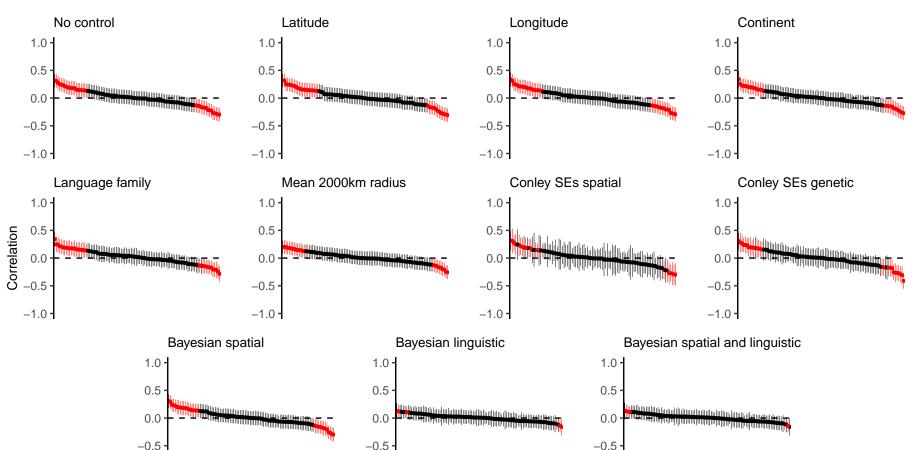
Simulation with strong cultural phylogenetic non-independence

-1.0 -



-1.0 -

-1.0