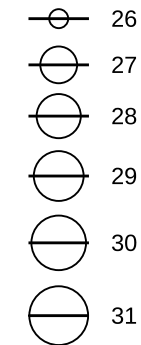


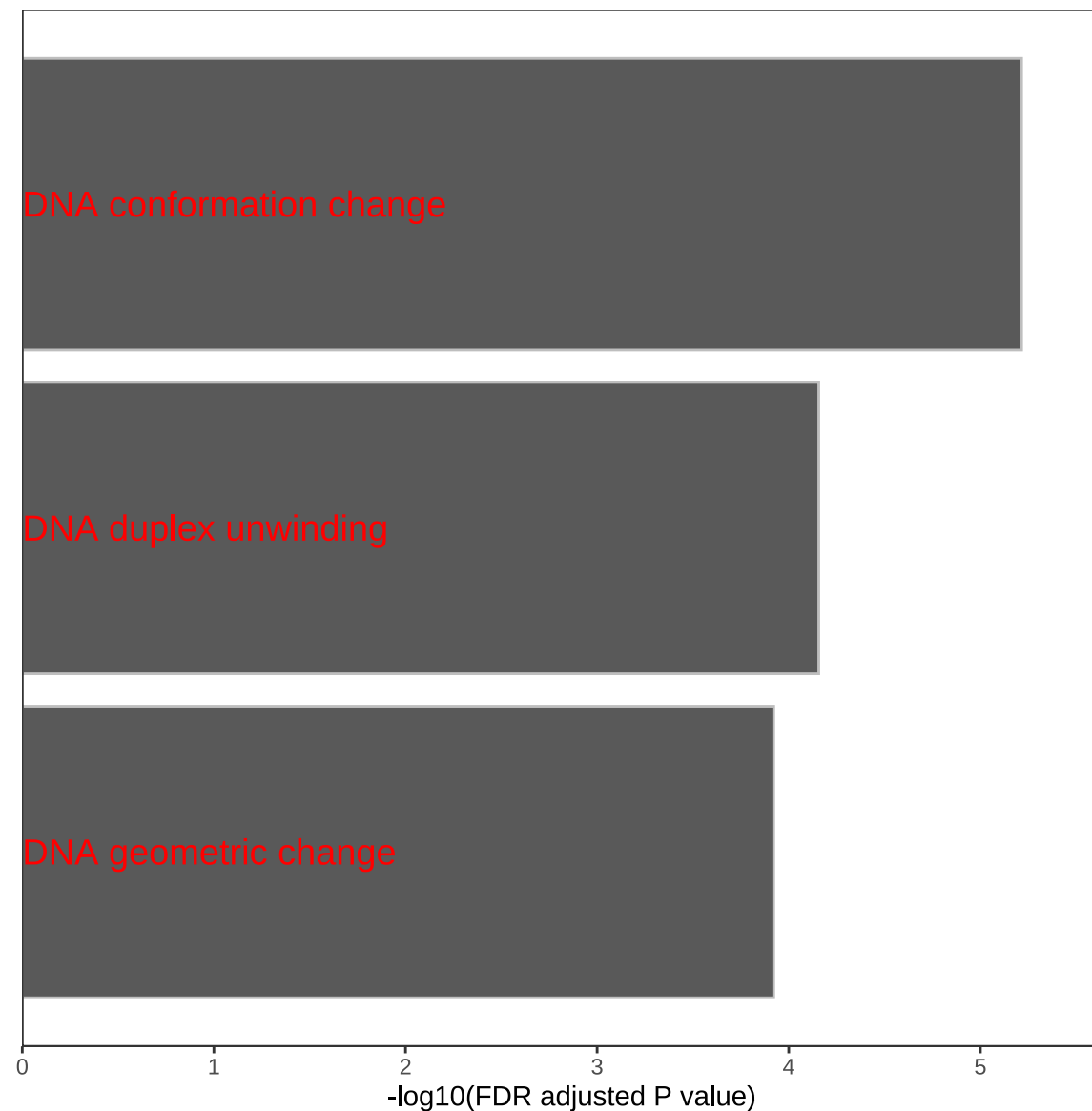
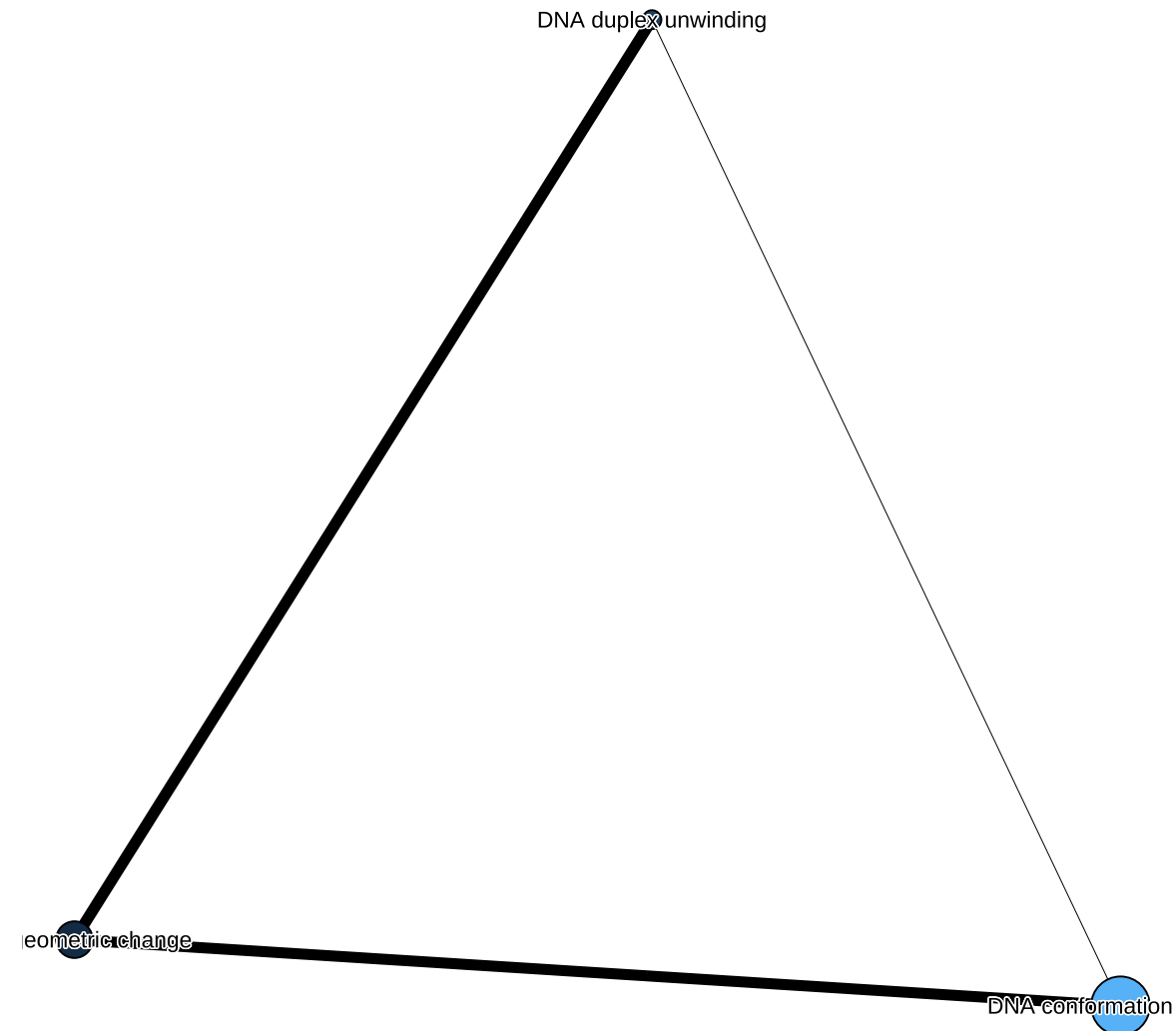
Count



sim



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



geometric change duplex
conformation
DNA unwinding