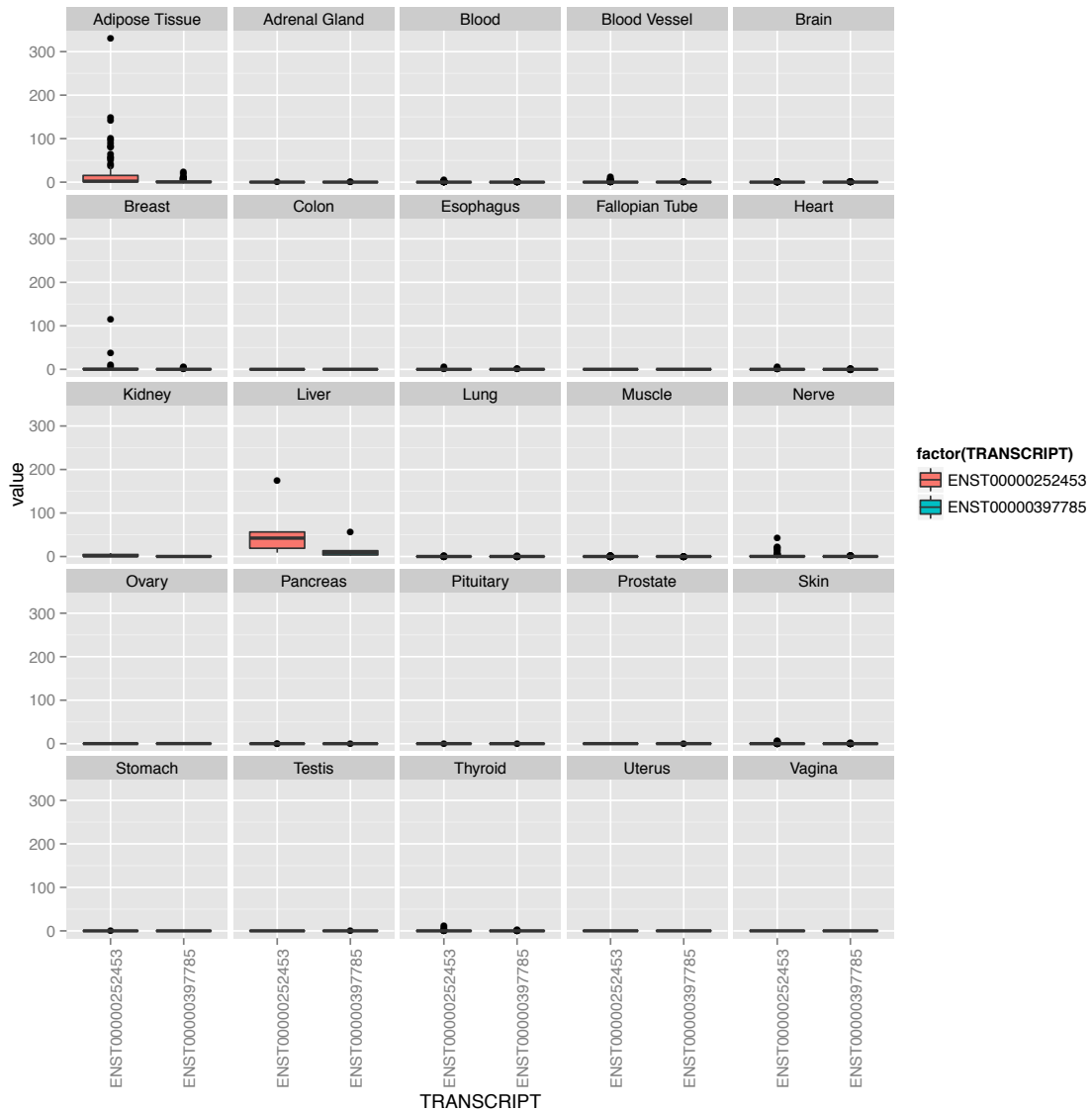


Supplementary Figure 1. *In-silico* predictions of the impact of premature termination codon p.Q121X suggests that the variant would trigger nonsense-mediated decay. Transcript isoform ENST00000252453 is highly expressed in liver with some usage of ENST00000397785 which is a lowly expressed isoform. ENST00000252453 supports the p.Q121X annotation and results in nonsense mediated decay using RNA sequencing transcript isoform data publically available from the Genotype-Tissue Expression (GTEx) project (<http://www.gtexportal.org/>) and a predictive model implemented in MAMBA (<http://www.well.ox.ac.uk/~rivas/mamba>). Y-axis is Isoform expression value (RPKM).



Supplementary Table 1. *ANGPTL8* p.Q121X and type 2 diabetes status by study

Study	p.Q121X carriers among cases	Percentage of cases who carry p.Q121X	p.Q121X carriers among controls	Percentage of controls who carry p.Q121X	Non-carriers among cases	Non-carriers among controls	Ref.
BioImage	1	0.12%	10	0.25%	814	4,062	[1]
CHARGE	15	0.16%	95	0.16%	9,509	60,623	[2]
T2D-GENES	8	0.18%	37	0.23%	4,477	15,907	[3]
Summary	24	0.16%	142	0.18%	14,800	80,592	

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