Goodness of fit COV mean trait_uncorrelated.partial.TreeBUGS + • • package) trait.partial.TreeBUGS interaction(method, pooling, simple.no.TreeBUGS condition strong_naming simple.complete.TreeBUGS weak_naming PB/MLE.no.MPTinR asymptotic.no.MPTinR asymptotic.complete.MPTinR -0.25 0.50 0.75 0.25 0.50 0.75 1.00 0.00 1.000.00