```
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GCTGGAGATCTGAGAACGGGCAGACTGC**CTCCTCAAGTGGGTC**CCTGACC**CCTGACCCCC
GAGCA**GCCTAACTGGGAGGCACCCCCCCAGCAGGGGCACACTGACACCTCACACGGCAGG
GTATTCCAACAGACCTGCAGCTGAGGGTCCTGTCTGTTAGAAGGAAAACTAACAAACAGA
AAGGACATCCACACCAAAAACCCATCTGTACATCACCATCATCAAAGACCAAAAGTAGAT
AAAACCACAAAG
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SUPPLEMENTARY FIG. S1.   DNA consensus of the L1PA2 5′UTR promoter region, generated by CLUSTALX alignment of 2230 sequences, obtained form Human Genome Build 16.