

Supporting Material - Experimental Materials and Methods

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1. Axolotls

Natural matings were established as described in [1]. One or two cell embryos were placed in 1 MBS + 4 % Ficoll (Sigma) and antibiotics, then injected in the animal hemisphere with 2–4 nl injections (one per blastomere in two cell embryos). Embryos were staged according to [2], which are approximately equivalent to the stages of *Xenopus* in [4]. Animal caps were cut from embryos at stage 9 and cultured for 48 hours.

2. Quantitative RT - PCR

qPCR was performed using the ABI 7500 Sequence Detection System (Applied Biosystems) with TaqMan probes and primers as detailed below. RNA was isolated from a minimum of 10 animal caps. qRT-PCR data were analysed by the comparative CT method [3]. Validation experiments were carried out on a 4-fold dilution series of cDNAs from 1 to 1/256 to ensure the PCR efficiencies of the target and endogenous reference, (ODC), were approximately equal. The

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data were analysed in excel (Microsoft) and graphs were plotted of the ratio of gene expression relative to uninjected animal caps. Error bars are one standard deviation of the sample.

2.1. Primers and probes

Primers and probes were designed using Primer Express version 3.0 software (Applied Biosystems) according to manufacturer's instructions. Probes (Sigma) are dual-labelled fluorogenic probes (5' FAM; 3' TAMRA) and HPLC purified.

| | Forward (5'-3') | Reverse (5'-3') |
|-------------|--------------------------------|----------------------------|
| AxNodal1 | CCCAGTGGATGAAACGTTTCAG | GGGTCCGGTGGTACAGCTT |
| AxBrachyury | CATTGACCACATGTACCAATTGC | GATCAAGGGTCAATCGTGAGTTC |
| AxMix | GTCCAGGATCCAGGTCTGGTT | GCTTCTGGGTGGATTTGATTTATAA |
| AxFGF8 | TGCAGGTCCTTGGAACAA | AAGGTGTCCGTTTCCACAATTAA |
| AxSox17 | TGGATACGACGCTCCACAGA | CTCCCTGTAGTGGCCGATGT |
| AxGsc | GCCTCTTCCAGGAGACCAAGT | TGGCTCTGCGGTTCTTGAAC |
| AxODC | ATGCCCGTCATGAGTAGTACCA | CCCGGACCCAGGTTACG |
| AxNcam | TGAATGTCGTTCAACGTGAGAGA | AAGAAAAGACTCTGGATGGACGTATC |
| | Probe (5'-3') | |
| AxNodal1 | CGACGAATCATGCCTACATGCAGAGC | |
| AxBrachyury | TACCCATAGTTCTTTTGTGCAGCATCCACG | |
| AxMix | AATAGGCGTGCCAAGTCCCCGCC | |
| AxFGF8 | ACGGCGACTCGCACGCCA | |
| AxSox17 | CATGAGCAGCAGTTCCAGCAGGACAAC | |
| AxGsc | CACCCGAGAGCAGCTGGCCC | |
| AxODC | GACAGTTCCAAGGTTTCATTCAATTGCTG | |
| AxNcam | AGACCCTGGCGTGACTGCTCACCA | |

2.2. Microscopy and photography

Animal caps were visualised under Nikon SMZ 1500 microscopes. Photographs were taken using a Nikon DXM 1200F camera.

References

- [1] J.B. Armstrong and G.M. Malacinski. *Developmental biology of the axolotl*. Oxford University Press, 1989.
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- [3] K.J. Livak and T. D. Schmittgen. Analysis of relative gene expression data using real-time quantitative pcr and the $2^{-\delta \delta c(t)}$ method. *Methods*, 25(4):402 – 408, 2001.
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