

Supplemental Table 6. Types of mutations found in genomes of *pol2rc-ΔN* strains.

Type of substitution	Number of mutations found in all genomes		Number of independent mutations (taking into account common origin of sequenced genomes)	
	Number of mutations	%	Number of independent mutations	%
GC->CG	116	39.7	89	45.2
AT->TA	50	17.1	39	19.8
AT->CG	12	4.1	12	6.1
AT->GC	35	12.0	25	12.7
GC->AT	28	9.6	15	7.6
GC->TA	49	16.8	16	8.1
Complex (AAA->TAT)	2	0.7	1	0.5
Total	292	100	197	100