for 1 h and washed away activin A. Then, we mixed and aggregated the activin A-treated dissociated cells with untreated dissociated cells at the ratio of 1:5 (1:5 aggregates) and 1:1 (1:1 aggregates), or the activin A-treated cells only (AA-aggregates) and cultured in RDX medium (Fukui *et al.*, 2003) on 96-well plates (Sumitomo Bakelite Co., Tokyo, Japan) in a humidified atmosphere of 5% CO<sub>2</sub> at 20°C.

### Histological examination

The aggregates were fixed with 4 % paraformaldehyde and then processed into paraffin for serial section (6 µm). After deparaffinization, the sections were stained with Alcian blue and PAS and counterstained with hematoxyline for light microscopy. For whole mount Alcian blue staining, the aggregates fixed with 4 % paraformaldehyde on culture day 7 were stained with Alcian blue for 16 h and destained in 80% methanol/20% acetic acid. The epithelium of the aggregates was manually removed with fine forceps under stereoscopic microscope for light microscopy.

### Cell-lineage analysis in the aggregates

Embryos at the 2-cell stage were injected with a total volume of 3 nl of 1% Texas Red-dextran-amine (TRDA, D-1863; Molecular Probes) or 1% fluorescein-dextran-amine (FDA, Molecular Probes) and raised up to stage-9 (Kuroda *et al.*, 2002). The TRDA-labeled animal caps were cut, dissociated in CMF-Steinberg's solution, treated with activin A and then aggregated with the untreated dissociated cells of FDA-labelled animal caps as described above. On culture day 7, the aggregates were fixed with 4 % paraformaldehyde and processed into paraffin for serial section (6 µm). The deparaffinized sections were stained with Alcian blue and analyzed with fluorescence microscope and the images were captured with a charge-coupled device camera (Hamamatsu Photonics, Hamamatsu, Japan) and AQUAMARINE software (Hamamatsu Photonics).

### RNA analysis

Total RNA was extracted from the aggregates by guanidine-thiocyanate-phenol-chloroform extraction method. One µg of total RNA were treated with ribonuclease (RNase)-free deoxyribonuclease (DNase) I (Invitrogen) and RNase inhibitor (TOYOBO). RT-PCR was then performed according to the instruction of GIBCO RT-PCR applications. Each cDNA was amplified by PCR using the following specific primer pairs: *goosecoid* (*gsc*) (Cho *et al.*, 1991), 5'-taacatttggctgcccagacc -3', 5'-cttgaactgtccacaacacg-3'; *X-Hoxa2* (Pasqualetti *et al.*, 2000), 5'-gacgattaaggacagcttataccaa-3', 5'-aactatagctgtctctccagcaa-3'; *Sox9* (Spokony *et al.*, 2002), 5'-cacatttgggaaaactgct-3', 5'-aaatcgcaagaaaagctgga-3'; *Xnot* (von Dassow *et al.*, 1993), 5'-atacatggttgccactga-3', 5'-cttctacagtccacatc-3'; *Hoxb9* (Wright *et al.*, 1990), 5'-ctgggacccaacattacac-3', 5'-tggtttgatgagtcgggtca-3'; *endodermin* (Sasai *et al.*, 1996), 5'-tattctgactctgaagtg-3', 5'-gagaactgccatgtgctc-3'. Primers for Pax6 (Onuma *et al.*, 2002) and an internal control, ornithine decarboxylase (ODC) (Furue *et al.*, 2002) were described previously. The expression rate is a percentage of the aggregation numbers in which a gene is detected divided by the experiment numbers. Significant differences were determined by Fisher's exact test.

### Sequences

PCR products were inserted directly into pGEM-T vector (Promega), according to the manufacture's instructions. The inserts were sequenced by using the dye terminator-cycle sequence method

with a DTCS kit (Beckman Coulter) and CEQ 2000 DNA analysis system (Beckman Coulter). The cloned PCR products were confirmed by sequence analysis as respective cDNA.

### In situ hybridization

Recombinant plasmids of *X-dll4*, *Col2*, *Sox9*, *gsc* and *Pax6* were used as templates for synthesis of RNA probes. The digoxigenin (DIG)-labeled RNA sense and antisense probes were prepared from template cDNA, according to the RNA-labeling kit instructions (Roche Molecular Biochemicals). We carried out *in situ* hybridization on Discovery™ (Ventana Medical Systems, Inc. USA), using with DIG-labeled probes, Ribomap™ and BlueMap™ kit (Ventna).

### Transplantation into embryos and adults

We made aggregates from GFP-transgenic *Xenopus* embryos as described above. We transplanted the 1:5 aggregates cultured for 24 h into the abdominal region of stage-23 embryos. The tissues were fixed with 4 % paraformaldehyde, then processed into paraffin for serial section (6 µm). The sections were stained with Alcian blue and PAS and counterstained with hematoxyline for light microscopy.

### Immunolocalization of amelogenin

The serial section were immunostained with an anti-amelogenin rabbit polyclonal antibody (Uchida *et al.*, 1991), visualized with peroxidase-conjugated Simple Stain MAX PO goat anti-rabbit IgG (Nichirei) and 3-amino-9-ethylcarbazol (AEC, Nichirei). Nuclei were counterstained with hematoxylin.

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