

Table S1. Ectopic expression of Brachyury: DeSeq analysis of differential expression

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Table S2. Differential expression of tissue markers in response to ectopic Brachyury, Foxa.a and/or Mnx expression

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Table S3. Summary of all differential expression results for notochord enriched genes

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Table S4. Transcription factors with decreased expression in response to loss of FGF signaling by MEK inhibition

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Table S5. Ectopic expression of Brachyury, Foxa.a and Mnx: DeSeq analysis of differential expression

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Table S6. In situ probe cloning primers

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