

Supplementary Table 4. Summary of significant KEGG pathways (FDR ≤ 0.05) in the HIV/HCV-f versus HIV/HCV-b comparison.

KEGG pathways	Hits	Genes	q-values
Epstein-Barr virus infection	6	BID, CDKN1A, HLA-A, NFKB2, PDIA3, RELB	0.025
p53 signaling pathway	4	BID, CDKN1A, PMAIP1, TP73	0.025

Statistical: *q*-values, *p*-values corrected for multiple testing using the false discovery rate (*FDR*) with Benjamini and Hochberg procedure. In red, up-regulated genes in HIV/HCV-f group; in green, down-regulated genes in HIV/HCV-f group.

Abbreviations: FDR, false discovery rate for multiple comparisons using Benjamini and Hochberg procedure; KEGG, Kyoto Encyclopedia of Genes and Genomes; HIV, human immunodeficiency virus; HCV, hepatitis C virus; HIV/HCV-b, HIV/HCV-coinfected patients at baseline; HIV/HCV-f, HIV/HCV-coinfected patients 24 weeks after SVR.