

# Coordinate involvement of Nodal-dependent inhibition and Wnt-dependent activation in the maintenance of organizer-specific *bmp2b* in zebrafish

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**ABSTRACT** A vertebrate signaling center, known in zebrafish as the organizer, is essential for axis patterning and formation and is regulated by multiple cell signaling pathways, including Wnt, Nodal, and Bmp. Organizer-specific Bmp2b plays important roles in the maintenance of the Bmp activity gradient and dorsal-ventral patterning. However, it is unknown how transcription of *bmp2b* in the organizer is regulated. In this study, we generated a *bmp2b* transgenic line *Tsgf-2.272bmp2b:gfp* that reproduced organizer-specific *bmp2b* expression. Dissection analysis revealed that a 0.273-kb minimal promoter was indispensable for *bmp2b* expression in the dorsal organizer. Reporter assays showed that organizer-specific *bmp2b* is negatively regulated by the Nodal signal and positively regulated by the Wnt signal in both embryos and cell lines. Promoter analysis and chromatin-immunoprecipitation (ChIP) indicated that one consensus Smad-binding element (SBE) (CAGAC) and one Lef/Tcf-binding element (LBE) (AGATAA) were present in the 0.273-kb promoter, and could be directly bound by Smad2 and  $\beta$ -catenin proteins. Together, these results suggest that maintenance of organizer-specific *bmp2b* expression involves opposite and concerted regulation by Nodal and Wnt signaling.

**KEY WORDS:** *zebrafish, organizer-specific bmp2b, transcriptional regulation, Nodal, Wnt*

## Introduction

A signaling center known as the organizer in zebrafish plays an essential role in germ layer formation and body axis patterning during early embryogenesis. Embryological manipulations have confirmed that analogous signaling centers are ubiquitously present in vertebrate embryos, and are known as “Spemann Organizer” in *Xenopus*, “Hensen’s node” in chicken, and “node” in mouse. These signaling centers are responsible for controlling cell fate, and studies using transplant experiments have demonstrated that their activity can induce a second body axis (Harland and Gerhart 1997; Kelly *et al.*, 2000; Medina *et al.*, 1997; Schier 2001; Yasuo

and Lemaire 2001).

In zebrafish, the organizer forms during the shield stage at 6 hours postfertilization (hpf) in a process that involves regulation by members of multiple cell signaling pathways. Among these, Wnt signal plays an important role during dorsal organizer generation. Recent study reported that, the maternal Wnt8a which functions as the dorsal determinant, is required for the primary dorsal center localization (Lu *et al.*, 2011), while the zygotic Wnt signal restricts

*Abbreviations used in this paper:* ChIP, chromatin-immunoprecipitation; LBE, Lef/Tcf-binding element; SBE, smad-binding element.

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organizer size after gastrulation (Ramel and Lekven 2004; Schier 2001). In addition, Nodal, Fibroblast growth factors (Fgf), Bone morphogenetic proteins (Bmps) and other factors are required for the formation of the dorsal organizer and correct axis patterning (Belo et al., 2009; Hikasa and Sokol 2013; Langdon and Mullins 2011; Maegawa et al., 2006; Schier 2001; Schier and Talbot 2001; Schier and Talbot 2005).

Bmps, which are members of the TGF- $\beta$  superfamily, exhibit a broad spectrum of biological activities and function as morphogens during embryonic and organ development, contributing to the formation of bone, blood vessels, heart, kidney, neurons, and liver (Langdon and Mullins 2011; Miyazono et al., 2010; Yamamoto and Oelgeschlager 2004). In *Xenopus* and zebrafish gastrulas, Bmps are distributed in a gradient along the dorsal-ventral axis, where they play an important role in germ layer induction and dorsoventral patterning. In the ectoderm, high levels of Bmp activity specify epidermal fates, intermediate levels lead to the formation of the central nervous system (CNS), and low levels are required for induction of the neural crest. In the mesoderm, high levels of Bmp facilitate formation of the lateral plate mesoderm (LPM), blood tissues, and kidneys, whereas lower Bmp activity is essential for notochord formation (De Robertis and Kuroda 2004; Heasman 2006; Langdon and Mullins 2011; Plouhinec et al., 2013).

Bmp activity gradients are maintained by a large network of molecular regulators that act at the transcriptional and translational levels. The Bmp antagonist Chordin (Chd) contributes to low Bmp activity on the dorsal side of early *Xenopus* and zebrafish embryos, where it is secreted and binds to Bmp protein, preventing its intracellular signal transduction. In contrast, in *Drosophila*, the Chordin homolog Sog, is produced on the ventral side of the embryo, where it binds and inhibits diffusion of the Bmp homolog Dpp from the dorsal side of the embryo. The complex is then transported back to the dorsal side of the embryo for degradation (Ben-Zvi et al., 2011; Ben-Zvi et al., 2008; O'Connor et al., 2006; Plouhinec et al., 2011). Other known regulators, such as the metalloproteinase Tolloid, can bind and digest Chd. In addition, twisted Gastrulation

(Tsg) functions extracellularly to facilitate formation of a Chd-Tsg-Bmp trimolecular complex. When this complex flows to the ventral side of the embryo, the high levels of Crossveinless-2 (CV2) there contribute to degradation of Bmp (Plouhinec et al., 2011; Umulis et al., 2009; Zakin and De Robertis 2010).

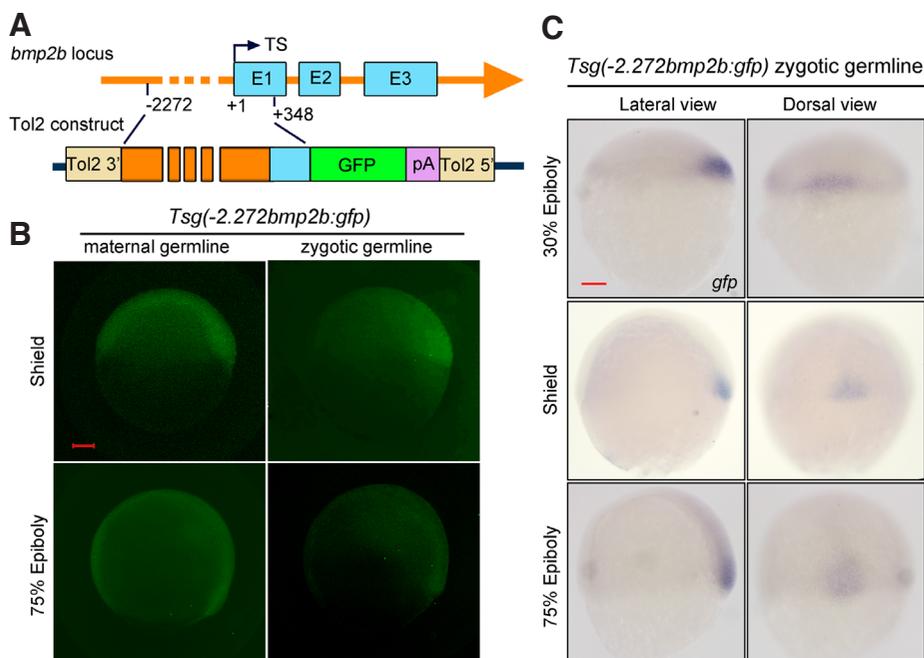
In early zebrafish gastrulas, different from other *bmp* genes with the expression in ventral mesoderm, *bmp2b* is also enriched at organizer region exclusively. Its asymmetric expression pattern is initiated by transcriptional repression by *bozozok* (*boz*) during the late blastula stage at approximately 4 hpf, the expression of *boz* in the organizer ceases from 50% epiboly stage onward (Leung et al., 2003; Solnica-Krezel and Driever 2001; Yamanaka et al., 1998). We recently showed that organizer-derived Bmp2b is required for maintenance of the Bmp activity gradient during embryonic development (Xue et al., 2014). However, the elements that mediate transcriptional regulation of organizer-specific *bmp2b* remain unknown.

In this study, we aimed to determine the functional elements that regulate organizer-specific *bmp2b* expression. We generated the novel transgenic line *Tsg(-2.272bmp2b:gfp)*, in which the 2.272-kb *bmp2b* promoter drives organizer-specific *bmp2b* expression. To determine the role of *bmp2b* promoter elements on its transcriptional regulation, we isolated a 0.273-kb minimal promoter from the main 2.272-kb sequence. The minimal promoter, which harbored one consensus Smad-binding element (SBE) and one Lef/Tcf binding element (LBE), responded to Nodal/Smad2 and Wnt/ $\beta$ -catenin signal through direct binding of these elements by Smad2 and  $\beta$ -catenin protein, respectively. Our results indicate that maintenance of organizer-specific *bmp2b* involves its direct repression by Nodal signal and activation by Wnt signal.

## Results

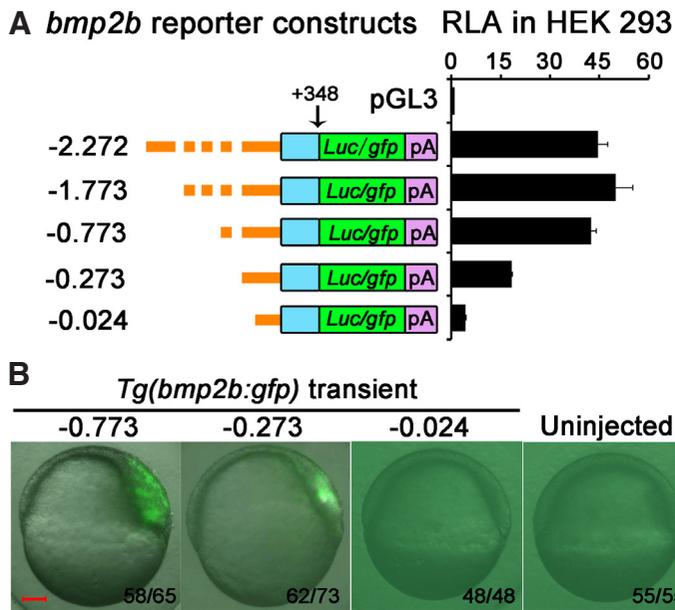
### Zygotic GFP expression of *Tsg(-2.272bmp2b:gfp)* transgenic embryos is specifically restricted to dorsal organizer region

We recently reported that a 2,620-base pair (bp) *bmp2b* pro-



**Fig. 1. Generation of the organizer-specific *bmp2b* transgenic line *Tsg(-2.272bmp2b:gfp)*.**

(A) Schematic diagram of the *bmp2b* locus and Tol2 construct. The cloned *bmp2b* promoter (-2272~+348) was inserted into the Tol2 construct for delivery into zebrafish genomic DNA. TS, transcription start site. (B) The GFP expression pattern of the *Tsg(-2.272bmp2b:gfp)* transgenic line. Embryos derived from *Tsg(-2.272bmp2b:gfp)* zebrafish crossed with wild-types at shield and 75% epiboly stage were photographed under fluorescence microscopy. In the maternal germline, embryos were derived from a female transgenic fish crossed with a wild-type male. In the zygotic germline, embryos were generated from a wild-type female crossed with a male transgenic fish. Embryos are displayed laterally, with ventral to the left. (C) Detection of the *gfp* transcripts by whole-mount in situ hybridization. *Tsg(-2.272bmp2b:gfp)* zygotic germlines were harvested at 30% epiboly, shield, and 75% epiboly stage. In the lateral view, the ventral side is oriented to the left. In the dorsal view, the animal pole is oriented to the top. Scale bars, 100  $\mu$ m.



**Fig. 2. Deletion analysis of *bmp2b* promoter.** (A) Illustration of reporter constructs and relative luciferase activity (RLA). Different truncated forms of the *bmp2b* promoter are shown with the retained promoter length indicated. The luciferase reporters driven by the *bmp2b* promoter were transfected into HEK293 cells and RLA was detected. (B) Transient GFP expression of indicated reporter constructs in embryos. Constructs containing the truncated form of the *bmp2b* promoter were injected into the embryo at the one-cell stage and GFP intensity was observed at the shield stage. Views of the embryos are lateral, with dorsal to the right. The ratio of embryos with different GFP expression pattern is indicated. Scale bar, 100  $\mu$ m.

moter region, which contains a 2,272-bp sequence upstream of its transcription start site and a 348-bp downstream sequence, has transcriptional activity specific to the dorsal region of the embryo (Xue *et al.*, 2014). To further explore the role of this promoter region on *bmp2b* expression during development, we isolated this promoter region and inserted it into a *Tol2*-mediated TSG vector to create a *Tsg(-2.272bmp2b:gfp)* construct (Fig. 1A). We then used this construct to create a *Tsg(-2.272bmp2b:gfp)* transgenic fish and got five  $F_0$  founder families, three males and two females, which could stably spread the transgene to next generation. Transgenic screening results showed that the *Tsg(-2.272bmp2b:gfp)* transgene was maternally expressed, as offspring of heterozygous females bred with wild-type males exhibited ubiquitous GFP expression. In contrast, offspring of heterozygous males bred with wild-type females showed specific but weak GFP expression on the dorsal side at both shield and 75% epiboly stages (Fig. 1B). We performed *in situ* hybridization to further characterize the zygotic *gfp* expression pattern. Dorsal *gfp* expression was initiated at 30% epiboly stage, was specifically expressed in the organizer region at shield stage, and was highest at 75% epiboly stage (Fig. 1C). This expression pattern was consistent with the expression pattern of GFP driven by the same promoter in our recent study (Xue *et al.*, 2014). These results suggest that the 2.272-kb *bmp2b* promoter region contains regulatory elements required for expression in the organizer region, but lacks elements necessary for expression in other domains, such as the ventral ectoderm. Importantly, the zygotic *Tsg(-2.272bmp2b:gfp)* germline stably expressed GFP in the dorsal organizer.

### Molecular dissection identifies a functional sequence of *bmp2b* promoter for dorsal organizer expression

In order to determine which region of the 2.272-kb *bmp2b* promoter is required for organizer-specific *bmp2b* localization, we created a *pGL3(-2.272bmp2b:luc)* construct in which the 2.272-kb *bmp2b* promoter drives expression of luciferase. A series of truncated forms of the constructs *Tg(-2.272bmp2b:gfp)* and *pGL3(-2.272bmp2b:luc)* were generated to dissect the 2.272-kb *bmp2b* promoter (Fig. 2). Deletion analysis identified a 0.273-kb proximal promoter region essential for GFP expression in embryos. Injection of the *Tg(-0.273bmp2b:gfp)* plasmid into embryos at the one-cell stage gave rise to transient GFP expression specifically (usually >80%) in the organizer region at the shield stage (Fig. 2B). This truncated promoter also drove transcription of the luciferase reporter in HEK293 cells transfected with the *pGL3(-0.273bmp2b:luc)* plasmid (Fig. 2A). Taken together, these data suggest that the 0.273-kb minimal promoter is sufficient to mediate expression of the reporter gene in the organizer region.

### Nodal and Wnt signals contribute differently to regulation of organizer-specific *bmp2b*

A question of interest to us is how organizer-specific *bmp2b* expression is controlled. We used the *pGL3(-0.273bmp2b:luc)* construct to perform a luciferase reporter assay in different cell lines. Luciferase expression was repressed by co-transfection of zebrafish Smad3b (Jia *et al.*, 2008) in Hep3B cells, suggesting that Nodal signaling inhibits *bmp2b* expression. While in HEK293 cells, luciferase expression was enhanced by co-transfection of human Lef1 which recruits  $\beta$ -catenin to Wnt target genes (Roel *et al.*, 2009) (Fig. 3A), suggesting that canonical Wnt signaling promotes *bmp2b* expression.

To substantiate the regulatory roles of these two signaling pathways on *bmp2b* expression, we knocked down a Nodal co-receptor gene *oep*, or overexpressed *lef1* in wild-type embryos (Gritsman *et al.*, 1999). We then performed *in situ* hybridization to examine the effects of these manipulations on *bmp2b* transcripts at the shield stage. Knockdown of *oep* led to an increase in *bmp2b* expression not only in the organizer but also in the ventral domain (Fig. 3B). Approximately 65% of embryos in which *lef1* was overexpressed showed a similarly increased pattern of *bmp2b* expression. The *gfp* expression in the organizer of *Tsg(-2.272bmp2b:gfp)* transgenic embryos was also enhanced by *oep* knockdown or *lef1* overexpression (Fig. 3C), consistent with the results from wild-type embryos and the cell reporter assays. These data indicate that organizer-specific *bmp2b* is inhibited by Nodal signal and activated by canonical Wnt signal. This suggests that the 0.273-kb minimal promoter contains responsive elements for Nodal and Wnt signals.

### Transcription of organizer-specific *bmp2b* is directly down-regulated by Nodal and up-regulated by Wnt signal

We next asked whether Nodal and Wnt signaling regulate organizer-specific *bmp2b* transcription directly. To address this question, we searched for Nodal/Smad2/3-responsive and Wnt/ $\beta$ -catenin-responsive elements in the 0.273-kb minimal promoter. The 0.273-kb proximal promoter region harbors one putative consensus Smad-binding element (SBE) (CAGAC, -36--32 bp) (Feng and Derynck 2005) and one putative Lef/Tcf-binding element (LBE) (AGATAA, -263--258 bp) (Blauwkamp *et al.*, 2008). When the SBE or LBE in the 0.273-kb proximal promoter region

**Fig. 3. The 0.273-kb *bmp2b* dorsal promoter is responsive to Nodal and Wnt/ $\beta$ -catenin signaling.**

(A) Luciferase reporter expression driven by the 0.273-kb *bmp2b* promoter in response to *Smad3b* or *Lef1* stimulation in Hep3B or HEK293 cells. RLA, Relative Luciferase Activity. \*  $P < 0.05$ ; \*\*  $P < 0.01$ ; NS,  $P > 0.05$ . (B,C) Expression of endogenous *bmp2b* or of *gfp* from *germlineTsg(-2.272bmp2b:gfp)* in *oep* morphants or *lef1*-overexpressing embryos at the shield stage. Embryos are laterally viewed with dorsal to the right. The ratio of embryos with the representative expression pattern is indicated. *std-MO*, standard control morpholino; *Aslef1* RNA, antisense *lef1* mRNA. MO and mRNA were injected at doses: 4 ng for *oep*-MO and *std*-MO, 100 pg for *lef1* mRNA and *Aslef1* mRNA. Scale bar, 100  $\mu$ m.

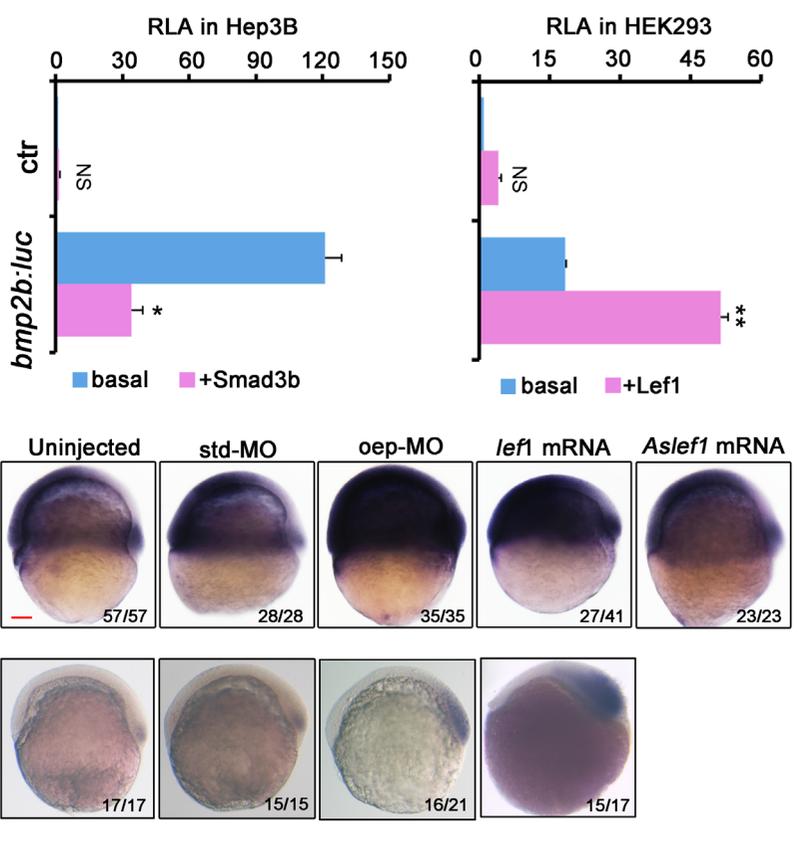
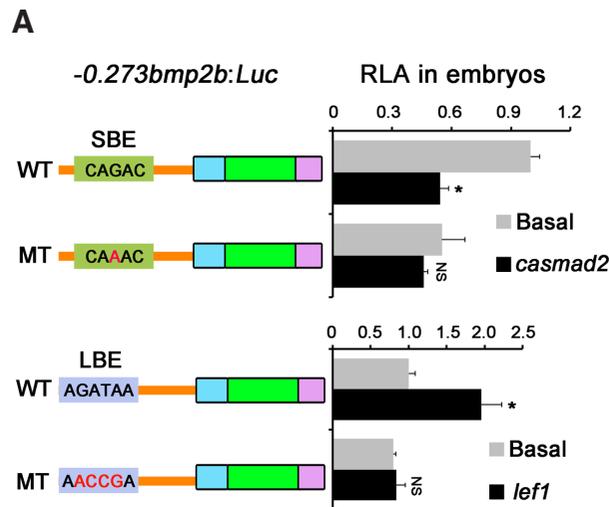
was mutated, luciferase reporter expression in embryos failed to respond to overexpression of constitutively active *smad2* (*casmad2*) (Liu et al., 2013) or to *lef1* overexpression (Fig. 4A). Thus, the SBE and LBE are required for the response of the *bmp2b* promoter to Nodal repression and Wnt activation, respectively.

Finally, we used chromatin-immunoprecipitation (ChIP) to investigate whether endogenous Smad2 or  $\beta$ -catenin could bind the SBE and LBE in the 0.273-kb proximal promoter region in zebrafish embryos. ChIP assays revealed that DNA immunoprecipitated using an anti-Smad2 antibody or an anti- $\beta$ -catenin antibody could be amplified using specific primers spanning those elements (Fig. 4B). This indicates that in embryos, the SBE and the LBE regions are bound by Smad2 and  $\beta$ -catenin, respectively.

Taken together, our data indicate that expression of *bmp2b* in the organizer is regulated negatively by Nodal/Smad2/3 signaling and positively by Wnt/ $\beta$ -catenin signaling in zebrafish embryos.

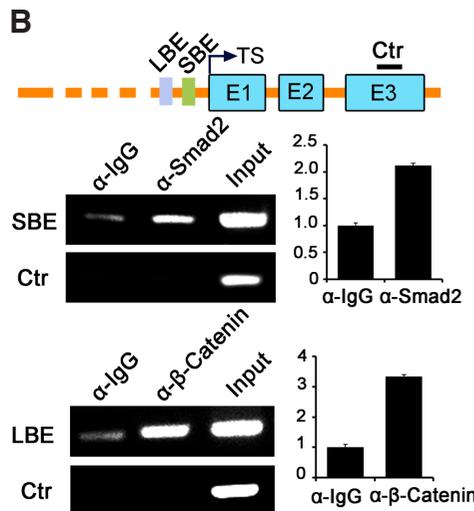
### Discussion

In this study, we identified a 2.272-kb *bmp2b* promoter that drives stable and specific GFP expression in the dorsal organizer region in transgenic gastrulas. In addition, we used molecular dissection

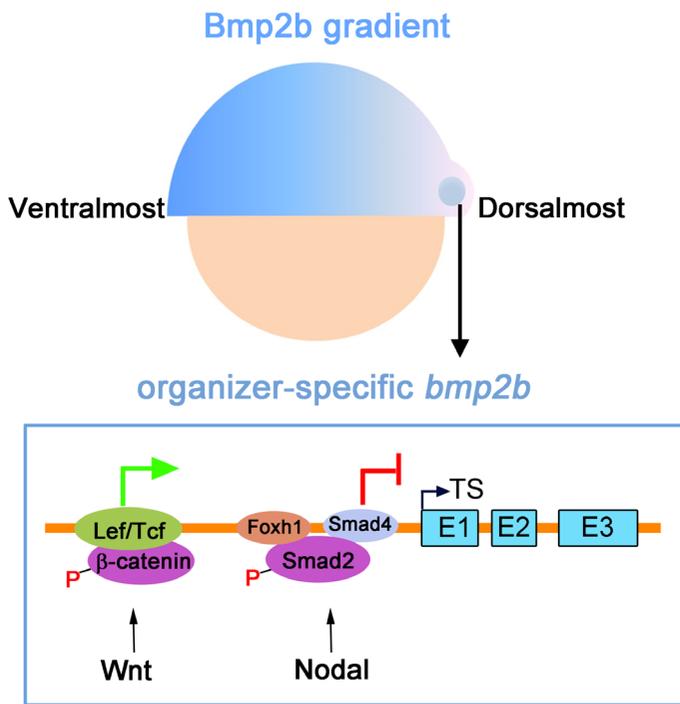


to show that a 0.273-kb minimal promoter was sufficient to express *bmp2b* in the organizer at the shield stage. This 0.273-kb minimal promoter contained a Smad2 binding site and a Lef1 binding site and was directly repressed by Nodal signal and activated by Wnt signal (Fig. 5).

In zebrafish, *bmp2b* mRNA is observed in the organizer region beginning at around 5.7 hpf, at the onset of organizer formation. *bmp2b* mRNA is further enriched in the organizer region and the dorsal side of the embryo throughout the gastrula stage. As mentioned earlier in the text, *baz*, a gene downstream of the Wnt signaling pathway, acts as the earliest repressor of *bmp2b* tran-



**Fig. 4. The 0.273-kb *bmp2b* dorsal promoter is directly controlled by Nodal and Wnt/ $\beta$ -catenin signaling.** (A) Effect of mutations in SBE or LBE of the 0.273-kb *bmp2b* promoter on luciferase reporter expression. The construct structures are illustrated on the left and RLA in injected embryos at the shield stage are shown on the right. (B) Smad2 or  $\beta$ -catenin occupancy of the *bmp2b* promoter. On the top, illustration of the amplified regions after ChIP; bottom, gel images (left) and quantification (right) of ChIP-PCR products. \*  $P < 0.05$ ; \*\*  $P < 0.01$ ; NS,  $P > 0.05$ .



**Fig. 5. Molecular regulatory model of organizer-specific *bmp2b*.** *Top image shows Bmp2b distribution at the shield stage. Embryos are viewed laterally with dorsal to the right. Bmp2b forms a gradient distribution along the ventral-dorsal axis, with the highest expression ventralmost, a gradual reduction in expression on the dorsal side, and specific enrichment at the dorsal organizer. Bottom image shows transcriptional regulation of organizer-specific *bmp2b* is mediated through activation by canonical Wnt signaling via Lef/Tcf1 binding, and repression by Nodal signaling via Smad2 binding on its promoter.*

scription in the dorsal field by directly binding to the *bmp2b* intron in the late blastula. When the organizer forms, *Boz* expression ceases, releasing repression of *bmp2b* and promoting its renewed expression in the organizer (Leung *et al.*, 2003; Solnica-Krezel and Driever 2001). Nodal and Wnt signals are required for organizer formation, and their functions are conserved across vertebrates (Schier 2001). Here, we focused on the transcriptional regulation of organizer-specific *bmp2b* at the shield stage. Our results confirmed that Nodal and Wnt signals are involved in the maintenance of organizer-specific *bmp2b* expression through direct binding of a Smad-binding site and Lef-binding site, respectively, in the 0.273-kb promoter (Fig. 4). The functional effects of Nodal and Wnt signals on organizer-specific *bmp2b* transcriptional regulation were opposite of one another, with Nodal signal repressing *bmp2b* expression and Wnt signal activating it. The interaction between these two signals deserves further investigation.

The 2.272-kb *bmp2b* promoter initiated *gfp* reporter gene expression in the dorsal side of the embryo at 30% epiboly stage (Fig. 1C). This was earlier than initiation of endogenous dorsal *bmp2b* expression (Xue *et al.*, 2014), suggesting that the 2.272-kb promoter lacks the binding elements of transcriptional repressor such as *Boz*, which is necessary for inhibition of dorsal *bmp2b* transcription at the early gastrula stage through targeting its first intron (Leung *et al.*, 2003). We also deleted a 50 bp region (-123--74 bp) of the 0.273-kb promoter to further explore its regulation. This deletion

led to the permanent absence of *gfp* expression (data not shown), suggesting that an important enhancer is embedded within this 50 bp sequence. As Fgf signal and other factors are essential for organizer formation (Kuo *et al.*, 2013; Maegawa *et al.*, 2006), it is likely that other signals or transcriptional factors work together with Nodal and Wnt signals to regulate organizer-specific *bmp2b* transcription. This hypothesis should be investigated in future studies.

One interesting result was that mutation of SBE included in the 0.273-kb *bmp2b* promoter caused almost loss of responsiveness to *casmaid2* repression. While, this mutation also reduced the luciferase reporter expression in embryos at basal level (Fig. 4A). This phenomenon may be explained that perhaps SBE not only functions as a repressor, but also an enhancer for some other factors, that is, there may exist some co-activators essential for transcriptional activity, which can form complex with Smad2/3 to function. Mutation of SBE both suppresses the binding of the complex to SBE and down-regulates the transcriptional activity of reporter gene.

As we mentioned in the introduction, maternal and zygotic Wnt signal function differently during dorsoventral patterning, previous study suggested that Wnt signal is inactive at the dorsal margin during late gastrulation (Lu *et al.*, 2011; Ramel and Lekven 2004; Shimizu *et al.*, 2012). One interesting issue is if Wnt signal continuously activates the expression of *bmp2b* at dorsal margin after shield stage? To explore this, we incubated wild-type embryos with a Wnt inhibitor, IWR-1-endo (1:5000, Selleck) (BurrIDGE *et al.*, 2014) from 64-cell stage. Compared with control group treated with DMSO, 100% of embryos treated with IWR showed observably reduced *bmp2b* expression in organizer region at shield stage (Fig. S1A), while not clearly decreased at dorsal margin at 75% epiboly stage (Fig. S1B). These results suggested that Wnt-signal is essential for *bmp2b* expression in organizer region but becomes weak and contributes less to *bmp2b* expression at dorsal margin during late gastrulation. We speculate that other genes expressing in this region compensate for the dorsal *bmp2b* maintenance at these stages.

In conclusion, we showed that organizer-specific *bmp2b*, which is essential for maintenance of the Bmp2b activity gradient, and more broadly, for dorsoventral patterning, was regulated by Nodal and Wnt signals in an opposite manner, thereby maintaining a balance of *bmp2b* expression in the dorsal organizer.

## Materials and Methods

### Zebrafish embryos and cell lines

Zebrafish embryos, the human 293 cell line, and the Hep3B cell line were used in this study. All animal studies were performed in accordance with guidelines approved by the institutional Human Ethics Review Committee and the Animal Care and Use Committee of Tsinghua University.

### Molecular cloning

A sequence between base pairs -2272 to +348 of the *bmp2b* locus was amplified by PCR from zebrafish genomic DNA and cloned into the *pGL3* basic vector (Promega) to form the construct *pGL3(-2.272**bmp2b**:luc)*. This insert was then subcloned into a modified version of the *pGL3* basic vector in which *Luciferase* was replaced by the enhanced green fluorescent protein (*GFP*) coding sequence to generate the construct *Tg(-2.272**bmp2b**:gfp)* for transient expression and deletion analysis. The primer sequences used for PCR amplification of DNA were the same as those used previously (Xue *et al.*, 2014).

To create the construct for antisense *lef1* mRNA, *lef1* coding sequence was amplified by PCR from cDNA, then was reversely inserted into *pX77*

vector. The primer sequences were: Forward-Spel: 5'-GGACTAGTatgc-cgcagttgtcagggtgga-3'; Reverse-EcoRI: 5'-CCGGAATTCcagatgtaccg-cgttttcattc-3'.

### Zebrafish strains and transgenesis

The tuebingen zebrafish strain was used in this study. To generate the transgenic line, the *bmp2b* promoter was cloned into a modified TSG vector (a map of the modified TSG vector used in this study can be provided upon request) (Han *et al.*, 2011). The plasmid *Tsg(-2.272bmp2b:gfp)* was then co-injected with Tol2 transposase mRNA into one-cell wild-type embryos. These founder fish were then mated to wild-type fish, and progeny that carried the transgene were identified by GFP expression. Embryos born from male founders showed specific but weak GFP expression that could be detected by *in situ* hybridization, while offspring of female founders displayed strong maternal GFP expression.

### RNA synthesis, morpholinos, microinjection and whole-mount *in situ* hybridization

mRNAs were synthesized *in vitro* from corresponding linearized plasmids using the mMessage mMachine kit (Ambion). Digoxigenin-UTP-labeled antisense RNA probes were generated by *in vitro* synthesis with a linearized plasmid as a template (Roche).

Whole-mount *in situ* hybridization was performed as previously described (Xiong *et al.*, 2006). Stained embryos were cleared in glycerol and photographed using a digital camera (SPOT Insight) under a Nikon SMZ 1500 microscope. The images were adjusted using Adobe Photoshop software.

The morpholinos used in this study were synthesized by Gene Tools, LLC and was as follows: oep-MO, 5'-GCCAATAAACTCCAAAACAACACTCGA-3' (Nasevicius and Ekker 2000). standard-MO, 5'-AAGGAAAAACGAAATG-GAAAGGAT-3'.

Approximately 1-1.5 nl of morpholino solution or DNA was injected into the yolk or cell of each embryo at the one-cell stage using MPPI-2 quantitative injection equipment (Applied Scientific Instrumentation Co.).

### Luciferase reporter and chromatin immunoprecipitation assay

Cell culture, transfection, and luciferase reporter assays were performed as previously described (Liu *et al.*, 2013). Shield stage embryos were harvested for embryonic reporter assays.

The chromatin-immunoprecipitation (ChIP) assay was performed as described previously (Liu *et al.*, 2011). For this experiment, approximately 1500 embryos at shield stage were harvested. A rabbit anti-Smad2/3 antibody (Cell Signaling, 1:50) and a rabbit anti- $\beta$ -catenin antibody (Abcam, 1:200) were used. The primer sequences for ChIP-PCR were: *bmp2b*-forward: 5'-ATACAAATGTAGATAATTTA-3'; *bmp2b*-reverse: 5'-TTTTTGTTGTCT-GATGATGT-3'; control-forward: 5'-ACACCCGTCTGGTGCAGGAC-3'; control-reverse: 5'-AGCCTCCTCGGATACTTC-3'. The ChIP-PCR products were examined by gel electrophoresis and band intensity was quantified using ImageJ software.

### Statistical analysis

Student's *t*-tests (two-tailed, unequal variance) were used to determine *p*-values of all groups compared in this study. Significance levels were denoted as \**p*<0.05 and \*\* *p*<0.01.

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### Author Contributions

Conceived and designed the experiments: YX CX AM YP. Performed

the experiments: YX CX CC WZ JX. Analyzed the data: YX CX AM. Contributed reagents/materials/analysis tools: YX CX AM YP. Wrote the paper: YX CX AM.

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