Groups ⁴	$r_g(\%)$
DRC_U (2)	100.0
A2 (1)	99.7
C (4)	99.5
B (4)	99.2
Н (3)	98.2
CRF02_AG (4)	97.9
DUG114 (1)	97.4

Table A5. Group ranking based on quartet scanning of HIV-1 group M including a divergent DRC lineage.

The number of sequences per group is indicated between brackets. The permutation *p*-value < 0.01 was based on d_{max} .