

Supplementary Material S2

Expression analysis of SelS

Legends:

Figure S2A:

Schema of SelS gene structure (gene name: VIMP), including the exons and junctions that were quantified in GTEx. The exons and junctions that do not belong to the canonical, most abundant isoform are highlighted.

Figure S2B:

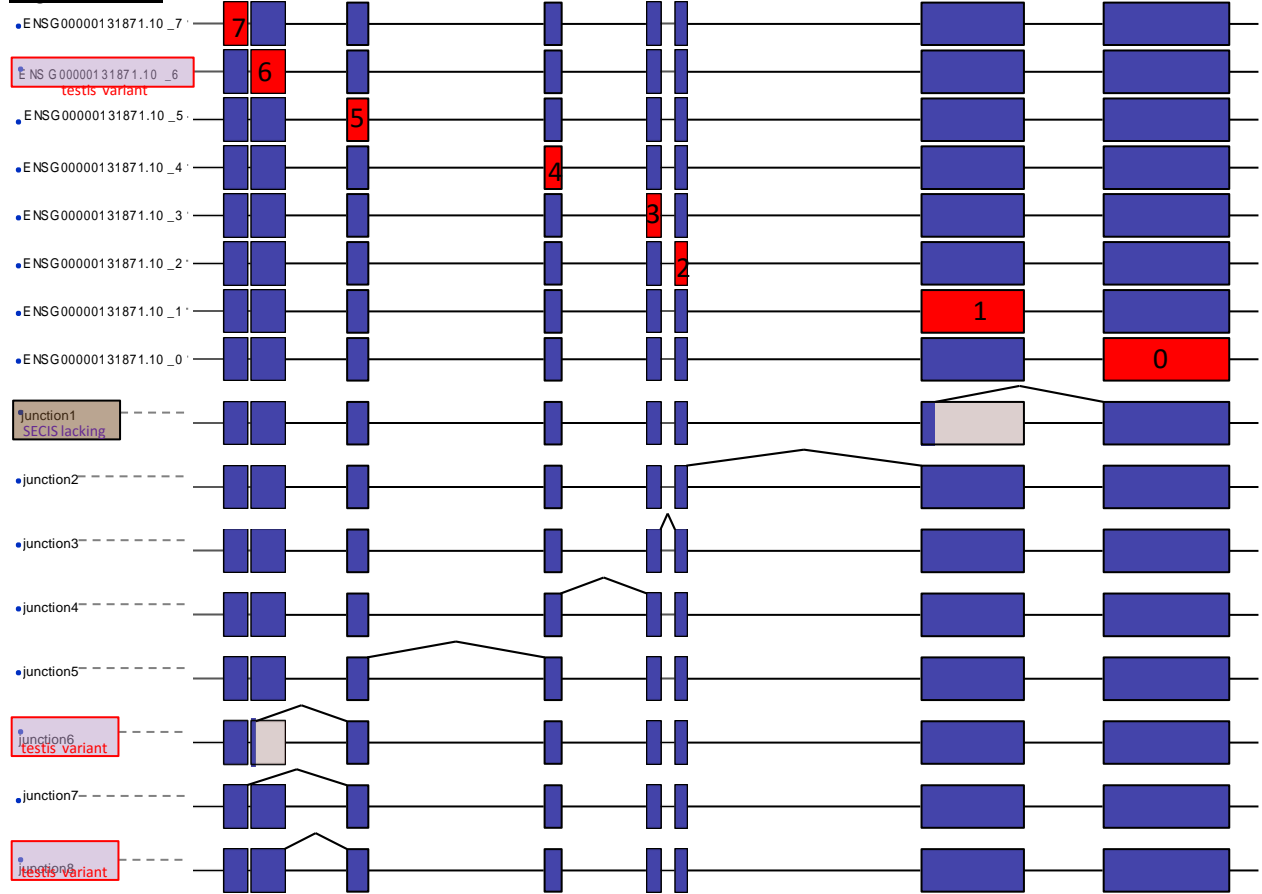
SelS exon quantification in GTEx samples, split by subtissue. Non-canonical exons are highlighted. The dotted red lines represent an arbitrary detection limit, an expression threshold of 0.1 RPKM.

Figure S2C:

SelS exon junction quantification in GTEx samples, split by subtissue. Non-canonical exons are highlighted. The dotted red lines represent an arbitrary detection limit, an expression threshold of 0.1 RPKM.

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Figure S2A:



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Figure S2B

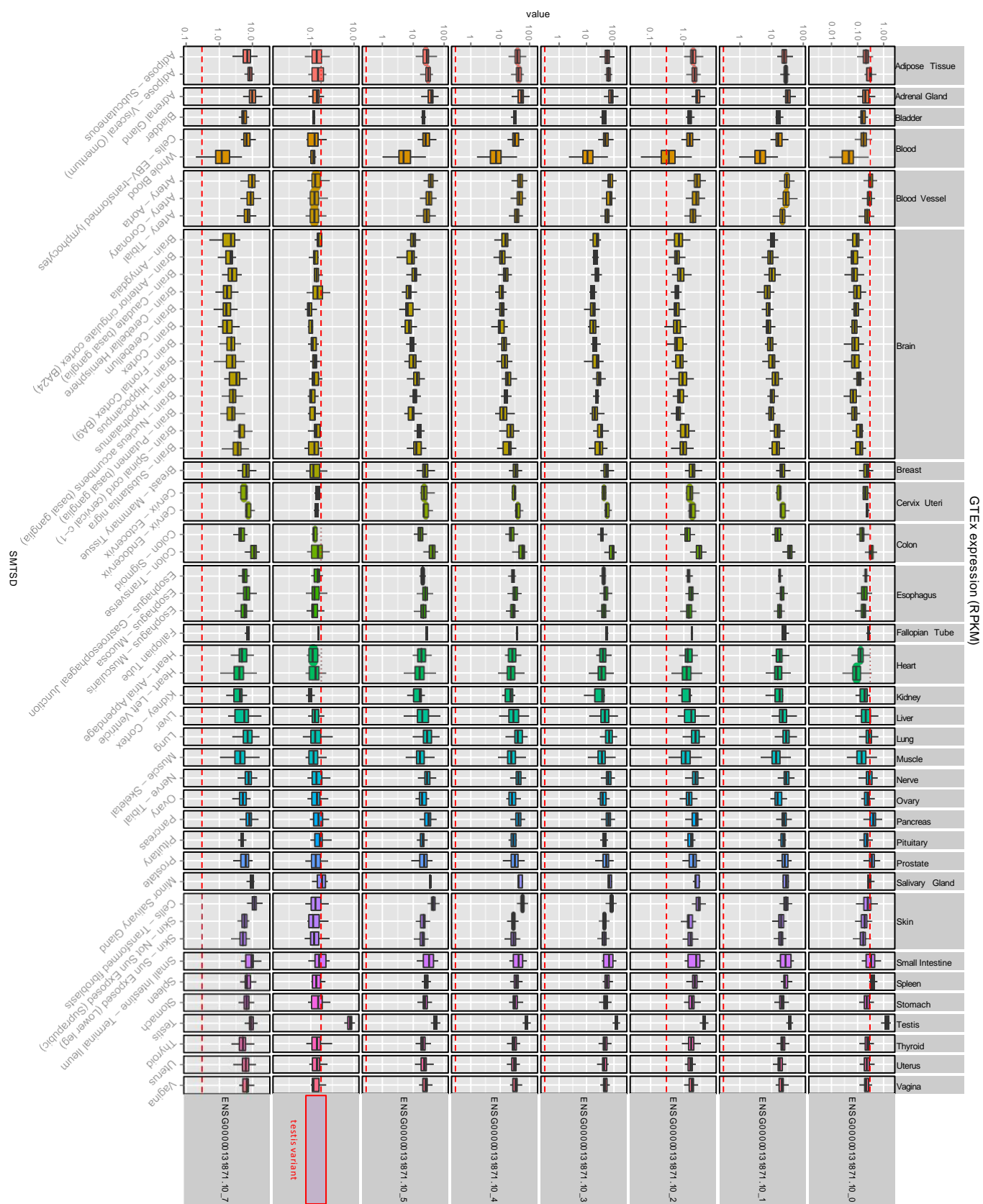


Figure S2C

GTEx expression (junction_reads / hb_size)

