Table A7. Group ranking based on quartet scanning of HIV-1 group M including a CRF02\_AE.

Groups <sup>4</sup>	$r_g$ (%)
CRF02_AE (4)	100.0
CRF02_AG (4)	99.4
B (4)	94.9
C (4)	94.8
A2 (1)	92.0
DUG114 (1)	92.0
H (3)	89.9

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