

SI DATASETS

Dataset S1: RNA-seq analysis of mouse germline from E9.5 to E11.5.

Dataset S2: Genes up-regulated following PGC colonization of the nascent gonads in mouse, and human one-to-one orthologs.

Dataset S3: Single-cell RNA-seq analysis of human germline as PGCs colonize the nascent gonads.

Dataset S4: Identification of SNPs that distinguish between 129S2 and 129S4 substrains.

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