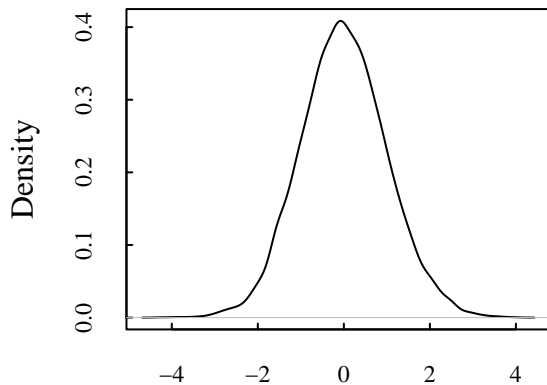


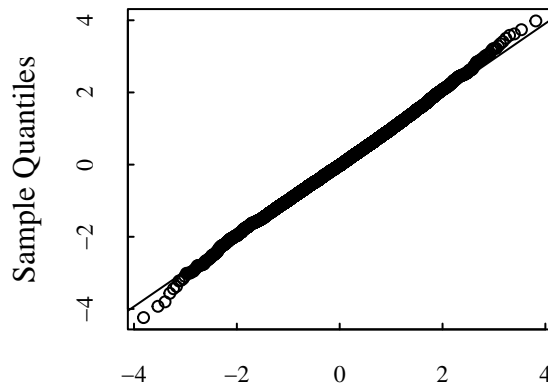
# Reptiles: diagnostic plots for the PGLS

**density.default(x = res)**

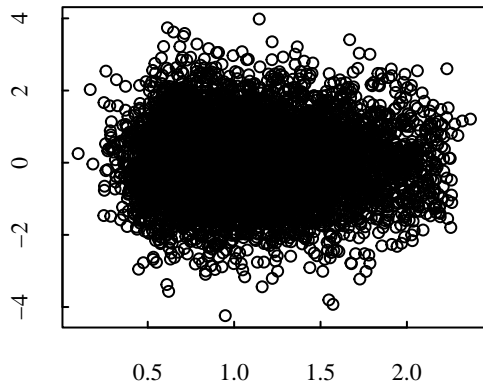


N = 7330 Bandwidth = 0.1498

**Normal Q-Q Plot**

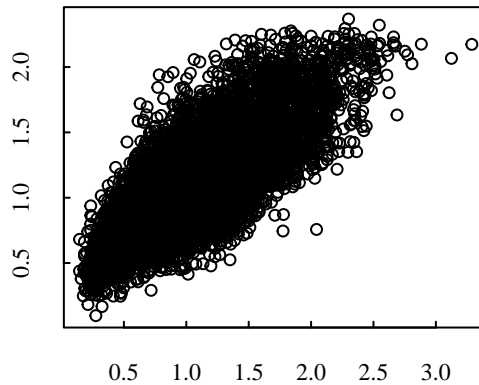


**Residual value (corrected for phylogeny)**



Fitted value

**Fitted value**



Observed value