

Supplemental Table 2. Gene Expression: Comparing Nanostring to RT-PCR¹

Genes	CR Flares		NR Flares	
	Nanostring	RT-PCR ²	Nanostring	RT-PCR
MX1	8.63	4.16	3.77	3.45
STAT1	4.42	4.66	2.89	4.67
CCL19	3.52	6.59	2.82	8.34
MME	0.4	0.59	0.15	-0.01
ITGB2	1.67	3.99	3.45	2.09

¹All expression values are reported as fold-change compared to normal controls.

OAZ1 was used as the internal control for RT-PCR.

²RT-PCR values were obtained using threshold cycle quantification method (C_T) and reported as $\Delta\Delta C_T$