

Supplemental Information for:

Vicriviroc Resistance Decay and Relative Replicative Fitness in HIV-1

Clinical Isolates Under Sequential Drug Selection Pressures

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Supplemental Figure S1. Protein alignment of native weeks 0, 28 and 48 full-length envelopes. The proportion of conserved amino acid identity, relative to p07Jenv0, is displayed above the sequences in bar format. Dots represent amino acids that are identical in all sequences; dashes signify length polymorphisms.

Supplemental Figure 1

