

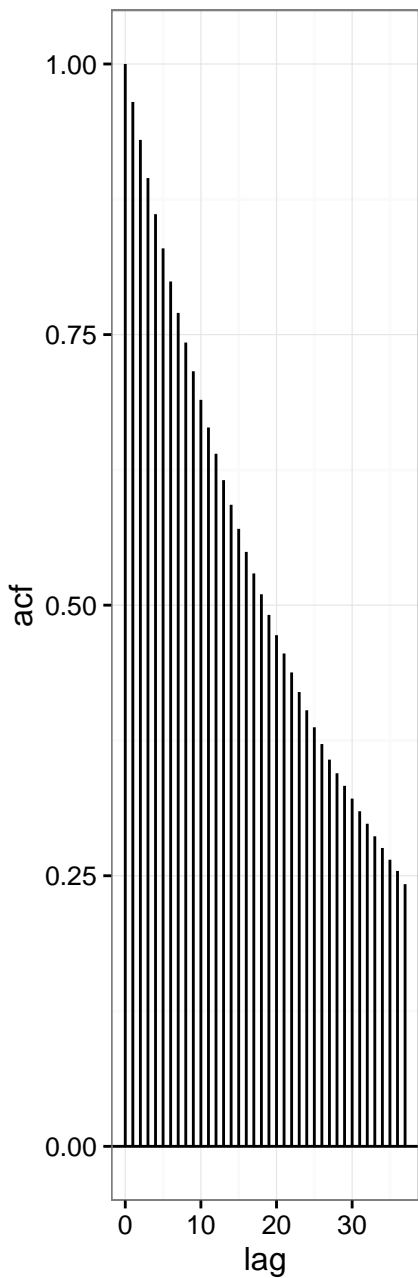
I ppl: 100
Unbalanced Replicates

K regions: 35
No mcmc: 1e+05
BurnIn: 20000
thinning: 15

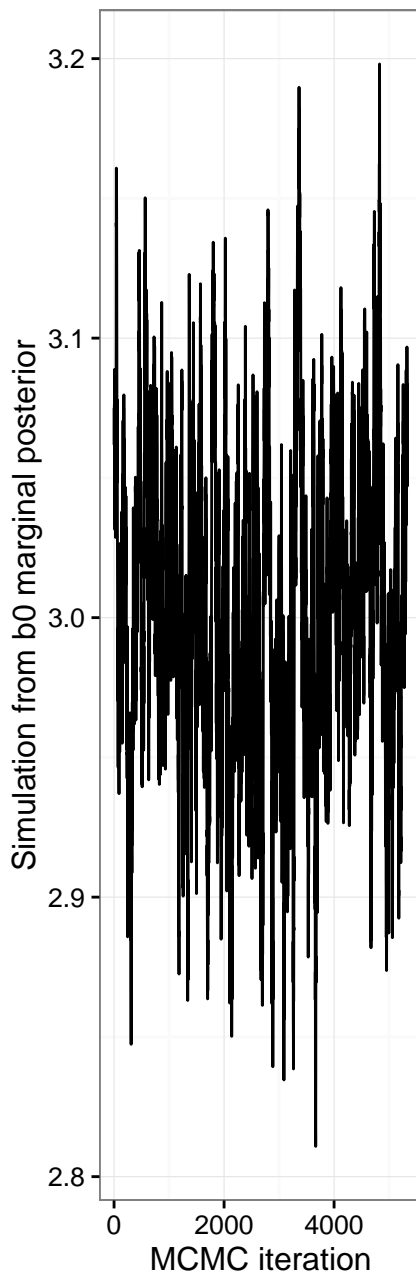
Length retained chain: 5333
Processed results from G1.pdf

	b0 estimate	
w.95.ci:2.8868	post.mean:3.0018	high.95.ci:3.1167
	b1 estimate	
w.95.ci:0.2814	post.mean:0.4673	high.95.ci:0.6203
	s2 estimate:	
w.95.ci:0.4951	post.mean:0.4992	high.95.ci:0.5031
	s2s estimate:	
w.95.ci:0.6445	post.mean:0.6929	high.95.ci:0.7453

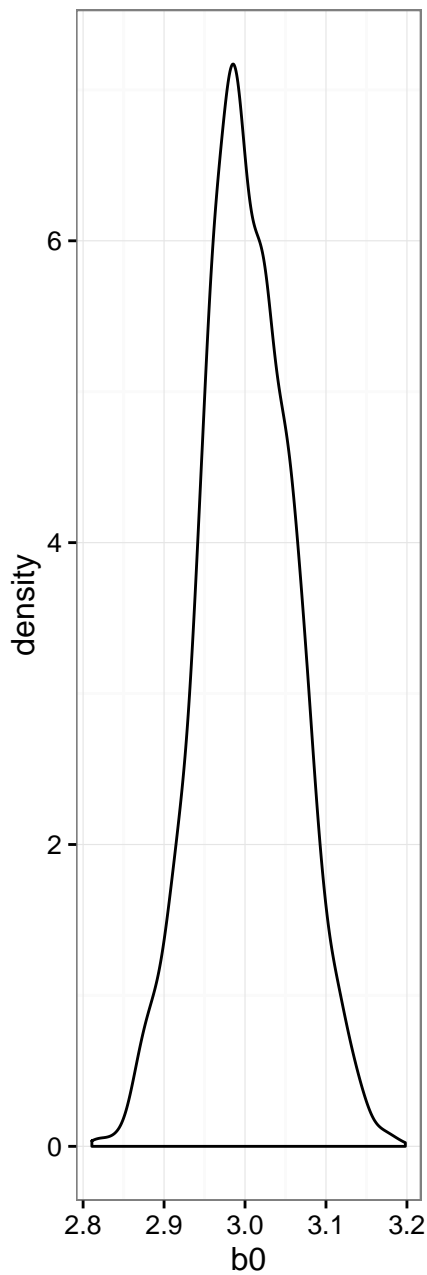
Autocorr for b0



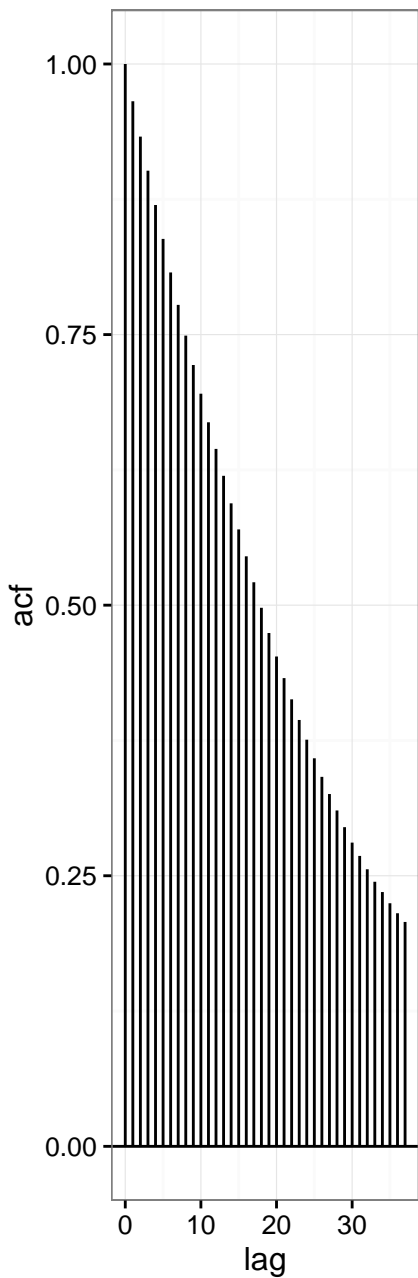
Trace plot for b0



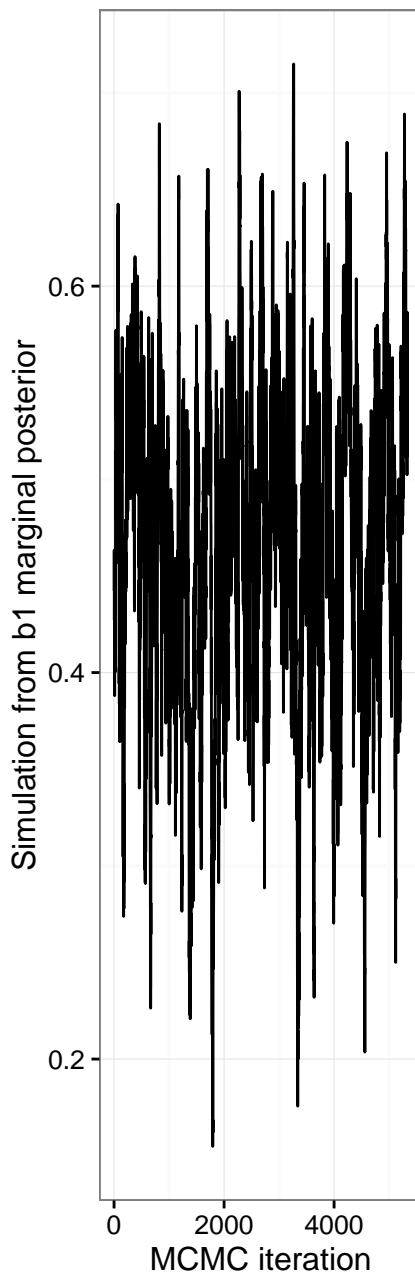
Density for b0 marginal



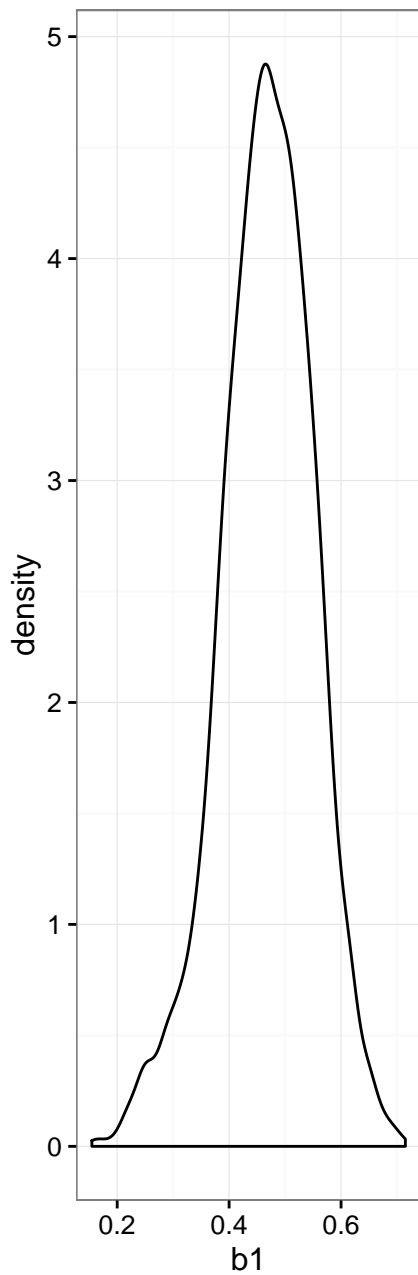
Autocorr for b1



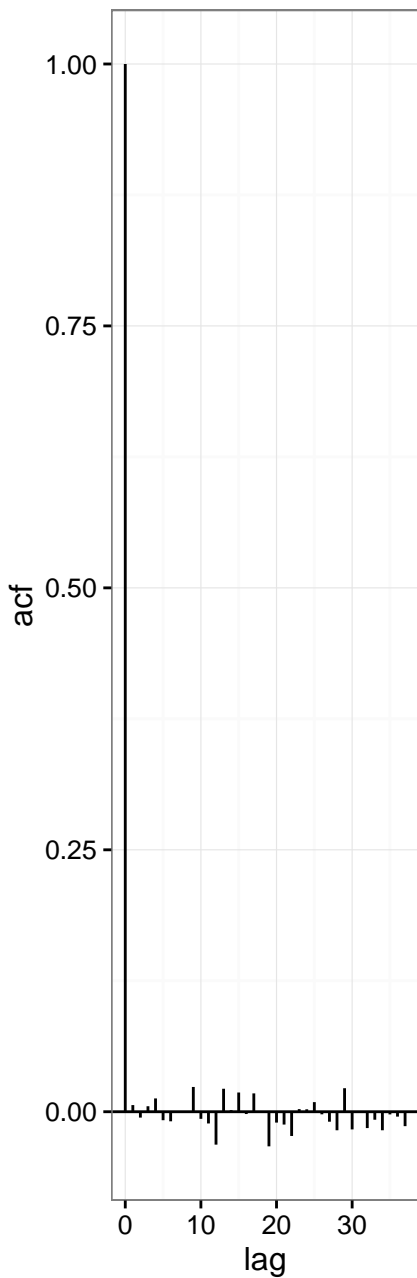
Trace plot for b1



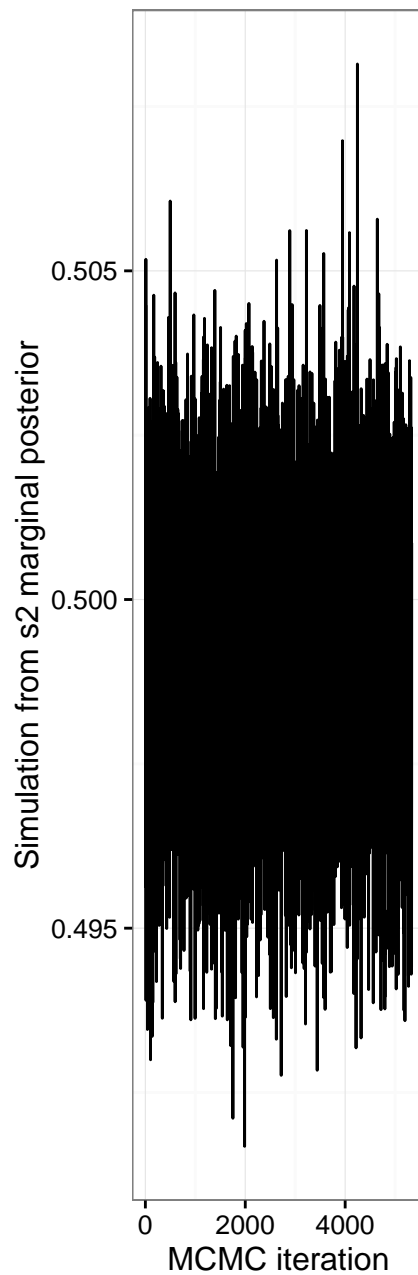
Density for b1 marginal



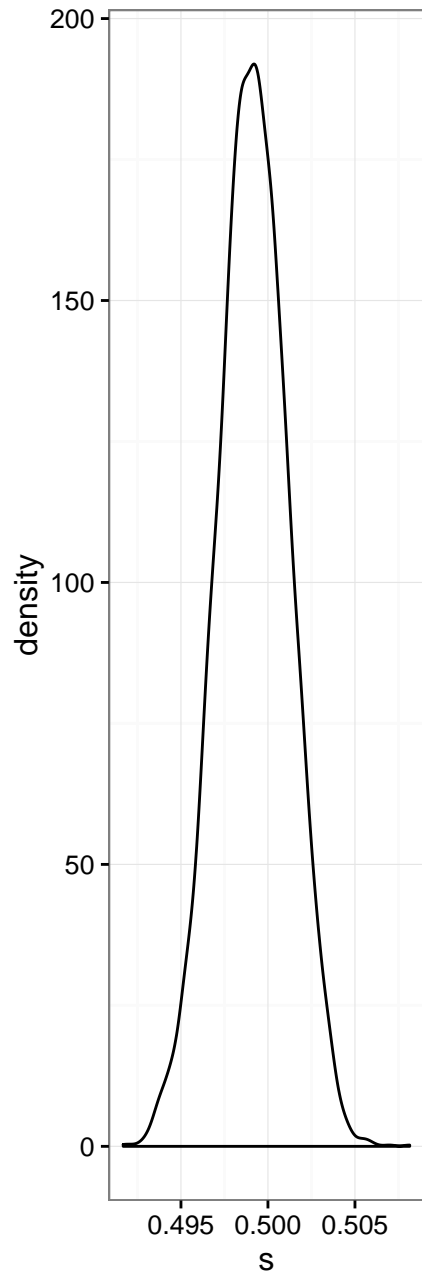
Autocorr for s



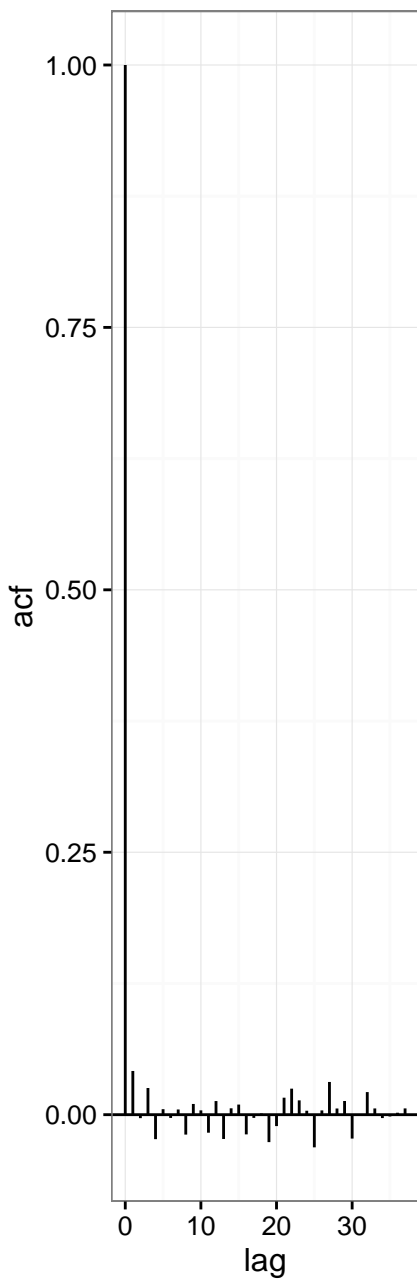
Trace plot for s2



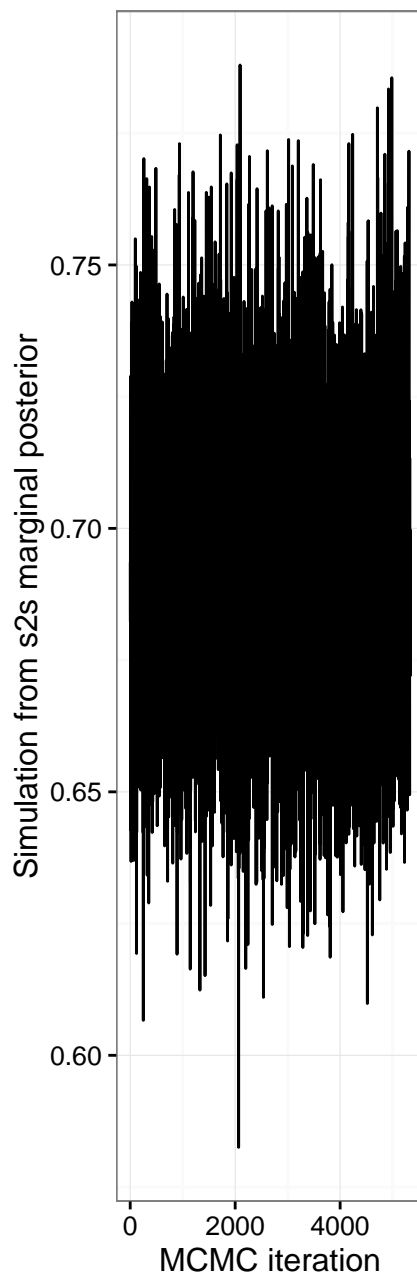
Density for s2 marginal



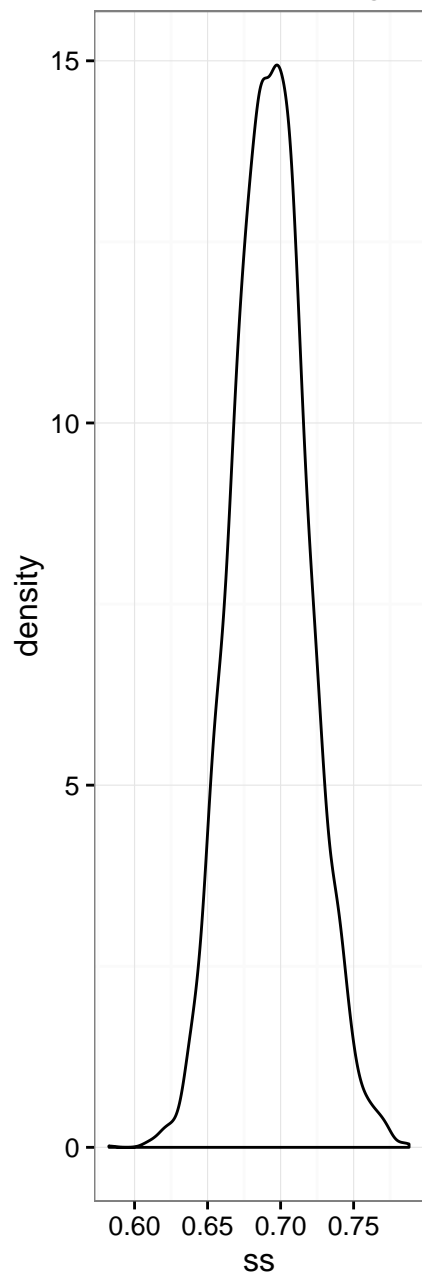
Autocorr for s2s



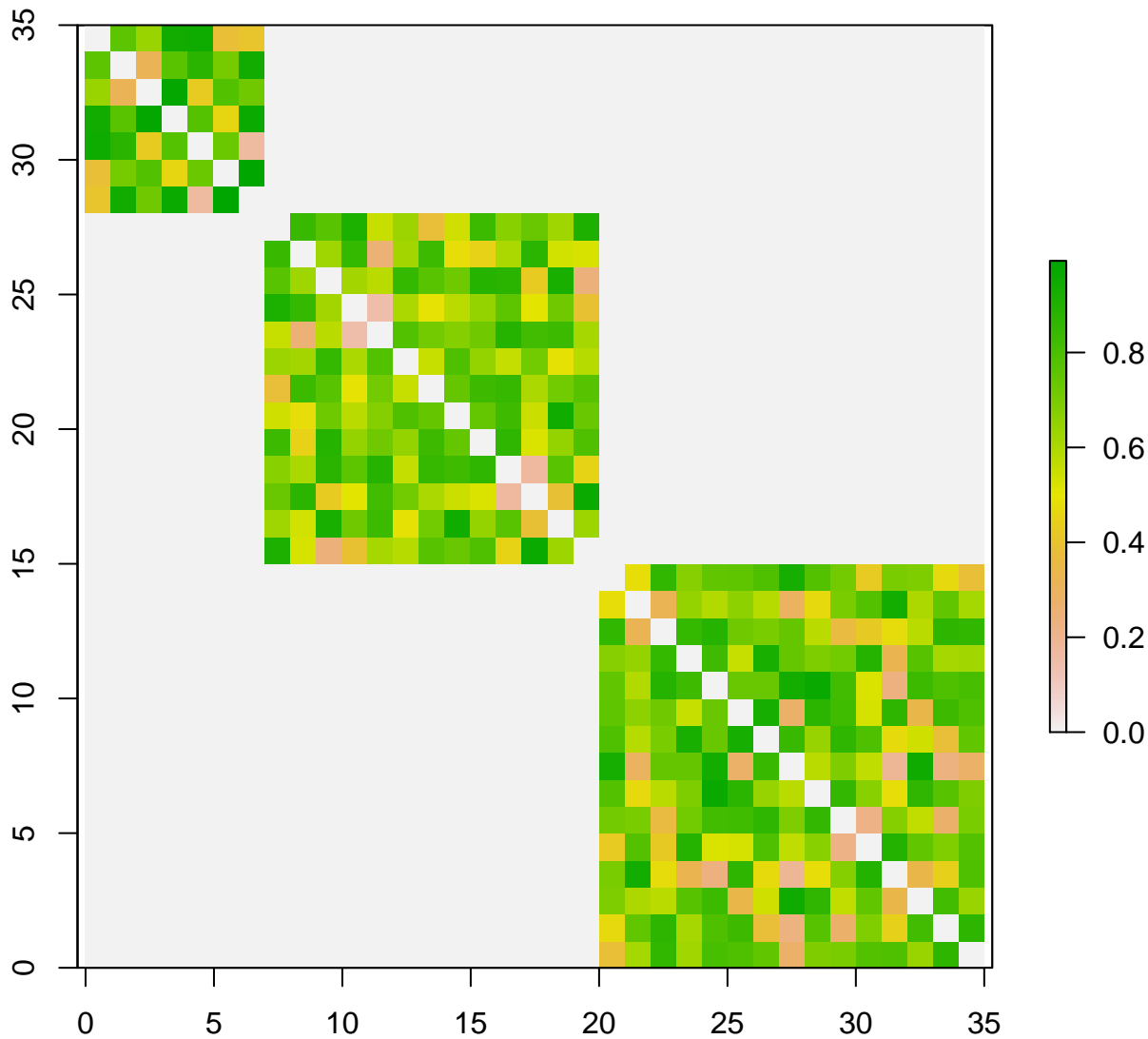
Trace plot for s2s



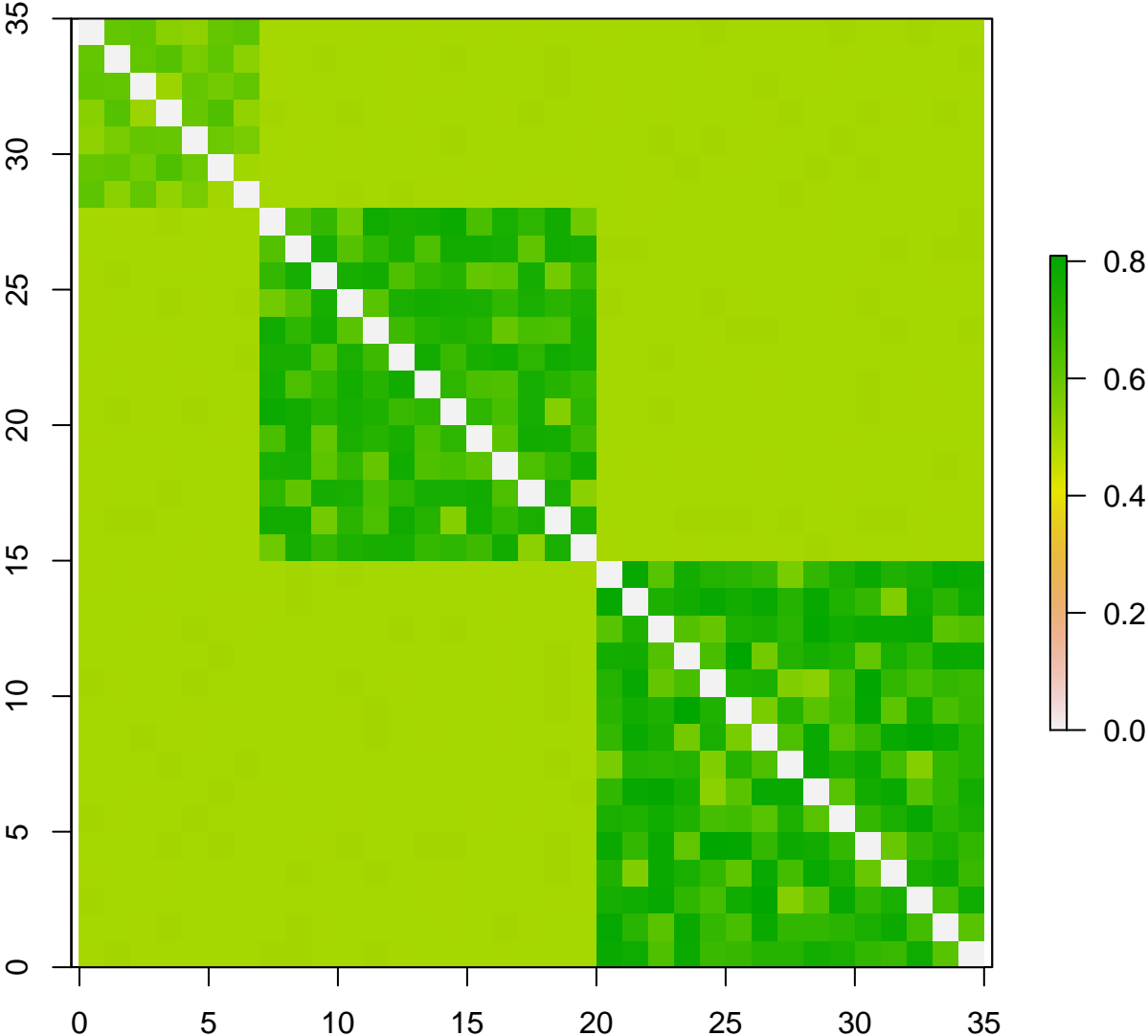
Density for ss marginal



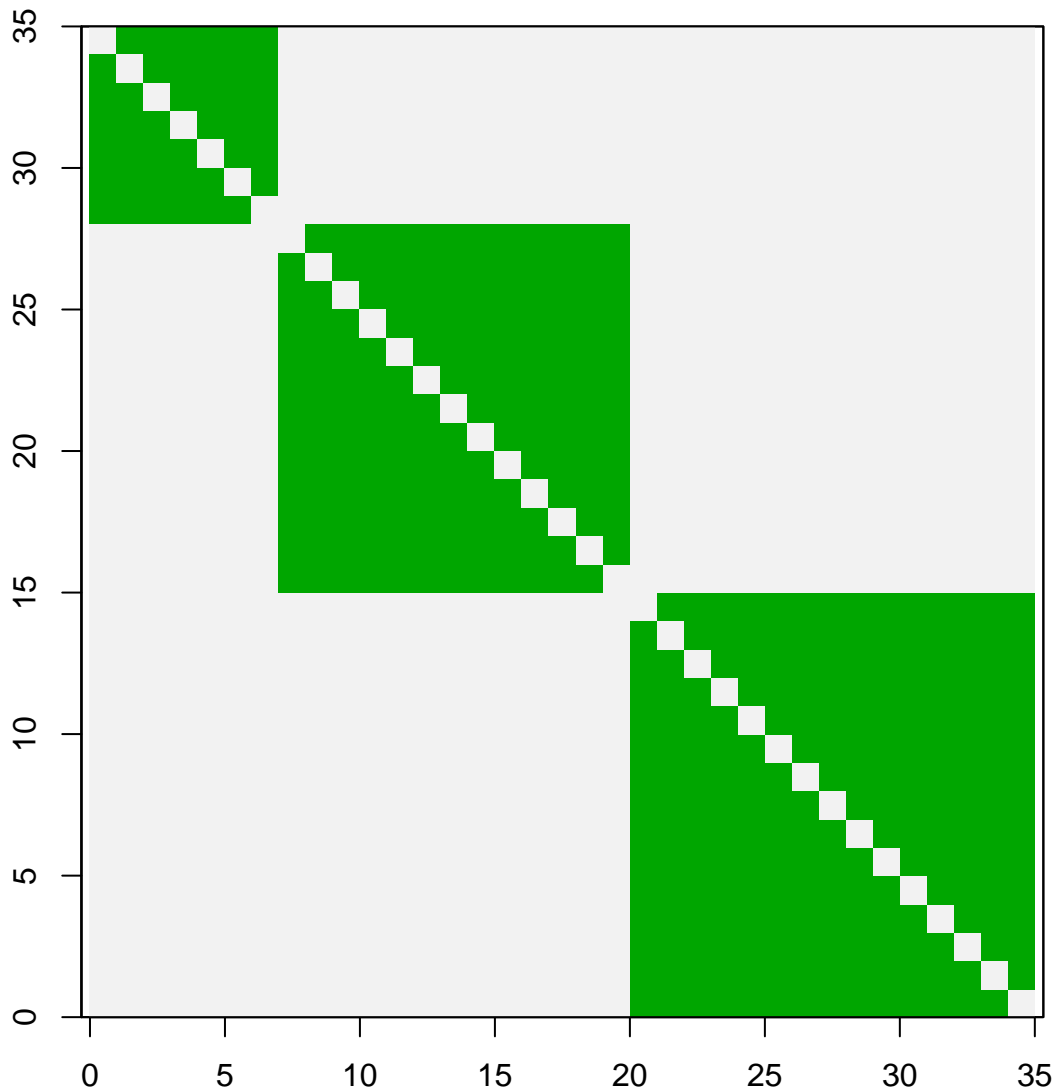
Estimated W – posterior mean



Acceptance rate for W



Start W



Solution W

