

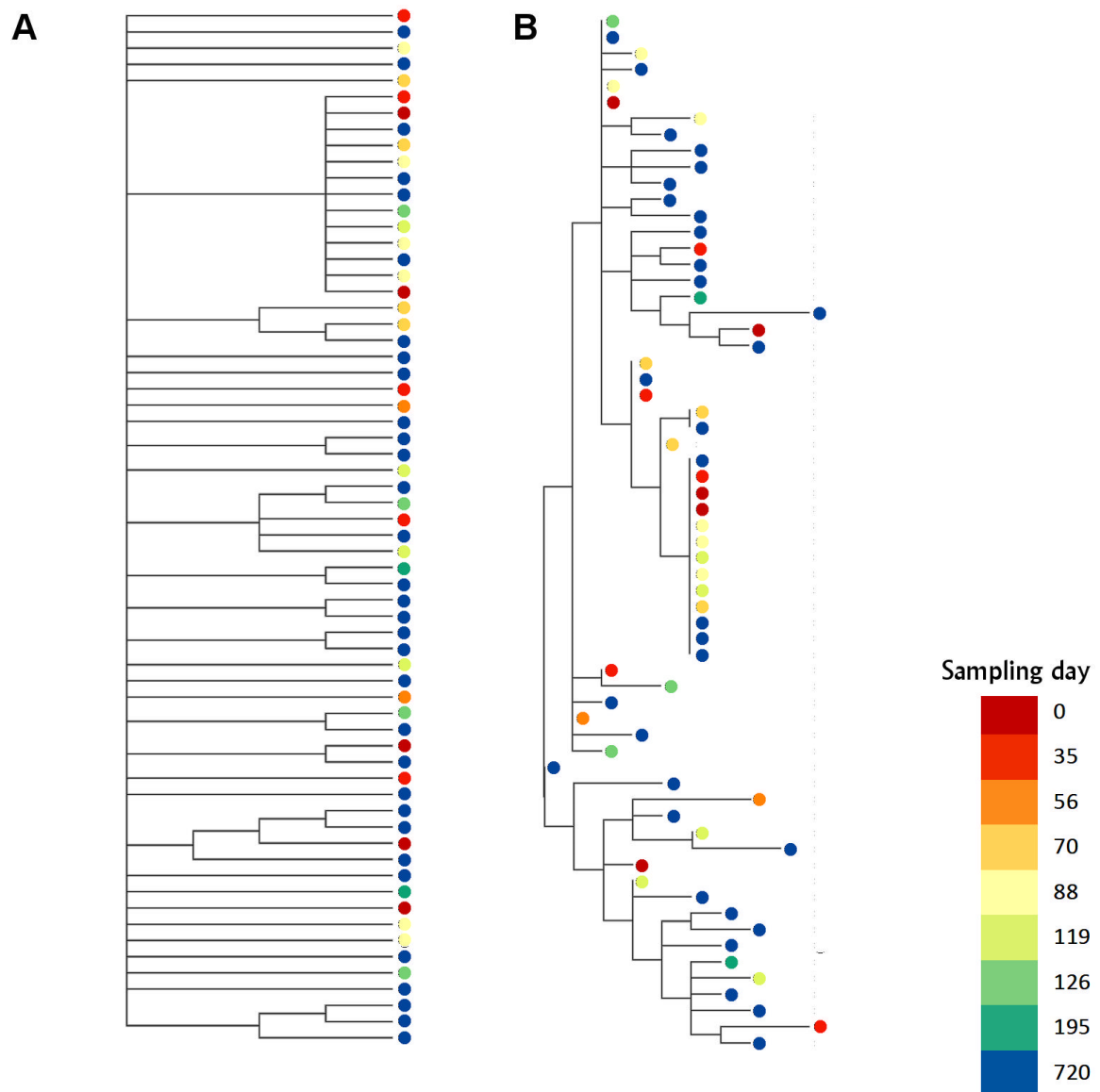
Supplementary material

No evidence of ongoing evolution in replication competent latent HIV-1 in a patient followed up for two years

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Supplementary figure 1

A. Maximum parsimony tree of *gag* sequences from reactivated patient viruses. The consensus tree inferred from 3 most parsimonious trees is shown. B. Maximum likelihood tree of *gag* sequences from reactivated patient viruses. Branch lengths show the number of substitutions per site.