

Table S1. Median percentages of T-cell phenotypes, HIV-negative women

Median (IQR)^a	HCV-	HCV+
CD4+ naive	28.9 (18.6-37.2)	15.9 (12.7-26.3)
CD4+ CM	19.4 (15.1-23.4)	16.8 (12.1-21.8)
CD4+ EM	38.3 (30.1-47.5)	50.1 (45.7-63.4)
CD4+ TE	12.0 (8.0-16.9)	7.9 (6.3-13.3)
CD4+ activated ^b	0.23 (0.08-0.37)	0.28 (-0.03-0.33)
CD4+ Treg	4.3 (3.4-5.5)	4.6 (3.4-5.5)
CD8+ naive	28.8 (19.7-40.2)	21.3 (14.6-24.9)
CD8+ CM ^b	0.44 (0.21-0.72)	0.34 (0.004-0.61)
CD8+ EM	27.4 (19.5-38.4)	38.5 (20.0-45.8)
CD8+ TE	34.8 (25.9-44.7)	36.0 (23.2-46.4)
CD8+ activated ^b	0.48 (0.25-0.74)	0.61 (0.20-0.79)

IQR: interquartile range

^aT cell phenotype abbreviations: effector memory (EM), central memory (CM) and terminally differentiated effector (TE)

^b Log(10) transformed phenotype

Table S2. Median percentages of T-cell phenotypes, HIV-positive women

Median (IQR)^a	HCV-	HCV+
CD4+ naive	30.6 (19.1-41.9)	26.6 (17.6-32.7)
CD4+ CM	15.8 (12.5-19.4)	14.9 (11.6-19.8)
CD4+ EM	40.2 (29.5-51.5)	45.6 (37.3-55.1)
CD4+ TE	10.6 (7.0-15.3)	9.3 (6.2-12.9)
CD4+ activated ^b	0.48 (0.31-0.68)	0.56 (0.38-0.72)
CD4+ Treg	4.9 (3.4-6.2)	6.0 (4.4-8.8)
CD8+ naive	17.6 (10.9-28.7)	17.1 (11.1-26.1)
CD8+ CM ^b	0.41 (0.20-0.65)	0.45 (0.25-0.68)
CD8+ EM	37.5 (27.4-48.7)	40.4 (30.2-47.3)
CD8+ TE	35.0 (26.9-45.8)	35.9 (26.0-45.1)
CD8+ activated ^b	0.83 (0.55-1.13)	0.87 (0.60-1.20)

IQR: interquartile range

^aT cell phenotype abbreviations: effector memory (EM), central memory (CM) and terminally differentiated effector (TE)

^b Log(10) transformed phenotype