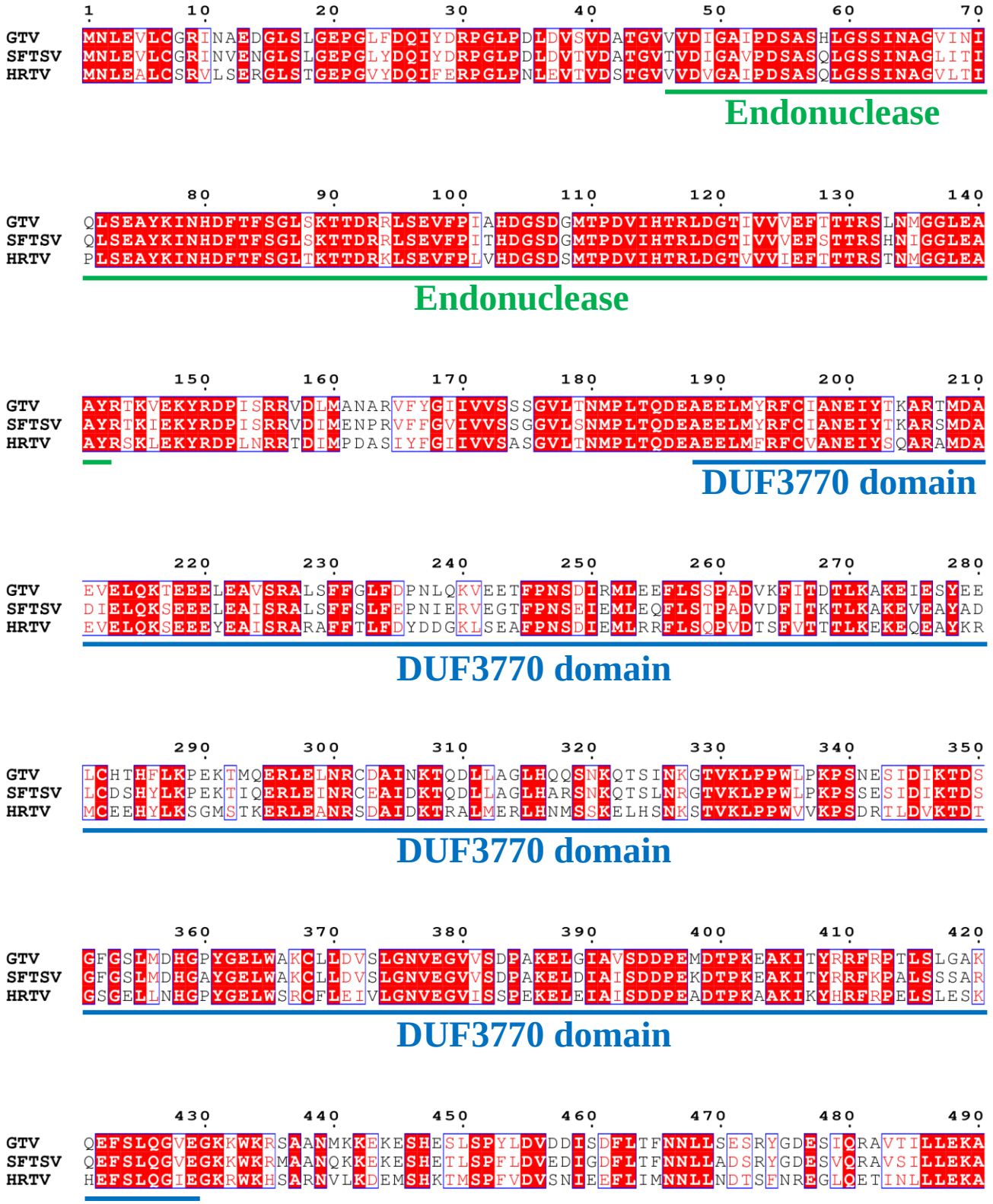
Figure S4 A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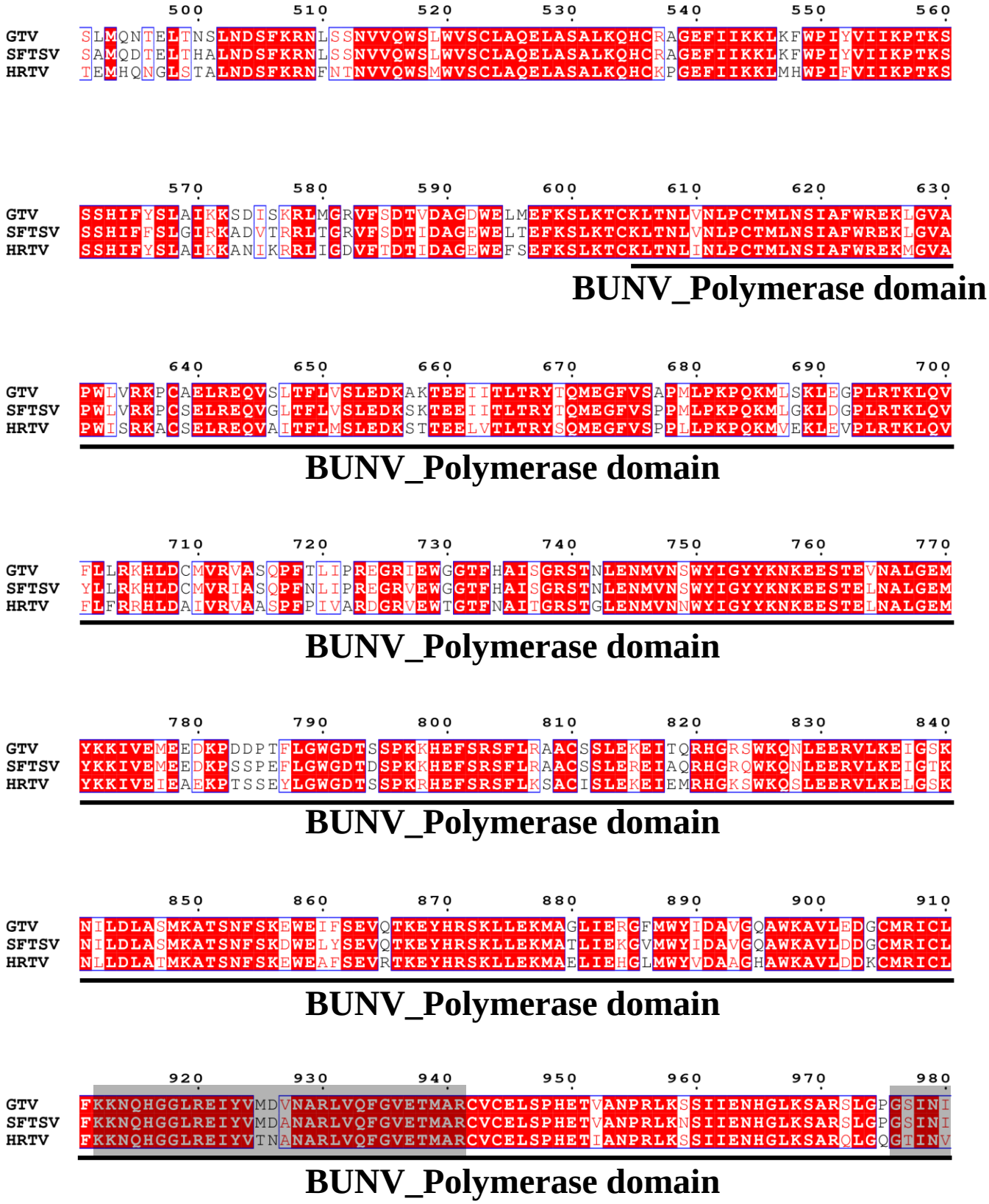
HRTV
Endonuclease
Endonuclease
DUF3770 domain
DUF3770 domain
DUF3770 domain
DUF3770 domain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SFTSV SAMQDTELTHALNDSFKRNLSSNVVQWSLWVSCLAQELASALKQHCRAGEFIIKKLKFWPIYVIIKPTKS
HRTV TEMHQNGLSTALNDSFKRNFNTNVVQWSMWVSCLAQELASALKQHCKPGEFIIKKLMHWPIFVIIKPTKS
570 580 590 600 610 620 630
GTV SSHIFYSLAIKKSDISKRLMGRVFSDTVDAGDWELMEFKSLKTCKLTNLVNLPCTMLNSIAFWREKLGVA
SFTSV SSHIFFSLGIRKADVTRRLTGRVFSDTIDAGEWELTEFKSLKTCKLTNLVNLPCTMLNSIAFWREKLGVA
HRTV SSHIFYSLAIKKANIKRRLIGDVFTDTIDAGEWEFSEFKSLKTCKLTNLINLPCTMLNSIAFWREKMGVA
BUNV_Polymerase domain
640 650 660 670 680 690 700
GTV PWLVRKPCAELREQVSLTFLVSLEDKAKTEEIITLTRYTQMEGFVSAPMLPKPQKMLSKLEGPLRTKLQV
SFTSV PWLVRKPCSELREQVGLTFLVSLEDKSKTEEIITLTRYTQMEGFVSPPMLPKPQKMLGKLDGPLRTKLQV
HRTV PWISRKACSELREQVAITFLMSLEDKSTTEELVTLTRYSQMEGFVSPPLLPKPQKMVEKLEVPLRTKLQV
BUNV_Polymerase domain
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SFTSV YLLRKHLDCMVRIASQPFNLIPREGRVEWGGTFHAISGRSTNLENMVNSWYIGYYKNKEESTELNALGEM
HRTV FLFRRHLDAIVRVAASPFPIVARDGRVEWTGTFNAITGRSTGLENMVNNWYIGYYKNKEESTELNALGEM
BUNV_Polymerase domain
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HRTV YKKIVEIEAEKPTSSEYLGWGDTSSPKRHEFSRSFLKSACISLEKEIEMRHGKSWKQSLEERVLKELGSK
BUNV_Polymerase domain
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920 930 940 950 960 970 980
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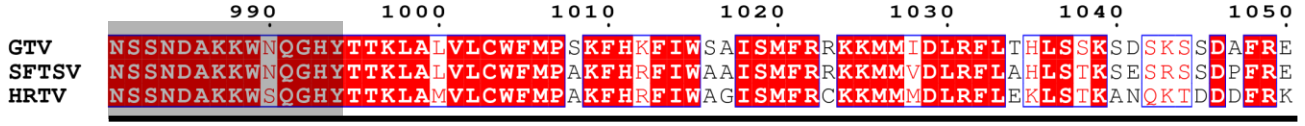

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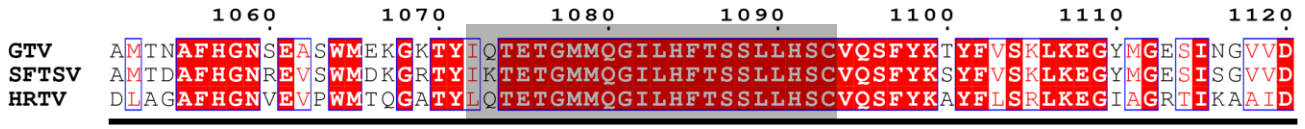

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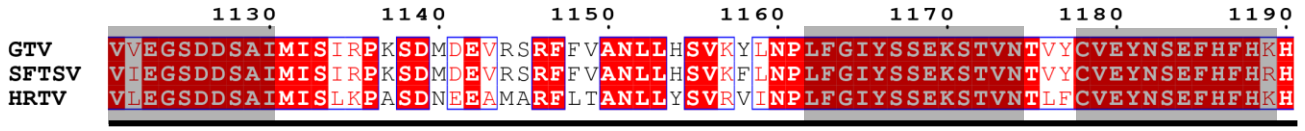

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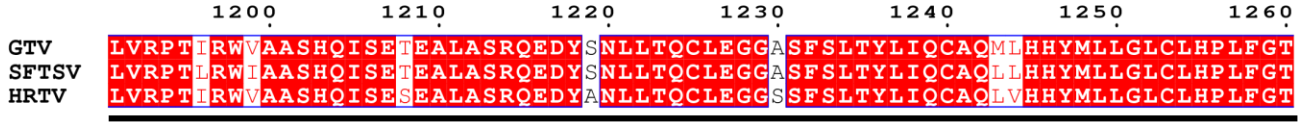

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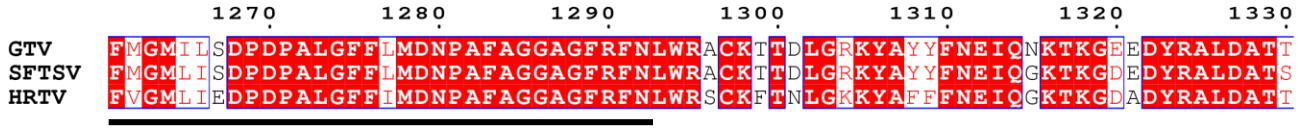

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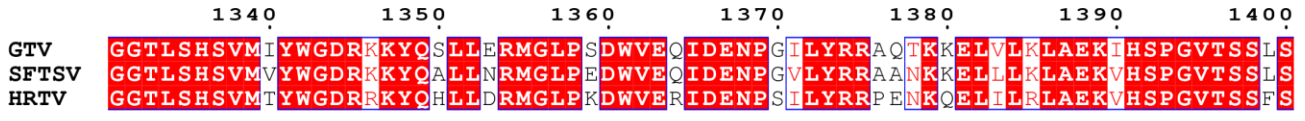

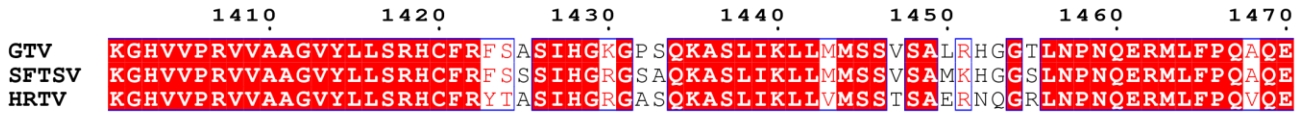

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SFTSV LEAEKSLDNQERLESISILKHVLFMVLNGPYTEEYKLEMIIEAFSTLVIPQPSEVIRKSRTMTLCLLSNY
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HRTV IEQRGGSILDQIEKAQSGTVGGFSKPQKPYRKQSGGIGYKGKGVWSGIMENTNVQILIDGDGSSNWIEEI
1760 1770 1780 1790 1800 1810 1820
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SFTSV RLSSDARLYDVIESIRRLCDDLGINNRVASAYRGHCMVRLSGFKIKPASRTDGCPVRIMERGFRIRELQN
HRTV RLSSESRLFDVIESVRRLCDDINVNNRVTSSFRGHCMVRLSNFKVKPASRVEGCPVRLMPSSFRIKELQN
1830 1840 1850 1860 1870 1880 1890
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1900 1910 1920 1930 1940 1950 1960
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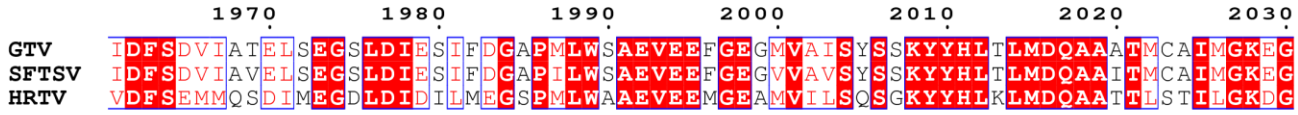
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1980
1990
2000
2010
2020
2030
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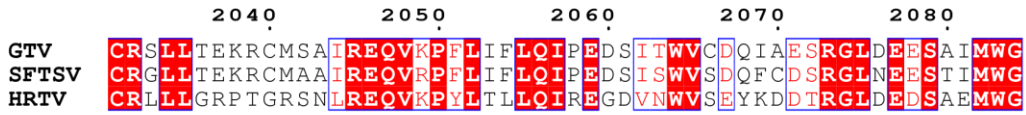
2040
2050
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2070
2080
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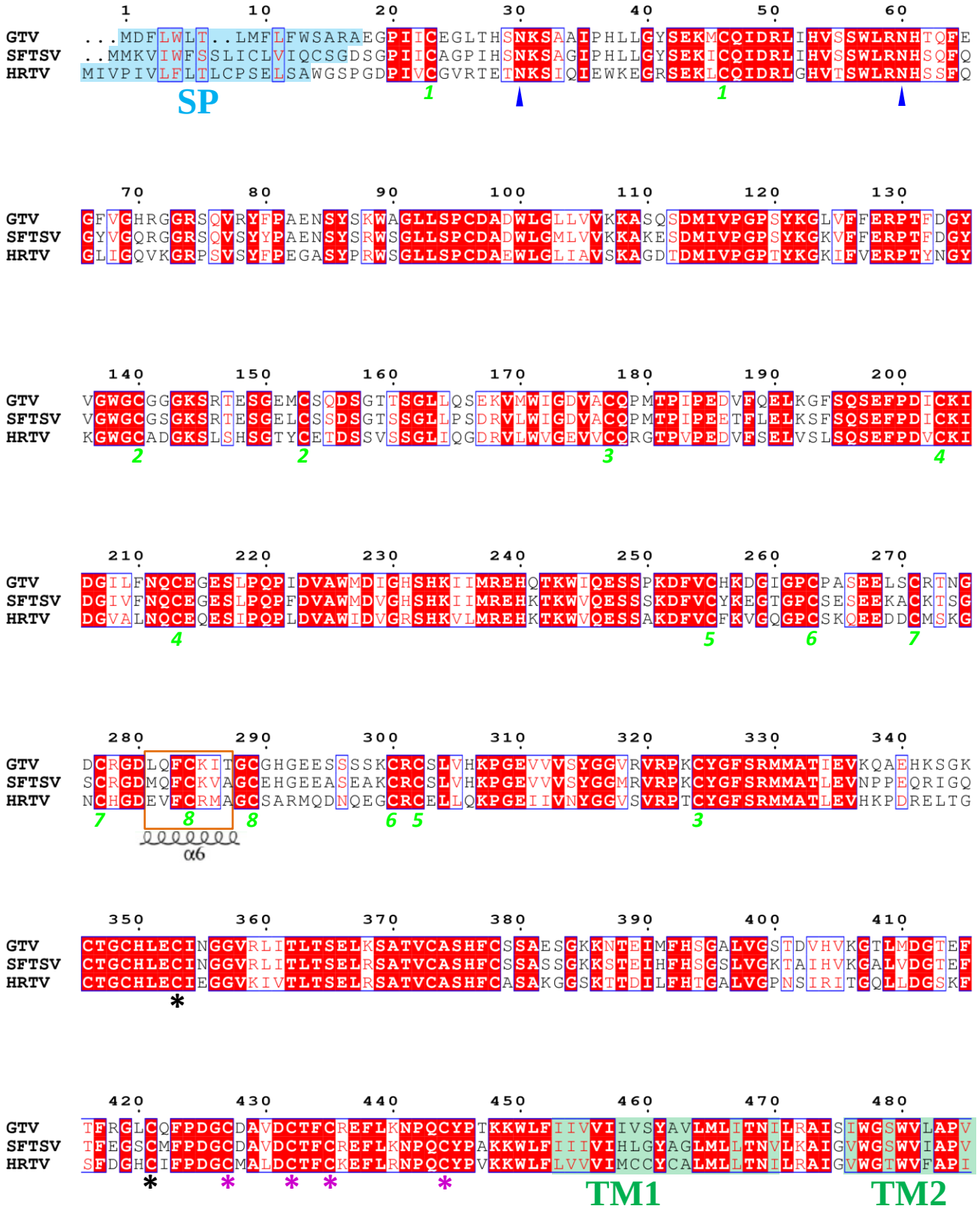
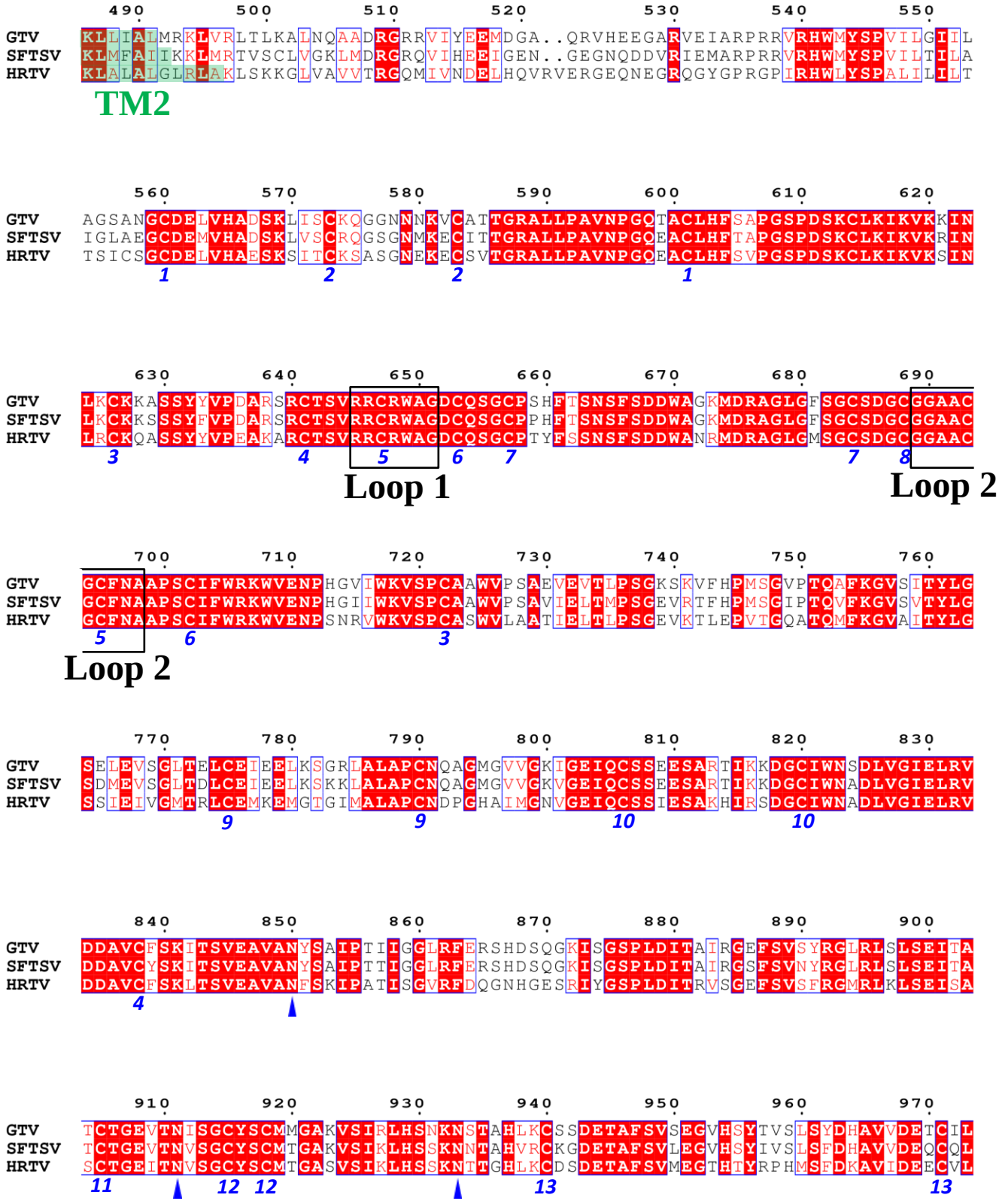
Figure S4 B

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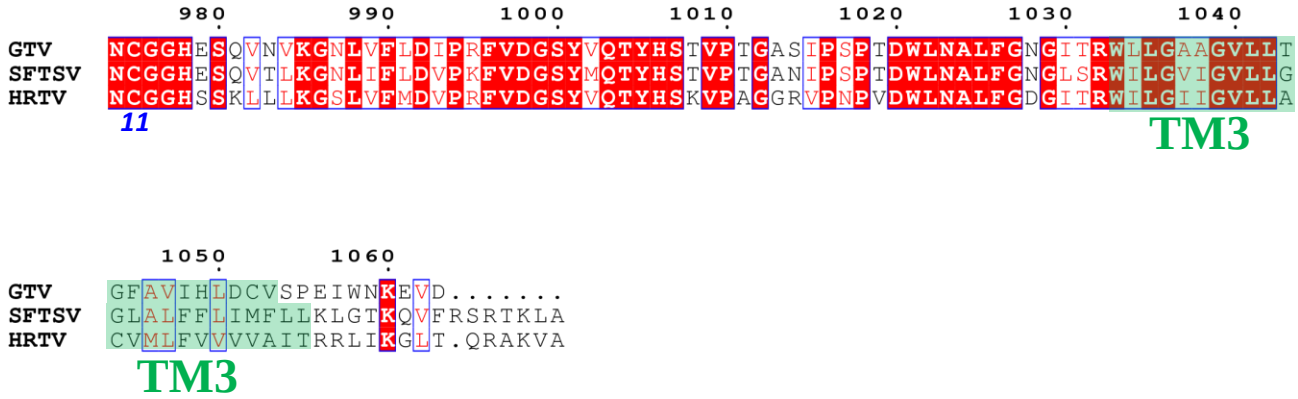
SFTSV
HRTV
TM2
Loop 1
Loop 2
Loop 2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1030
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TM3
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GTV
SFTSV
HRTV
TM3

Figure S4 C

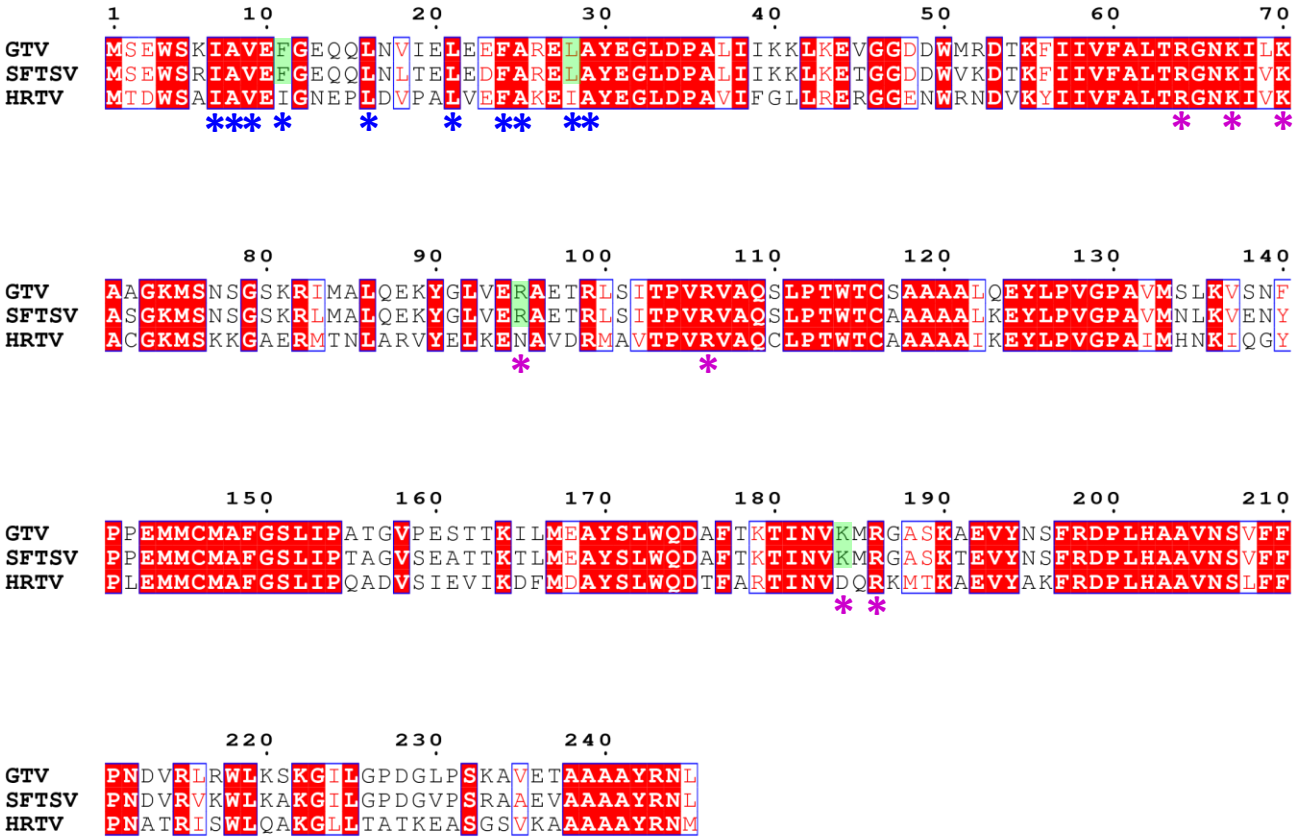
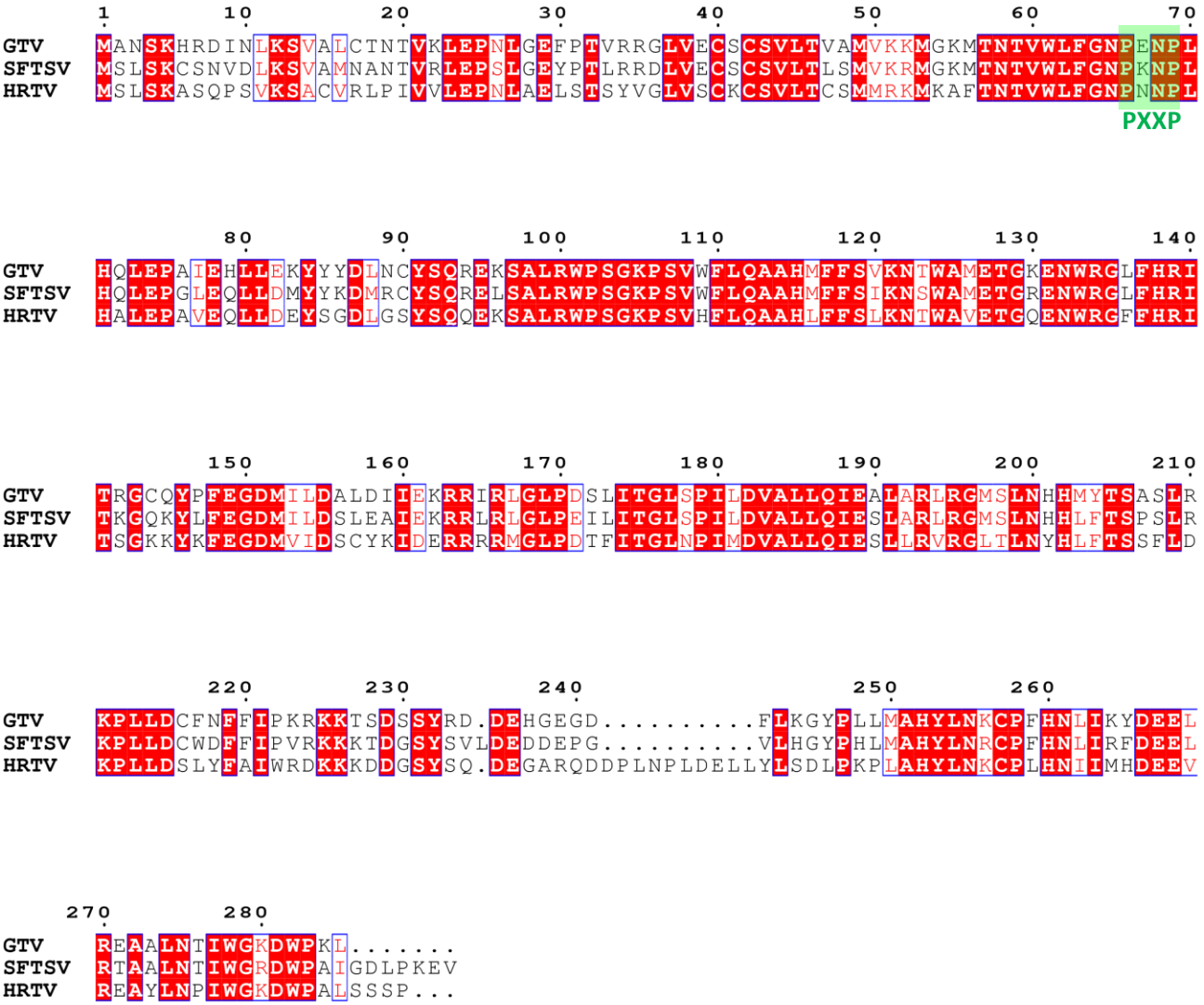
Figure S4 D