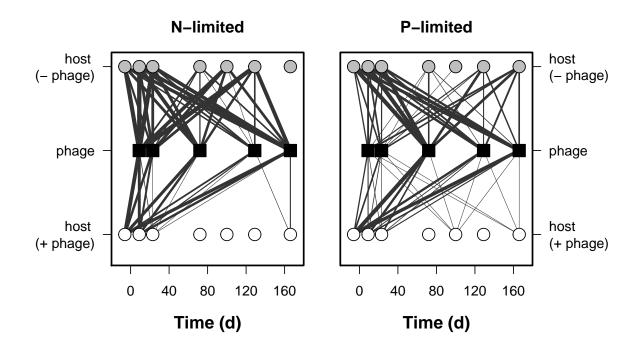
Evolutionary Dynamics

1 Data Reduction and Figures

1.1 Evolutionary Networks

1.1.1 Figure 2: Co-evolutionary dynamics



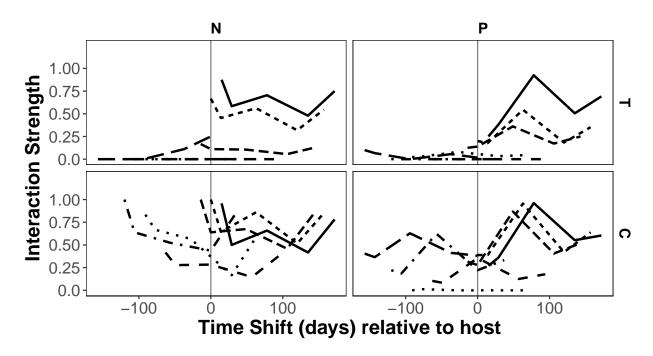
1.1.2 RMANOVA for Interaction Strengths

	AIC	BIC	logLik	L.Ratio	p-value
model.ar	169.1	260.1	-65.57	NA	NA
${f model.arma1}$	115.5	211.3	-37.76	55.63	8.769e-14
${f model.arma2}$	98.61	199.2	-28.31	18.91	1.37e-05

	Std .	t-value	p-value
(Intercept)	0.07713	9.872	7.099e-22
trtT	0.08638	-3.702	0.0002276
limP	0.09298	-4.93	0.007875
phage.time	0.0006172	-1.916	0.05569
bac.time	0.001005	0.1231	0.9021
trtT:limP	0.1078	1.856	0.06381
trtT:phage.time	0.0007027	1.008	0.3136
limP:phage.time	0.0007399	3.511	0.0004695
trtT:bac.time	0.001097	-3.091	0.00206
limP:bac.time	0.001136	-0.02436	0.9806
phage.time:bac.time	8.462 e-06	-1.811	0.07056
trtT:limP:phage.time	0.0008774	-0.9831	0.3258
trtT:limP:bac.time	0.001294	1.933	0.05351
trtT:phage.time:bac.time	$9.205\mathrm{e}\text{-}06$	2.014	0.04427
limP:phage.time:bac.time	9.576 e - 06	0.4493	0.6533
trtT:limP:phage.time:bac.time	1.086e-05	-1.801	0.07197

1.2 Evolutionary Time-shifts

1.2.1 Figure 3 | Figure S



1.3 Time-shift statistics

	AIC	BIC	logLik	L.Ratio	p-value
${f model.ar}$	169.1	260.1	-65.57	NA	NA
${f model.arma1}$	115.5	211.3	-37.76	55.63	8.769e-14
${f model.arma2}$	98.61	199.2	-28.31	18.91	1.37e-05

	Std.Error	t-value	p-value
(Intercept)	0.03853	16.31	1.635 e-52
\mathbf{trtT}	0.04353	-10.05	0.0001667
limP	0.04541	-6.084	0.001735
time.shift	0.0003838	-2.211	0.02726
trtT:limP	0.05345	4.45	0.006701
${ m trtT:} { m time.shift}$	0.0004238	4.997	$7.002\mathrm{e}\text{-}07$
limP:time.shift	0.0004391	3.912	$9.853\mathrm{e}\text{-}05$
${ m trtT:} { m limP:} { m time.shift}$	0.0005044	-3.341	0.0008681