



Figure 3 - figure supplement 1 | The common evolutionary rate of gene expression levels presented in Fig. 3 is robust to changes in correlation metrics or expression threshold. a,b, The gene expression levels used in Fig. 3 were correlated with alternative correlation metrics, Kendalls τ (a) and Pearson's R (b). The resulting evolutionary rates remained statistically indistinguishable. **c,** The gene expression level of all genes were analyzed rather than excluding the values below 5 TPM as was done in Fig. 3. The resulting evolutionary rates remained statistically indistinguishable.