

Ferequency of stop codon/frameshift mutation (per codon)

0.02

0.01

0.00

Gag_MA

Gag_p30

RnaseH

rve

RVP

RVT_1

TLV_coat

domains

subfamily

FcERV_y6

MLERV1.1

MLERV1.2

MLERV1.3

