Distribution of p-values for summary statistic sigsq.est var.contrast cor.contrast.var 200 -150 **-**100 -50 · trait count leafN cor.contrast.asr cor.contrast.nh ks.contrast seedMass sla 200 150 -100 -50 0 0.50 0.75 0.25 0.50 0.75 1.00 0.00 0.25 1.00 0.00 0.25 0.50 0.75 1.00 0.00 p-value