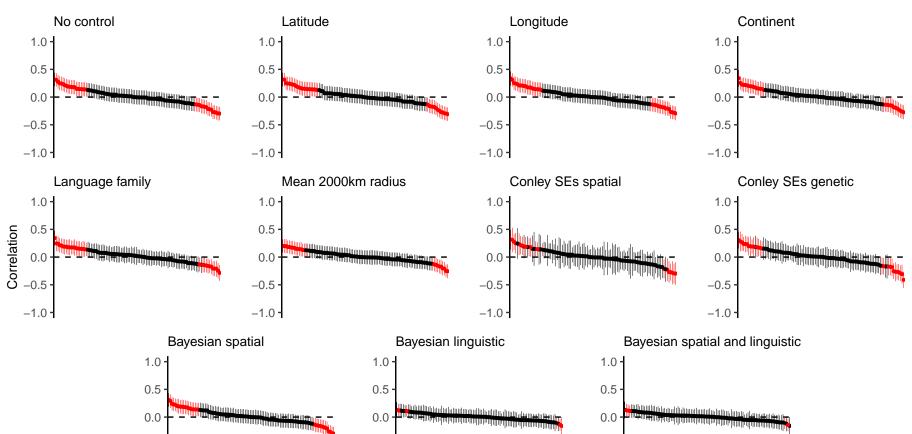
Simulation with strong cultural phylogenetic non-independence

-0.5

-1.0 -



-0.5

-1.0 -

-0.5

-1.0