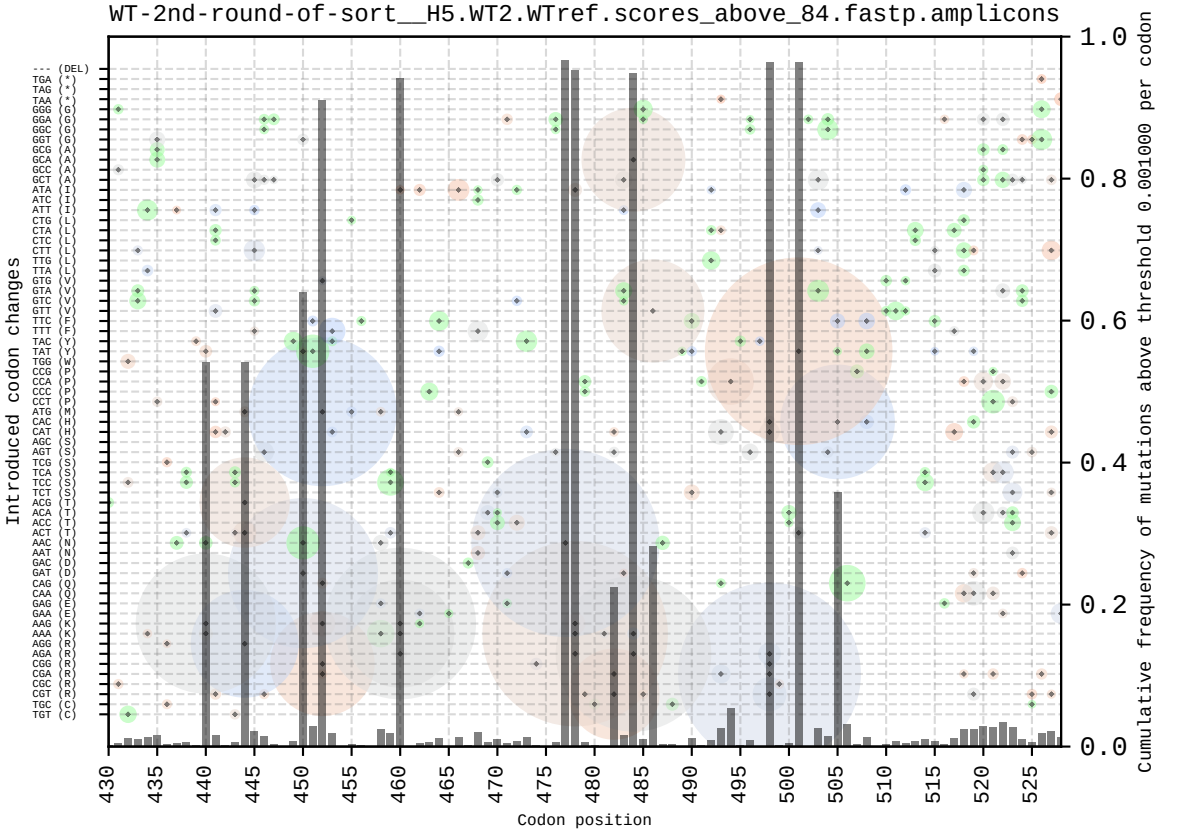
WT-2nd-round-of-sort__H5.WT2.WTref.scores_above_84.fastp.amplicons
Introduced codon changes
--- (DEL)
TGA (*)
TAG (*)
TAA (*)
GGG (G)
GGA (G)
GGC (G)
GGT (G)
GCG (A)
GCA (A)
GCC (A)
GCT (A)
ATA (I)
ATC (I)
ATT (I)
CTG (L)
CTA (L)
CTC (L)
CTT (L)
TTG (L)
TTA (L)
GTG (V)
GTA (V)
GTC (V)
GTT (V)
TTC (F)
TTT (F)
TAC (Y)
TAT (Y)
TGG (W)
CCG (P)
CCA (P)
CCC (P)
CCT (P)
ATG (M)
CAC (H)
CAT (H)
AGC (S)
AGT (S)
TCG (S)
TCA (S)
TCC (S)
TCT (S)
ACG (T)
ACA (T)
ACC (T)
ACT (T)
AAC (N)
AAT (N)
GAC (D)
GAT (D)
CAG (Q)
CAA (Q)
GAG (E)
GAA (E)
AAG (K)
AAA (K)
AGG (R)
AGA (R)
CGG (R)
CGA (R)
CGC (R)
CGT (R)
TGC (C)
TGT (C)
430
435
440
445
450
455
460
465
470
475
480
485
490
495
500
505
510
515
520
525
Codon position
0.0
0.2
0.4
0.6
0.8
1.0
Cumulative frequency of mutations above threshold 0.001000 per codon