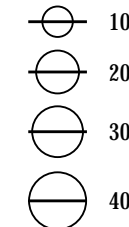


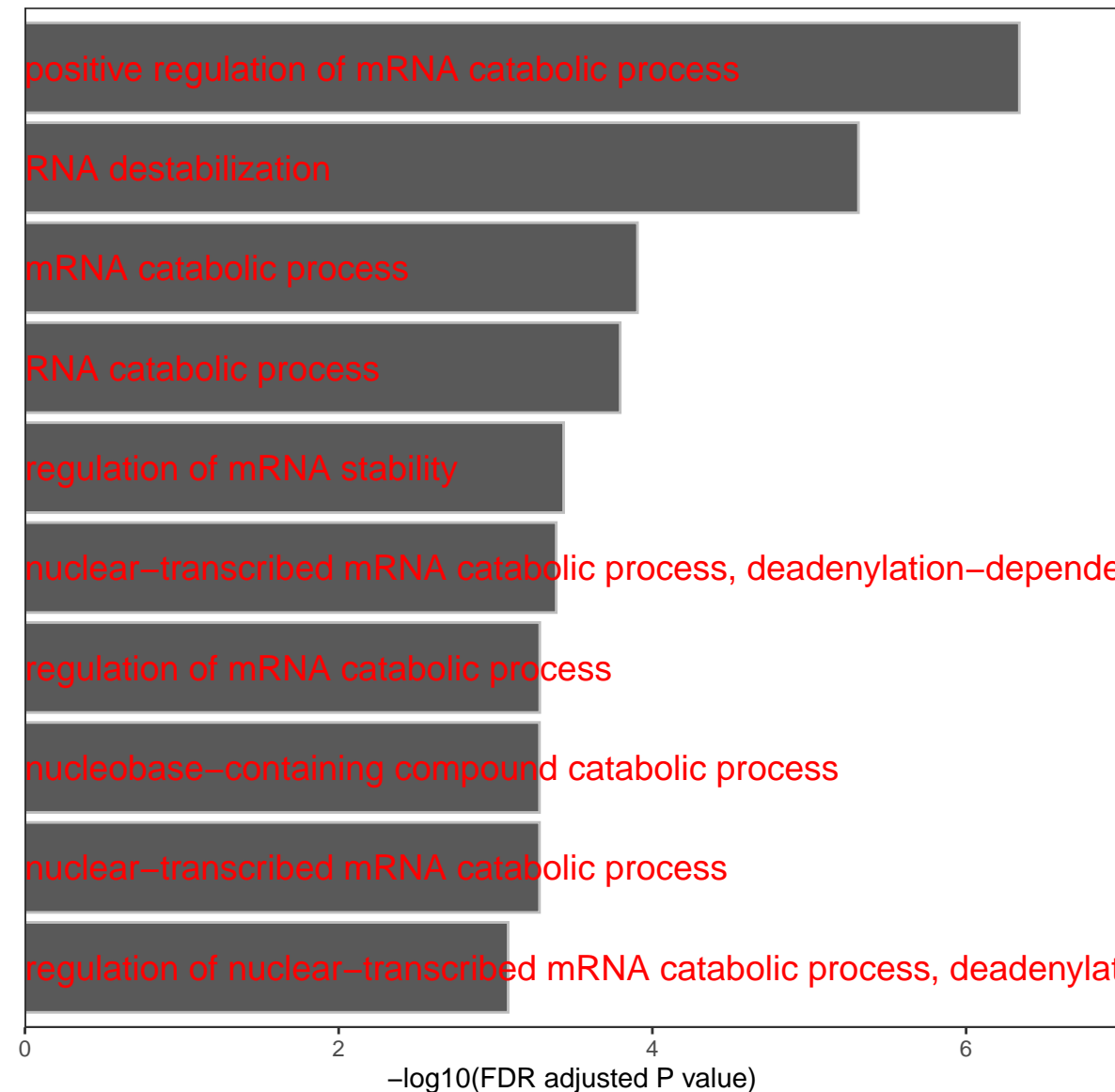
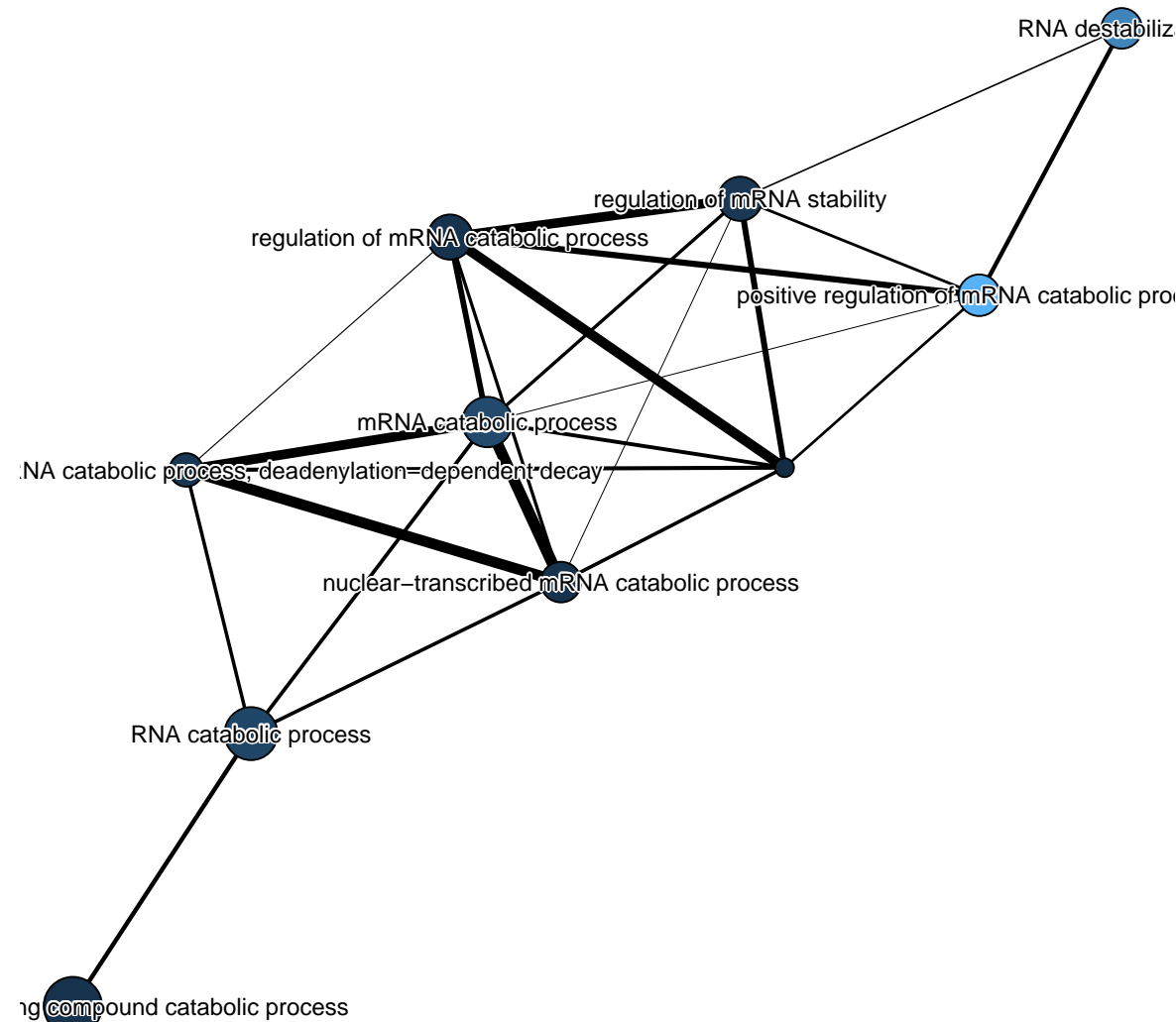
Count



$-\log(p.adjust, 10)$



sim



process mRNA

decay compound destabilization

stability **catabolic** RNA

deadenylation-dependent

nuclear-transcribed

nucleobase-containing regulation