## Table S3

Number of reads and hybrids identified within biological replicates and individual infection groups.

Lytic							
	Replicate 1	Replicate 2	Replicate 3	Total			
Total Number of Reads	33,155,072	18,657,088	24,443,598	76,255,758			
Total Number of Hybrids	75,222	145,353	133,142	353,717			
Total miRNA Hybrids	31,277	67,120	57,984	156,381			
Total Host miRNA Hybrids	26,946	59,127	50,501	136,574			
Total Viral miRNA Hybrids	4,331	7,993	7,483	19,807			

Reactivation							
	Replicate 1	Replicate 2	Replicate 3	Total			
Total Number of Reads	36,040,560	17,794,998	20,190,940	74,026,498			
Total Number of Hybrids	129,172	83,977	113,026	326,175			
Total miRNA Hybrids	43,425	25,936	39,812	109,173			
Total Host miRNA Hybrids	36,476	21,801	33,980	92,257			
Total Viral miRNA Hybrids	6,949	4,135	5,832	16,916			

Latency							
	Replicate 1	Replicate 2	Replicate 3	Total			
Total Number of Reads	9,038,009	6,467,044	8,322,503	23,827,556			
Total Number of Hybrids	20,906	18,403	24,934	64,243			
Total miRNA Hybrids	8,146	9,100	11,945	29,191			
Total Host miRNA Hybrids	7,302	8,108	10,629	26,039			
Total Viral miRNA Hybrids	844	992	1,316	3,152			