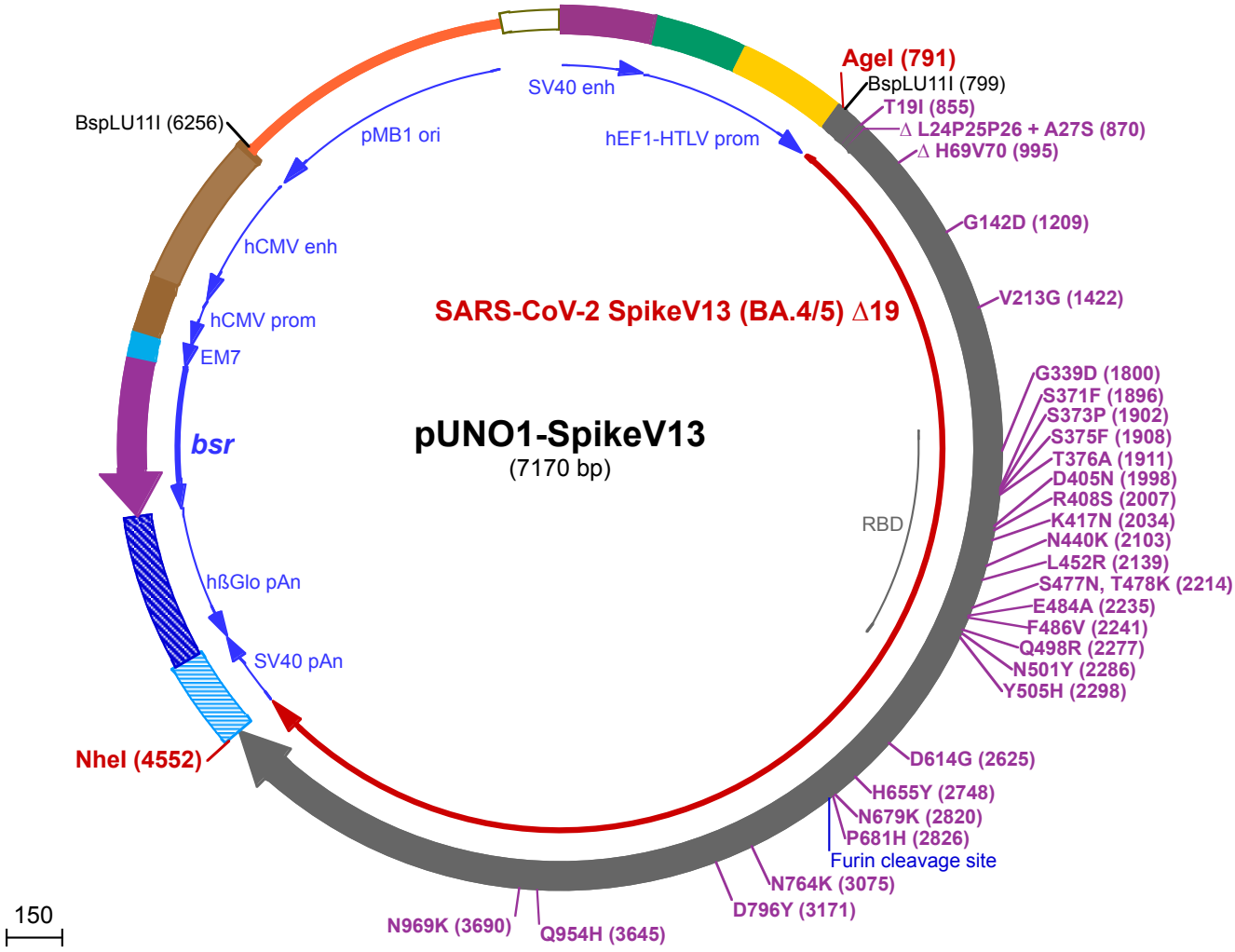
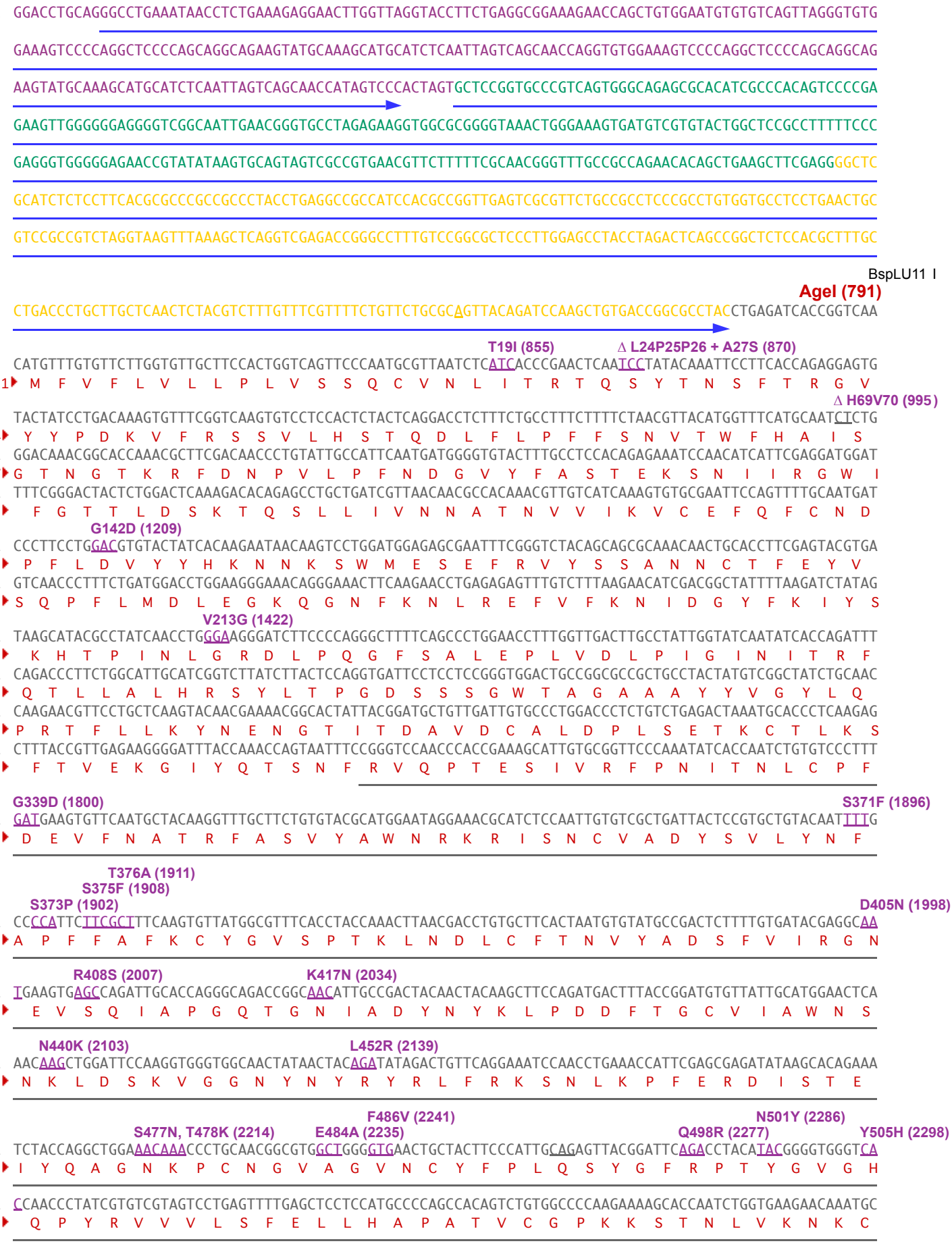

AgeI (791)
BspLU11I (799)
T19I (855)
Δ L24P25P26 + A27S (870)
Δ H69V70 (995)
G142D (1209)
V213G (1422)
G339D (1800)
S371F (1896)
S373P (1902)
S375F (1908)
T376A (1911)
D405N (1998)
R408S (2007)
K417N (2034)
N440K (2103)
L452R (2139)
S477N, T478K (2214)
E484A (2235)
F486V (2241)
Q498R (2277)
N501Y (2286)
Y505H (2298)
D614G (2625)
H655Y (2748)
N679K (2820)
P681H (2826)
Furin cleavage site
N764K (3075)
D796Y (3171)
Q954H (3645)
N969K (3690)
NheI (4552)
BspLU11I (6256)
SV40 enh
pMB1 ori
hEF1-HTLV prom
hCMV enh
hCMV prom
EM7
bsr
hßGlo pAn
SV40 pAn
SARS-CoV-2 SpikeV13 (BA.4/5) Δ19
pUNO1-SpikeV13
(7170 bp)
RBD
150


BspLU11 I
AgeI (791)
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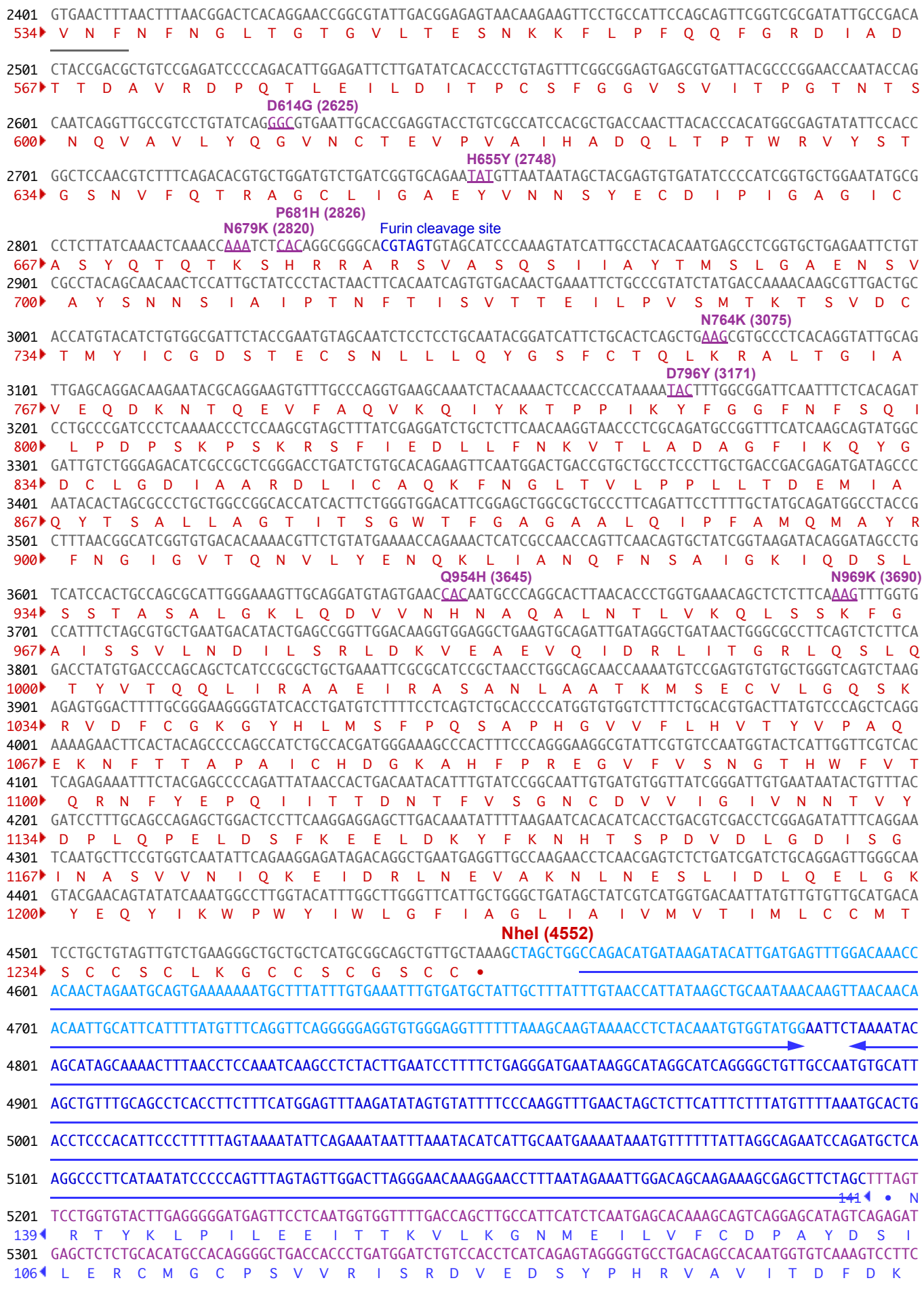
D614G (2625)
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Furin cleavage site
N764K (3075)
D796Y (3171)
Q954H (3645)
N969K (3690)
NheI (4552)


5401 TGCCCGTTGCTCACAGCAGACCCAATGGCAATGGCTTCAGCACAGACAGTGACCCTGCCAATGTAGGCCTCAATGTGGACAGCAGAGATGATCTCCCCAG
72 Q G N S V A S G I A I A E A C V T V R G I Y A E I H V A S I I E G T
5501 TCTTGGTCCTGATGGCCGCCCCGACATGGTGCTTGTTGTCCTCATAGAGCATGGTGATCTTCTCAGTGGCGACCTCCACCAGCTCCAGATCCTGCTGAGA
39 K T R I A A G V H H K N D E Y L M T I K E T A V E V L E L D Q Q S
5601 GATGTTGAAGGTCTTCATGATGGCCCTCCTATAGTGAGTCGTATTATACTATGCCGATATACTATGCCGATGATTAATTGTCAAAACAGCGTGGATGGCG
6 I N F T K M
5701 TCTCCAGCTTATCTGACGGTTCACTAAACGAGCTCTGCTTATATAGACCTCCCACCGTACACGCCTACCGCCCATTTGCGTCAATGGGGCGGAGTTGTTA
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