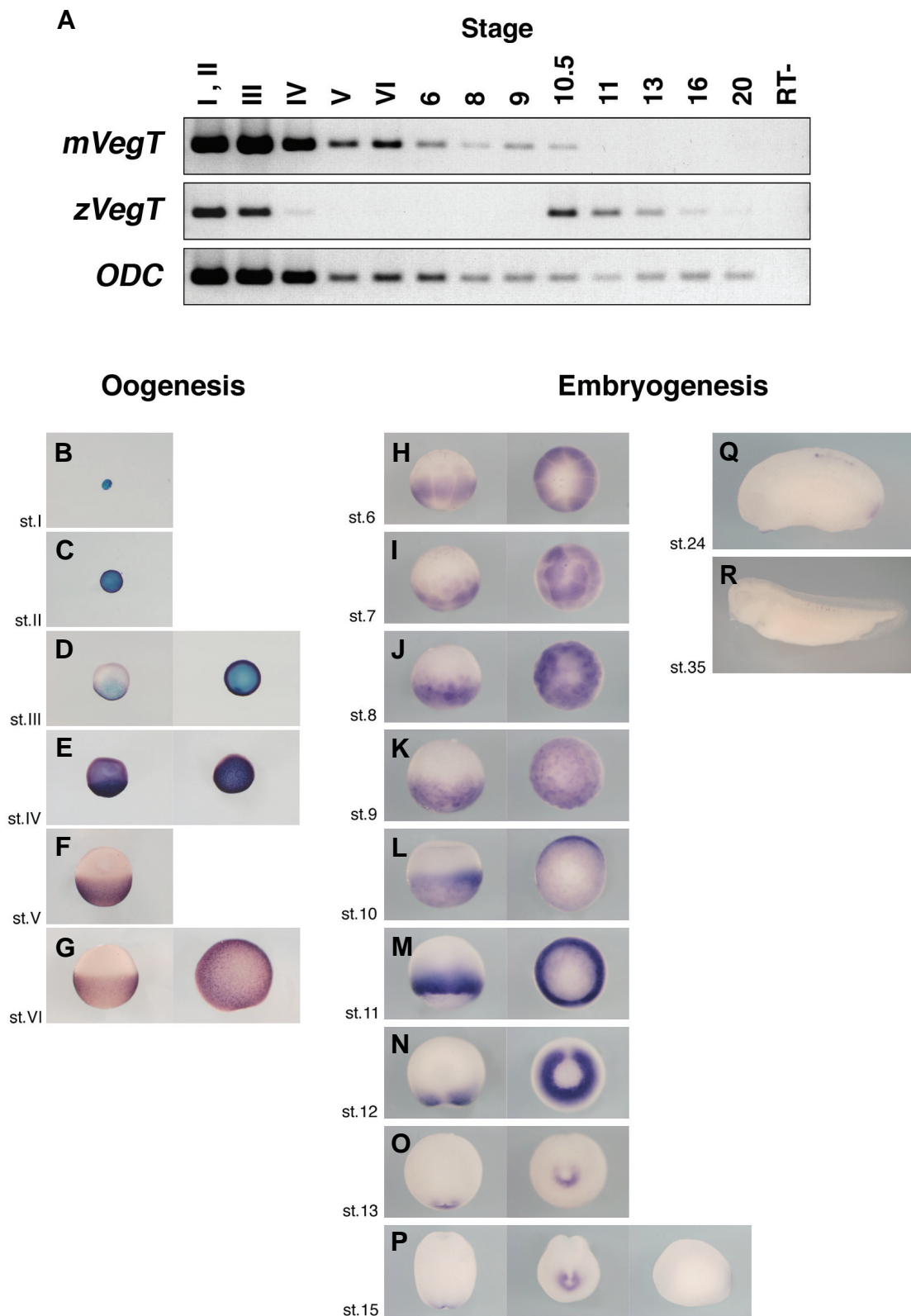


**SUPPLEMENTARY MATERIAL**

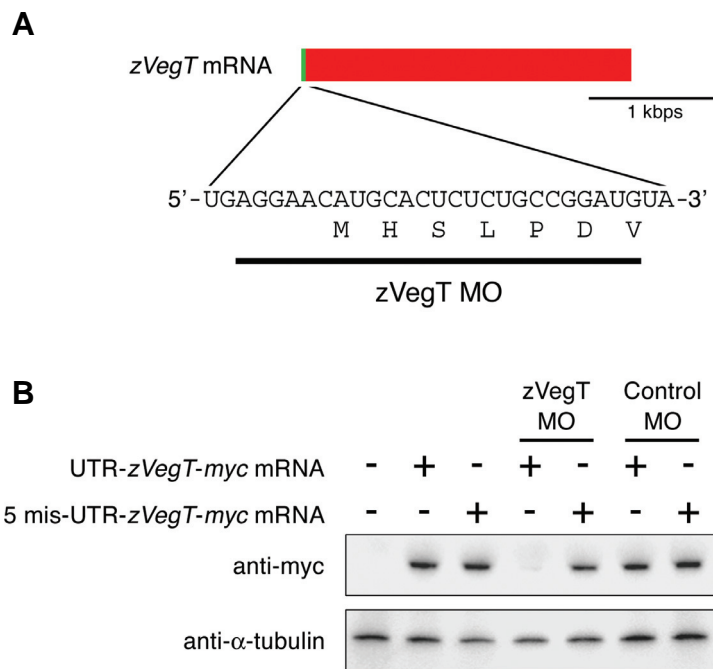
**corresponding to:**

**Zygotic *VegT* is required for *Xenopus* paraxial mesoderm  
formation and regulated by Nodal signaling  
and Eomesodermin**

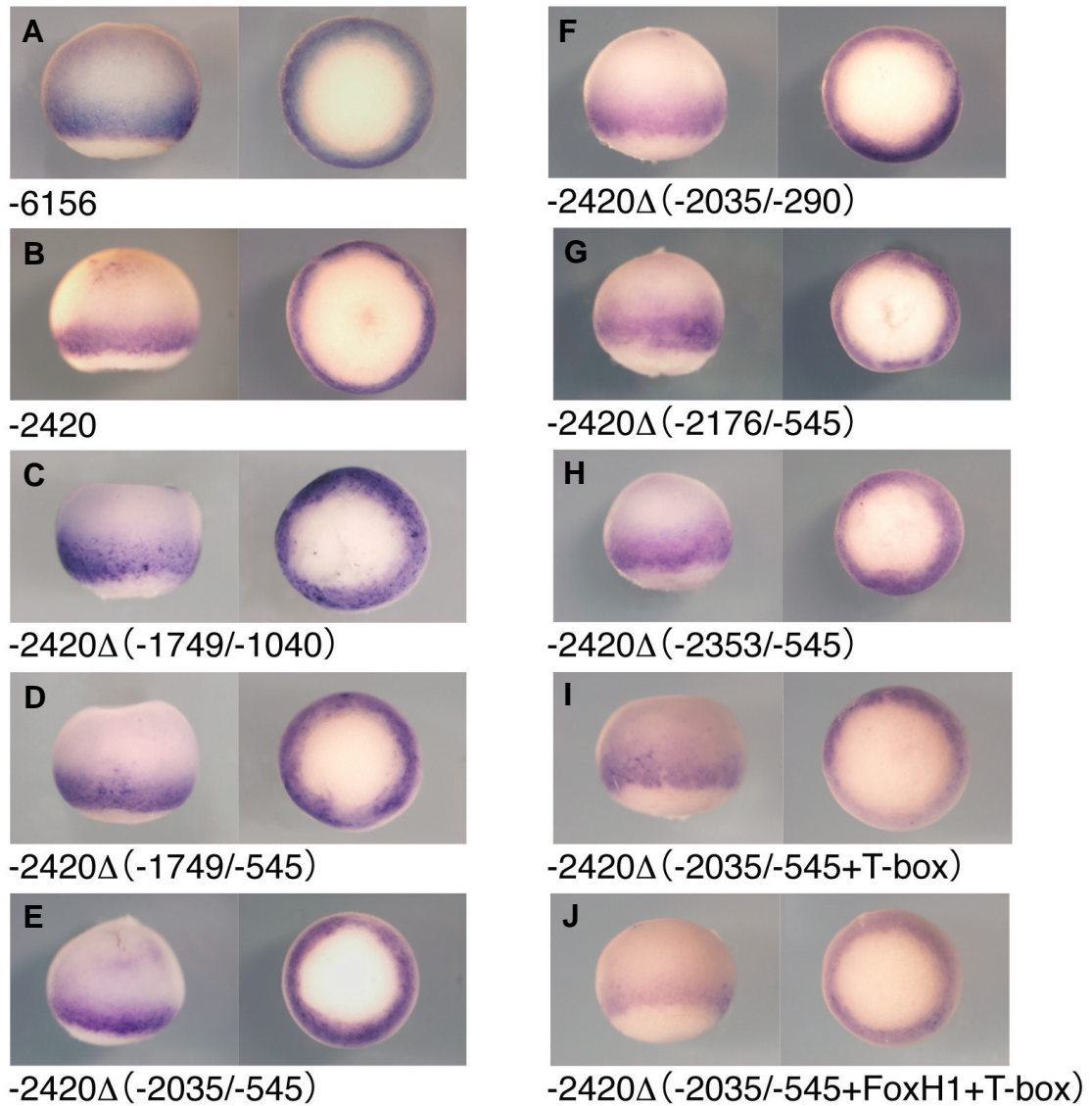
MASAKAZU FUKUDA, SHUJI TAKAHASHI, YOSHIKAZU HARAMOTO, YASUKO ONUMA, YEON- JIN KIM,  
CHANG-YEOL YEO, SHOICHI ISHIURA AND MAKOTO ASASHIMA



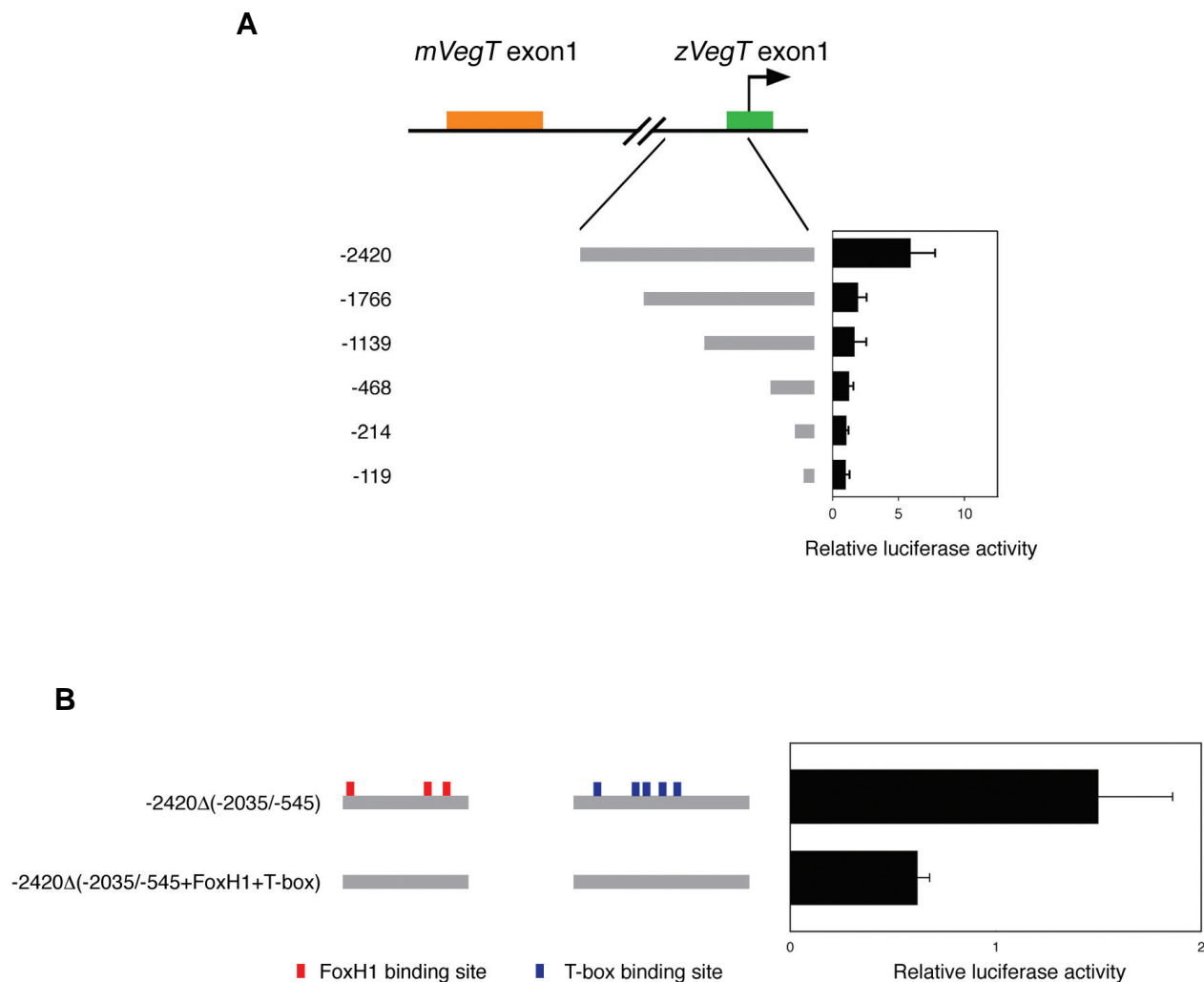
**Supplementary Fig. S1. Temporal and spatial gene expression patterns of *VegT* during oogenesis and embryogenesis of *X. tropicalis*. (A)** Temporal expression was analyzed by semi-quantitative RT-PCR. Primer pairs specific for *X. tropicalis* ODC were described previously (Haramoto et al., 2004). The primers designed for *X. tropicalis* *mVegT* are as follows: 5'-ATGAGAACTGCTGTCAGGAACAC-3' and 5'-TGAAACCTGGGCTTGATGCG-3'. (B-R) Spatial expression was analyzed by WISH. Roman figure shows stage of oogenesis.



**Supplementary Fig. S2. zVegT MO specifically inhibits the translation of zVegT.** (A) zVegT MO was designed to bind the translational start region of *X. tropicalis* zVegT. (B) 1 ng of UTR-zVegT-myc mRNA or 5mis-UTR-zVegT-myc mRNA was coinjected with 40 ng of zVegT MO or Control MO into both blastomeres at the two-cell stage in *X. laevis* embryos. The injected embryos were cultured until stage 9 for Western blot analysis. To generate pCS2-UTR-zVegT-myc and pCS2-5mis-UTR-zVegT-myc, the UTR and coding regions of *X. tropicalis* zVegT were amplified by PCR using different primer pairs and subcloned into the pCS2-MT vector, respectively. The construct pCS2-5mis-UTR-zVegT-myc contains five silent mutations in the MO-binding region. These mutations were generated by PCR using the following primers: forward, 5'-GAAGATGCATtCaCTtCCGGAcG-3' and reverse, 5'-CCAACAGCTGTATGGAAAGAGAG-3' (mutated bases are shown in lower case letters). Cloning products were verified by sequence analysis. For Western blot analysis, the zVegT protein with 6myc-epitope tags was detected using the anti-c-myc (9E10) HRP-conjugated monoclonal antibody (Santa Cruz Biotechnology). The internal control, α-tubulin, was detected using the monoclonal anti-α-tubulin antibody (mouse IgG1 isotype) (Sigma). Anti-mouse IgG, HRP-linked antibody (Cell Signaling Technology, Inc.) was used as the secondary antibody. zVegT MO inhibited translation from UTR-zVegT-myc mRNA, but not from 5mis-UTR-zVegT-myc mRNA.



**Supplementary Fig. S3. *X. laevis* transgenic embryos expressed the EGFP reporter gene in the marginal region.** Reporter EGFP expression was detected by WISH (Figs. 3A and 5 A,B). Panels at the left and right indicate lateral and vegetal views with dorsal side up, respectively. All embryos show equivalent expression patterns for endogenous zVegT.



**Supplementary Fig. S4. Transcriptional activity of several DNA reporter constructs in *X. laevis* embryos.** To generate firefly luciferase reporter constructs, -2420, -1766, -1139, -468, -214, -119, -2420Δ(-2035/-545), and -2420Δ(-2035/-545+FoxH1+T-box), the 5' upstream regions were amplified by PCR and inserted into the pGL3-basic vector (Promega). Schematic diagrams representing the cis-regulatory region of each construct are shown on the left of the graphs. The FoxH1 and T-box binding sites are indicated as red and blue boxes, respectively. Reporter activity was measured by the Dual-Luciferase Reporter Assay System (Promega). Each experiment was repeated at least three times in duplicate. Specific pGL3-zVegT promoter constructs (50 pg) were injected along with the Renilla luciferase construct pRL-TK (50 pg) into the marginal zone of both blastomeres at the two-cell stage, and embryos were harvested at stage 11.