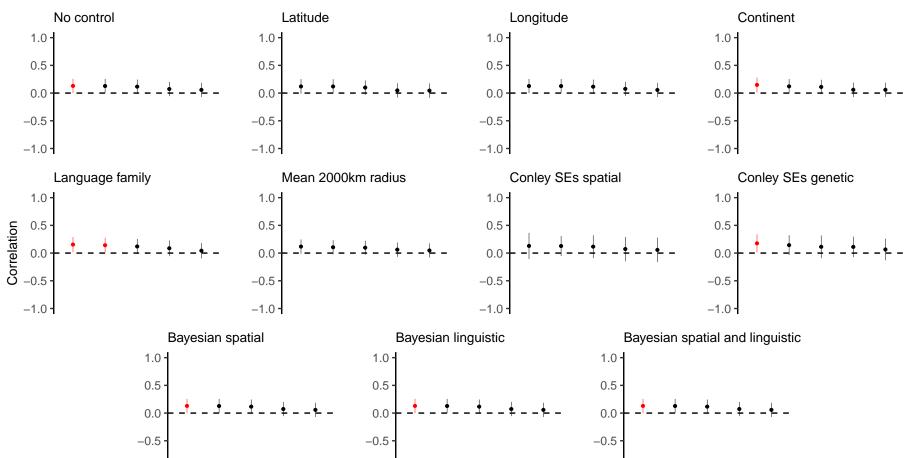
Simulation with strong cultural phylogenetic non-independence

-1.0 **-**



-1.0 **-**

-1.0 -