



Fig. S1. Hepatitis C virus (HCV) NS5B phylogenetic tree comparing patients A and B with 73 HCV genotype 1a strains obtained from human immunodeficiency virus (HIV)-positive men who have sex with men (MSM). Robust MSM-specific clusters are depicted in grey. The insert shows sequential sequences of patients A and B over time. Symbols: HIV-positive MSM (\square), patient A (\star), patient B (\diamond).