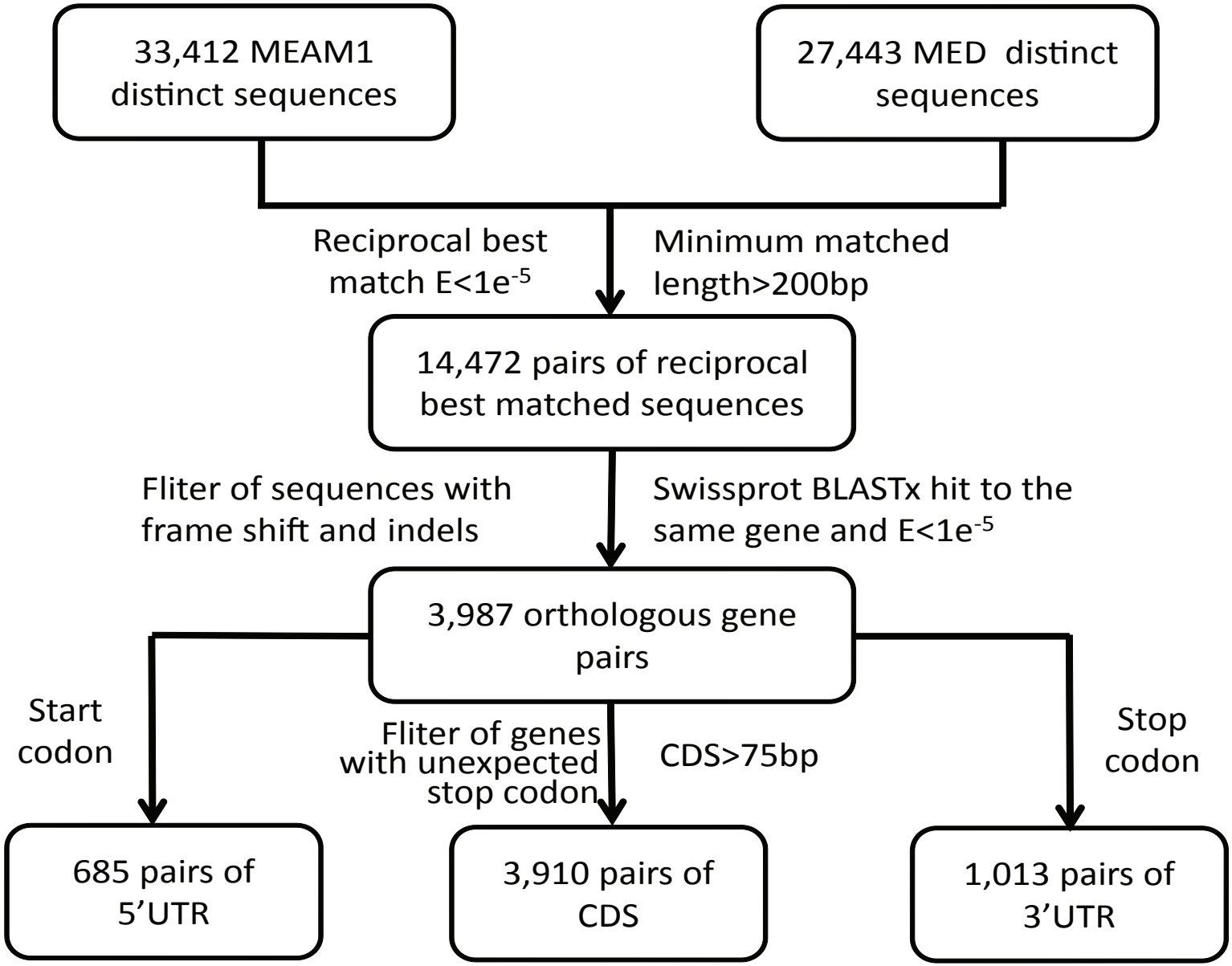
Figure S2


33,412 MEAM1 distinct sequences
27,443 MED distinct sequences
Reciprocal best match E<1e-5
Minimum matched length>200bp
14,472 pairs of reciprocal best matched sequences
Fliter of sequences with frame shift and indels
Swissprot BLASTx hit to the same gene and E<1e-5
3,987 orthologous gene pairs
Start codon
Fliter of genes with unexpected stop codon
CDS>75bp
Stop codon
685 pairs of 5'UTR
3,910 pairs of CDS
1,013 pairs of 3'UTR