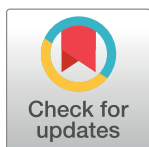


CORRECTION

Correction: Normal Levels of Sox9 Expression in the Developing Mouse Testis Depend on the TES/TESCO Enhancer, but This Does Not Act Alone

Nitzan Gonen, Alexander Quinn, Helen C. O'Neill, Peter Koopman, Robin Lovell-Badge

Errors were introduced during the production process resulting in the incorrect labelling of the genotypes in [Fig 5C](#). Please view the correct [Fig 5](#) here.



OPEN ACCESS

Citation: Gonen N, Quinn A, O'Neill HC, Koopman P, Lovell-Badge R (2017) Correction: Normal Levels of *Sox9* Expression in the Developing Mouse Testis Depend on the TES/TESCO Enhancer, but This Does Not Act Alone. PLoS Genet 13(2): e1006584. doi:10.1371/journal.pgen.1006584

Published: February 1, 2017

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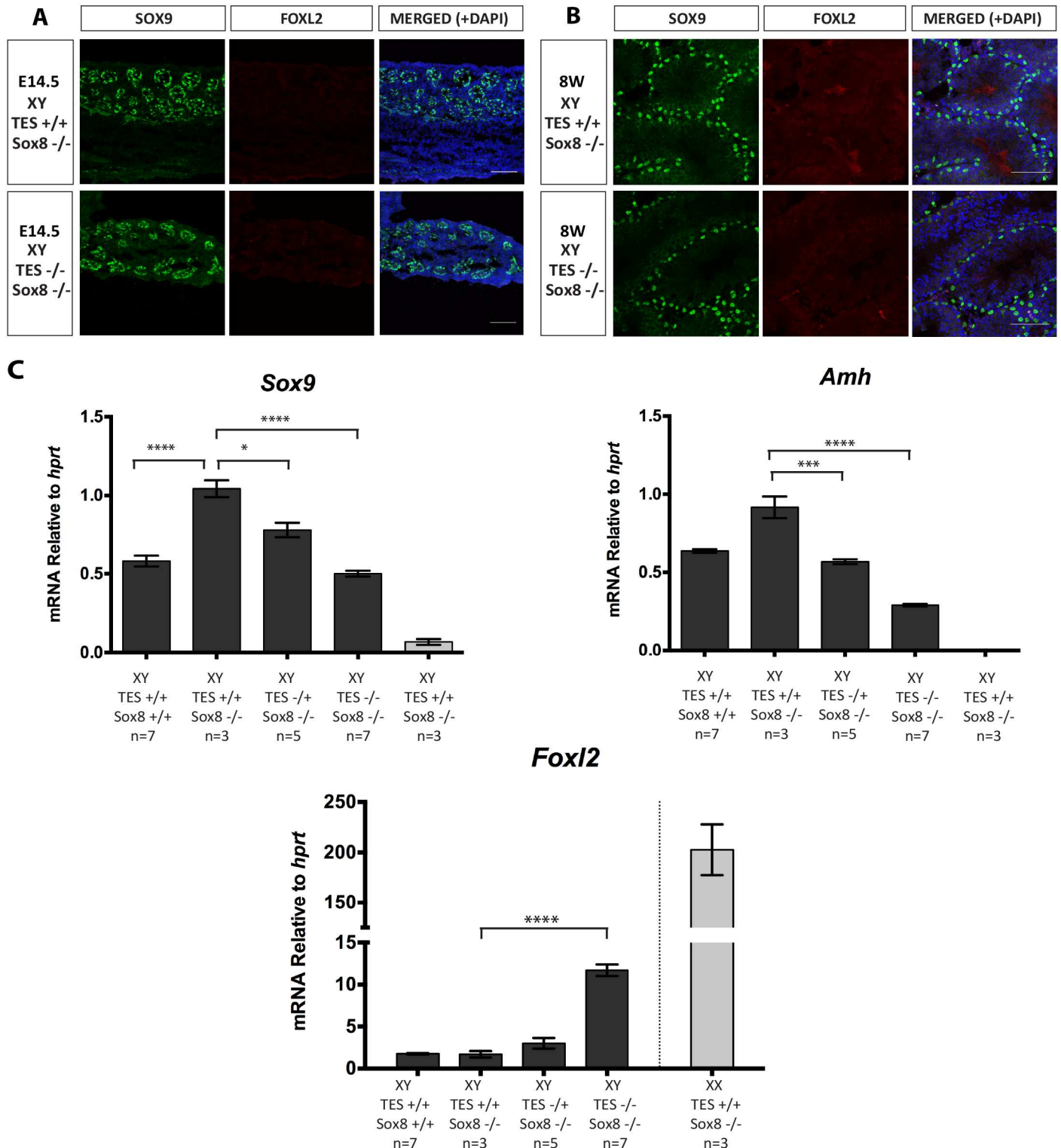


Fig 5. Immunofluorescence and real-time quantitative RT-PCR analysis of mice carrying TES deletion on Sox8-null background. (A) Immunostaining of 14.5 dpc XY testes of wild type and TES^{-/-} embryos on a Sox8-null background. (B) Immunostaining of 8 week-old XY testes of wild type and TES^{-/-} mice on a Sox8-null background. Testes were stained for SOX9 (green), FOXL2 (red) and DAPI (blue). (C) Gene expression in XY TES deleted gonads with Sox8-null background at 14.5 dpc. Data are presented as mean 2^{-ΔΔCt} values normalized to *Hprt*. Sample size represents number of individuals and is indicated below each genotype. Error bars show SEM of 2^{-ΔΔCt} values. P value is presented above the relevant bars (unpaired, two-tailed t-test on 2^{-ΔΔCt} values). Dark grey bars: XY; light grey bars: XX.

doi:10.1371/journal.pgen.1006584.g001

Reference

1. Gonen N, Quinn A, O'Neill HC, Koopman P, Lovell-Badge R (2017) Normal Levels of *Sox9* Expression in the Developing Mouse Testis Depend on the TES/TESCO Enhancer, but This Does Not Act Alone. *PLoS Genet* 13(1): e1006520. doi: [10.1371/journal.pgen.1006520](https://doi.org/10.1371/journal.pgen.1006520) PMID: [28045957](https://pubmed.ncbi.nlm.nih.gov/28045957/)