## Figure S7. Statistical testing of TCR data from CPI colitis and controls.

A. Comparisons of the number of expanded clusters (node size >1), normalized to the total

number of cells per sample, across disease states in biopsies (top) and blood (bottom).

						Overal	colitis	colitis	hc vs.
Biopsies	level	Overall	cpi_colitis	hc	uc	۱p	vs uc	vs hc	uc
n		14	9	3	2				
	1	55.00	87.00	27.00	34.00				
ncluster (m	nedian	[33.25,	[62.00,	[25.00,	[29.50,				
[IQR])		99.75]	106.00]	28.50]	38.50]	0.011	0.033	0.012	0.564
normalized	l.ncluster	0.08 [0.06,	0.09 [0.08,	0.05 [0.04,	0.06 [0.06,				
(median [I0	QR])	0.09]	0.11]	0.05]	0.06]	0.009	0.034	0.013	0.083

						Overal	colitis	colitis	hc vs.
Blood	level	Overall	cpi_colitis	hc	uc	۱p	vs uc	vs hc	uc
N		14	8	3	3				
	•	13.00	13.00	10.00	27.00	13.00			
ncluster (m	ledian	[4.00,	[5.75,	[6.50,	[15.00,	[4.00,			
[IQR])		22.50]	18.75]	26.00]	36.50]	22.50]	0.22	0.838	0.658
		0.02	0.03			0.02			
normalized	.ncluster	[0.01,	[0.01,	0.02 [0.01,	0.02 [0.01,	[0.01,			
(median [IC	QR])	0.03]	0.03]	0.03]	0.03]	0.03]	0.838	0.683	0.827

**B.** Gini coefficient for all T cells per sample in blood and biopsies across disease states.

						Overall	colitis	colitis	hc vs.
	level	Overall	cpi_colitis	hc	uc	р	vs uc	vs hc	uc
N		14	8	3	3				
Gini coeff	icient in								
blood (me	edian	0.03 [0.01,	0.03 [0.02,	0.01 [0.01,	0.12 [0.06,				
[IQR])		0.09]	0.05]	0.05]	0.12]	0.544	0.683	0.414	0.544

14	9	3	2				
0.17 [0.12,	0.18 [0.17,	0.07 [0.07,	0.14 [0.13,				
0.24]	0.34]	0.08]	0.14]	0.016	0.099	0.013	0.016
	0.17 [0.12,	0.17 [0.12, 0.18 [0.17,	0.17 [0.12, 0.18 [0.17, 0.07 [0.07,	0.17 [0.12, 0.18 [0.17, 0.07 [0.07, 0.14 [0.13,	0.17 [0.12, 0.18 [0.17, 0.07 [0.07, 0.14 [0.13,	0.17 [0.12, 0.18 [0.17, 0.07 [0.07, 0.14 [0.13,	0.17 [0.12, 0.18 [0.17, 0.07 [0.07, 0.14 [0.13,

**C.** Descriptive statistics of the number of individual expanded TCR $\alpha\beta$  clusters shared by

combinations of CD4T RM, CD8T RM, and CD8T GZMB cells in blood and biopsies.

	Both blood and biopsies	Blood	Biopsies
# of Expanded clusters	1248	230	1035
(cluster size >1)			
# of clusters shared by	66	0	66 (6.4%)
CD4T RM, CD8T GZMB,			
CD8T RM			
# of Shared between CD4T	81	0	81 (7.8%)
RM, CD8T GZMB			
# of Shared between CD4T	155	0	155 (15.0%)
RM, CD8T RM			
# of Shared between CD8T	203	0	202 (19.5%)
GZMB, CD8T RM			

D. Descriptive statistics of the number of individual expanded clusters shared by combinations

of CD4T RM, CD8T RM, and CD8T GZMB cells in biopsies across disease states.

Biopsies	cpi_colitis	uc	hc
# of Expanded clusters	887	68	80
(cluster size >1)			
# of clusters shared by	64 (7.2%)	2 (2.9%)	0
CD4T RM, CD8T GZMB,			
CD8T RM			
# of Shared between CD4T	78 (8.8%)	2 (2.9%)	1 (1.3%)

RM, CD8T GZMB			
# of Shared between CD4T RM, CD8T RM	146 (16.4%)	7 (10.3%)	2 (2.5%)