

Regulation of Plant Stem Cell Quiescence by a Brassinosteroid Signaling Module

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In the originally published version of this paper, Table S1 was mistakenly reported to have contained primer sequences, microarray data, and quantification of QC division phenotypes. However, Table S1 actually only contained the quantifications of phenotypes, whereas the primer sequences are in the Supplemental Experimental Procedures. In addition, a separate Excel file with microarray data was missing from the paper, and this is now provided as Table S2 and available at GEO with the accession number GSE67144. This error has now been corrected in the online version of the paper. The authors apologize for any confusion this error may have caused.

Weaning Triggers a Maturation Step of Pancreatic β Cells

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In the originally published version of this paper, the order of gene sets in Figure 7A has been flipped, making it incorrectly seem as if mitotic gene sets are induced upon weaning more than G1/S gene sets. The mistake has been corrected online, and the final version of Figure 7 is shown here. The authors apologize for the error and any inconvenience that may have resulted.

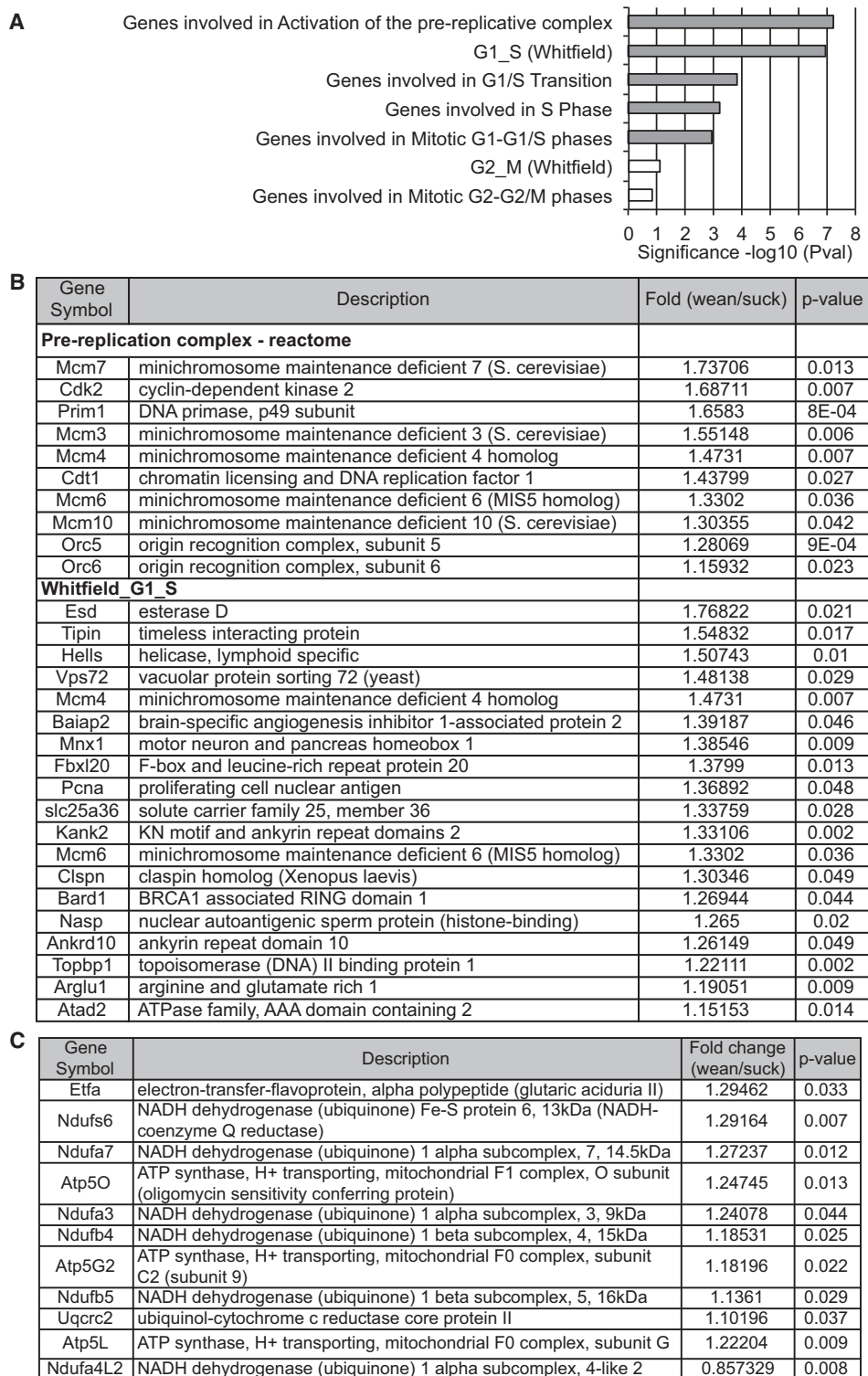


Figure 7. Transcriptome Changes in β Cells upon Premature Weaning

(A) Cell-cycle-related gene sets that are significantly induced (gray bars) or unchanged (white bars) in β cells of prematurely weaned p25 mice compared with β cells from suckling littermates. Note that gene sets that have changed are preferentially associated with early phases of the cell cycle and not with G2 and mitosis.

(B) Cell-cycle-related genes induced in GFP-positive β cells from prematurely weaned mice. Gene sets used were from Reactome pathway analysis and Whitfield et al. (2002).

(C) Oxidative phosphorylation-related genes induced in GFP-positive β cells from prematurely weaned mice.

See also Figures S4 and S5 and Table S1.