Limitation	Chemostat replicate	Treatment	Mean Density (± SEM)			Stability	
			Synechococcus	SRIM-	-8 phage	Synechococcus	SRIM-8 Phage
NL		Coevolved	1.69E+07 (± 1.306E+07) 2.52E+08	(± 1.390E+08)	0.75	1.05
PL		Coevolved	1.11E+07 (± 1.089E+07) 2.39E+08	(± 9.488E+07)	0.59	1.45
NL		Control	1.31E+07 (± 3.694E+06			2.05	
PL	•	Control	1.78E+07 (± 9.016E+06			1.14	•
NL	NL2	Coevolved	1.76E+07 (± 1.207E+07) 1.52E+08	(± 7.919E+07)	0.84	1.11
NL	NL3	Coevolved	1.61E+07 (± 1.004E+07) 2.92E+08	(± 1.458E+08)	0.93	1.16
NL	NL5	Coevolved	1.70E+07 (± 1.648E+07) 3.13E+08	(± 1.609E+08)	0.59	1.12
PL	PL2	Coevolved	1.29E+07 (± 1.428E+07) 1.61E+08	(± 5.907E+07)	0.52	1.58
PL	PL4	Coevolved	1.08E+07 (± 6.656E+06) 2.43E+08	(± 8.836E+07)	0.94	1.59
PL	PL5	Coevolved	9.40E+06 (± 1.038E+07) 3.13E+08	(± 1.105E+08)	0.52	1.63
NL	NL1	Control	1.31E+07 (± 3.694E+06			2.05	
PL	PL1	Control	2.54E+07 (± 1.026E+07			1.43	
PL	PL3	Control	9.86E+06 (± 4.016E+06			1.42	