

**Table S2 GenBank flags used in transcript and protein RefSeq entries**

<b>GenBank flags for exceptional cases</b>
gene /exception="dicistronic gene" [transcript RefSeq entries only]
CDS /exception="nonconsensus splice site"
CDS /trans_splicing
CDS /note="non-AUG ([ <i>codon specified</i> ]) translation initiation"
CDS /transl_except=(pos:x..y,aa:Met)
CDS /transl_except=(pos:x..y,aa:Sec)
CDS /transl_except=(pos:x..y,aa:OTHER)
CDS /ribosomal_slippage
CDS /transl_table=5
CDS /transl_except=(pos:x,aa:TERM)
CDS /note="TAA stop codon is completed by the addition of 3' A"
CDS /note="start codon not determined"