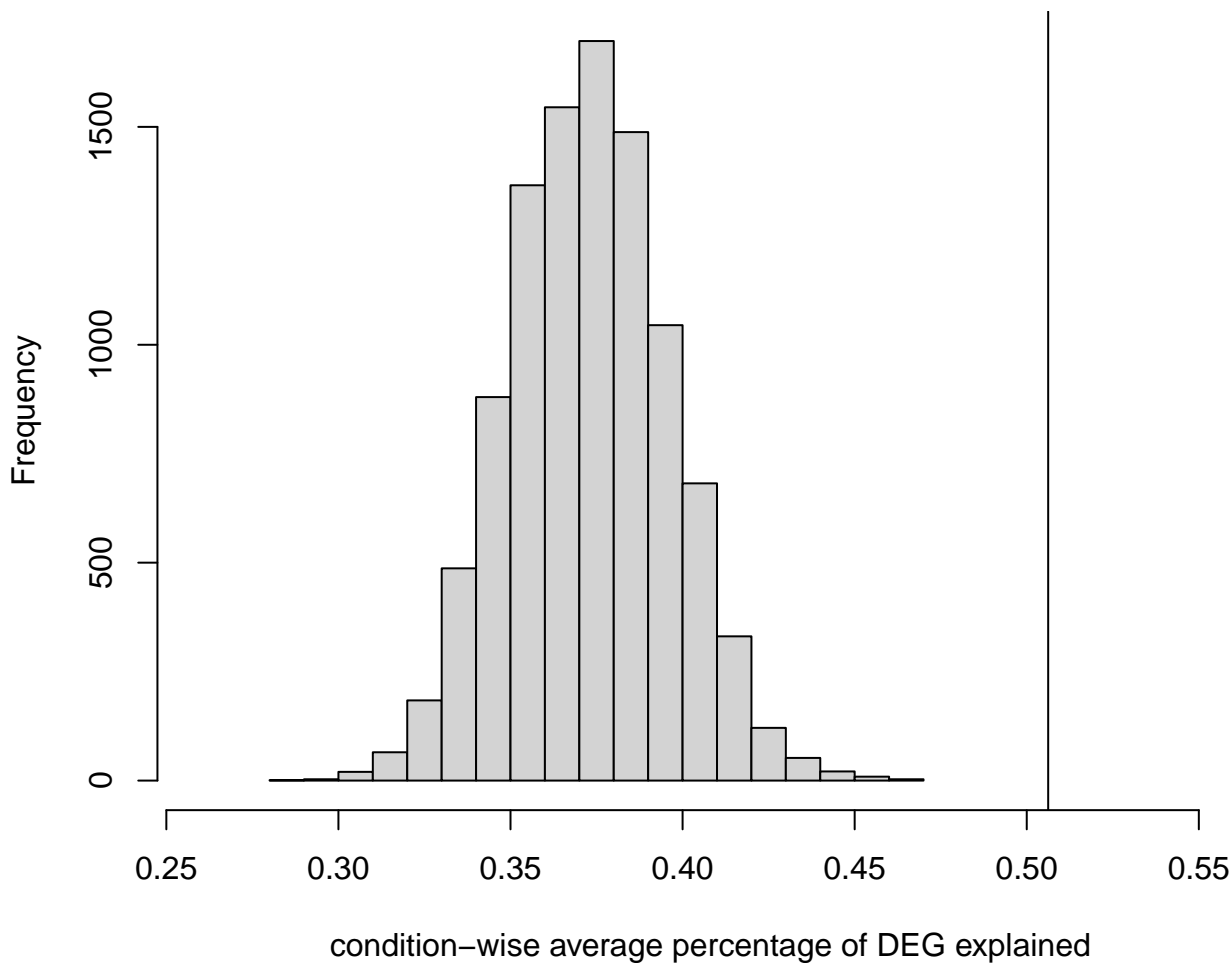
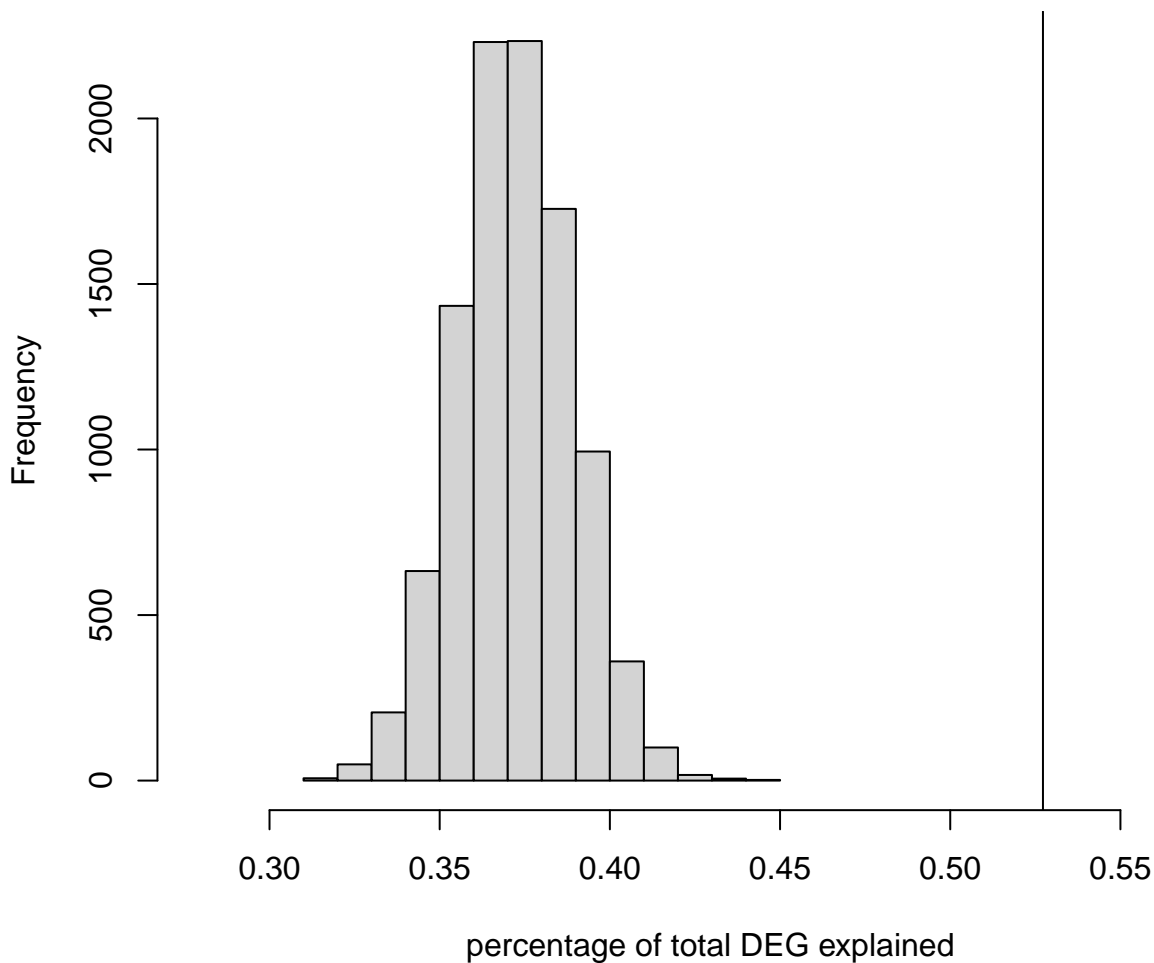


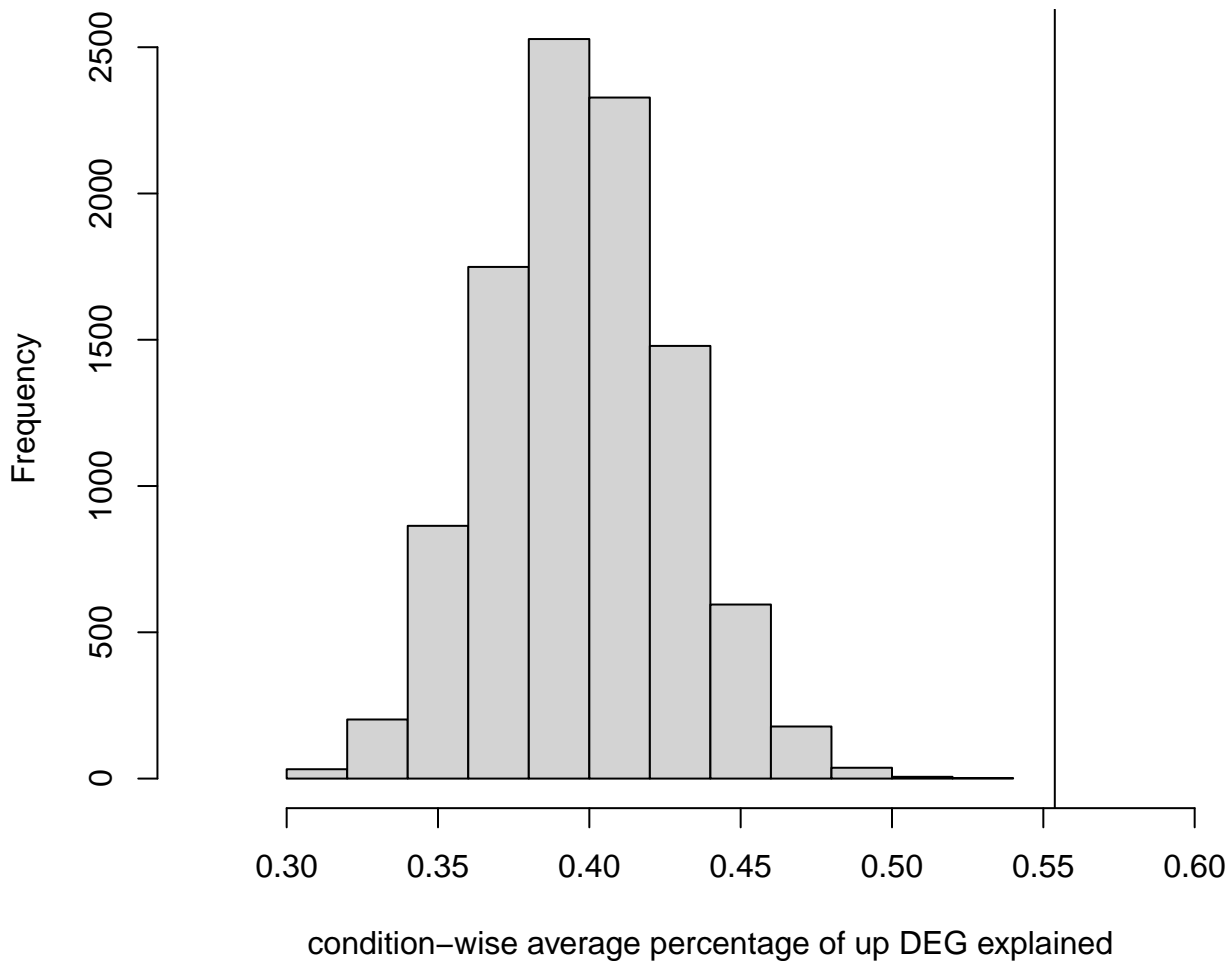
CR model – full data
p < 1e-04



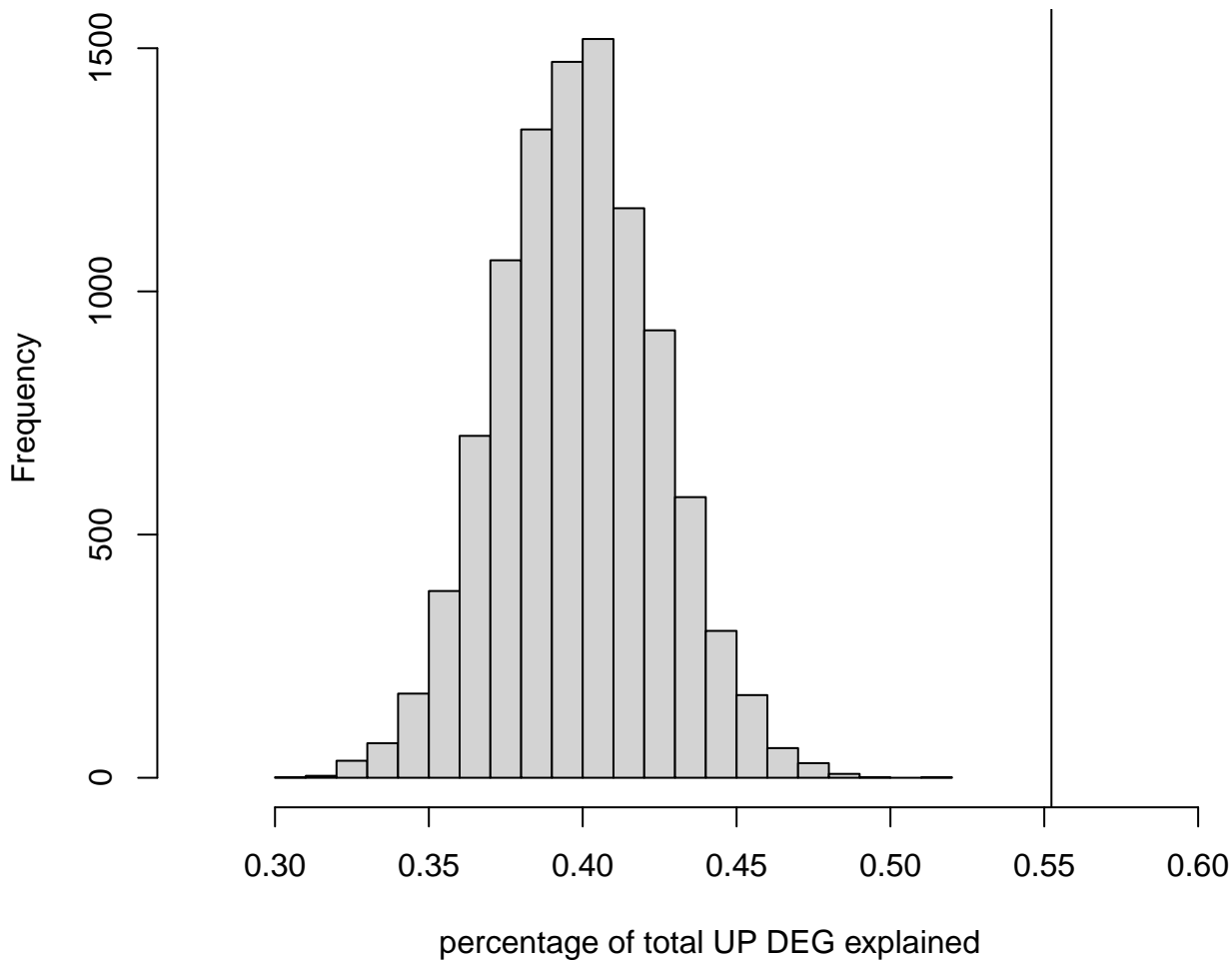
CR model – full data
 $p < 1e-04$



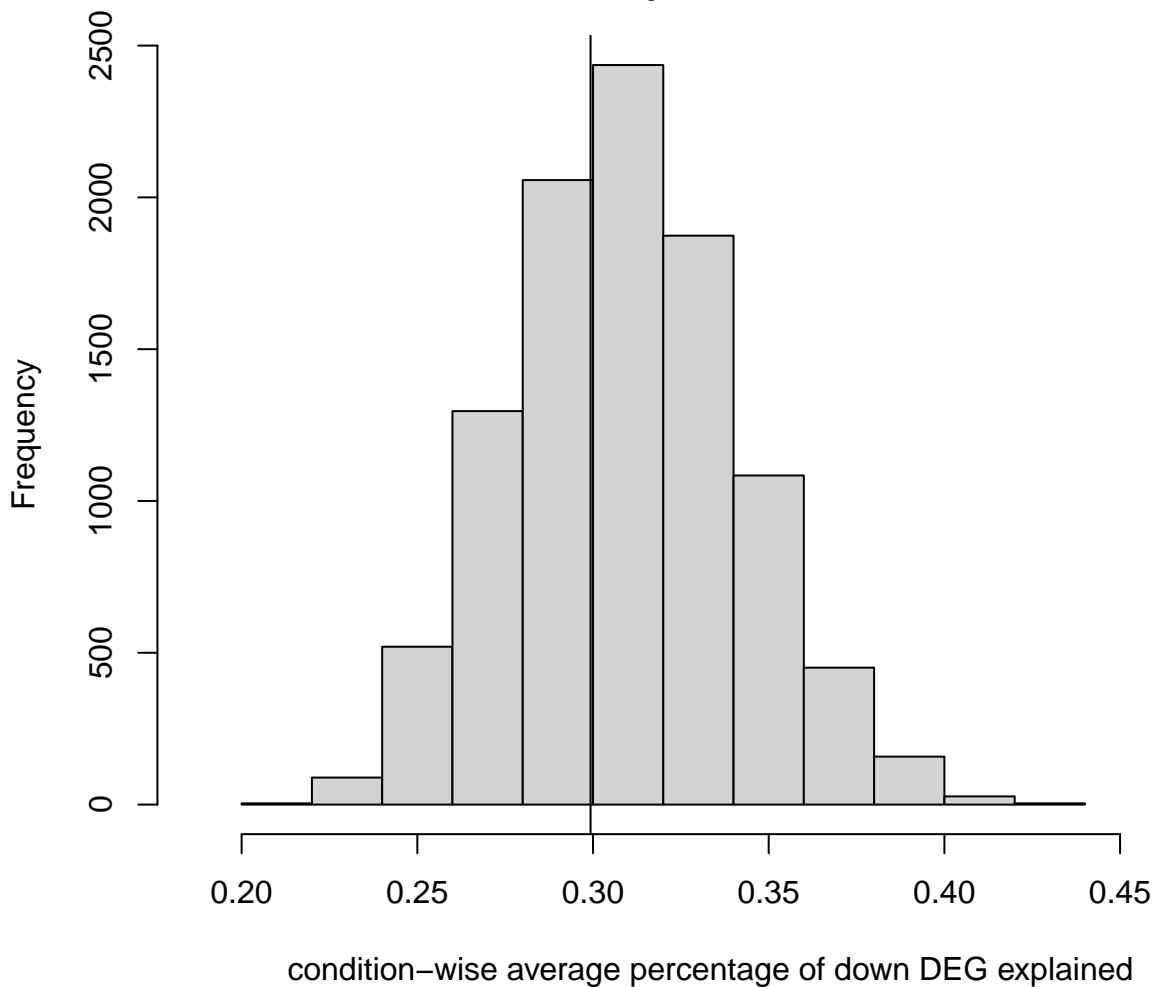
CR model – full data
p < 1e-04



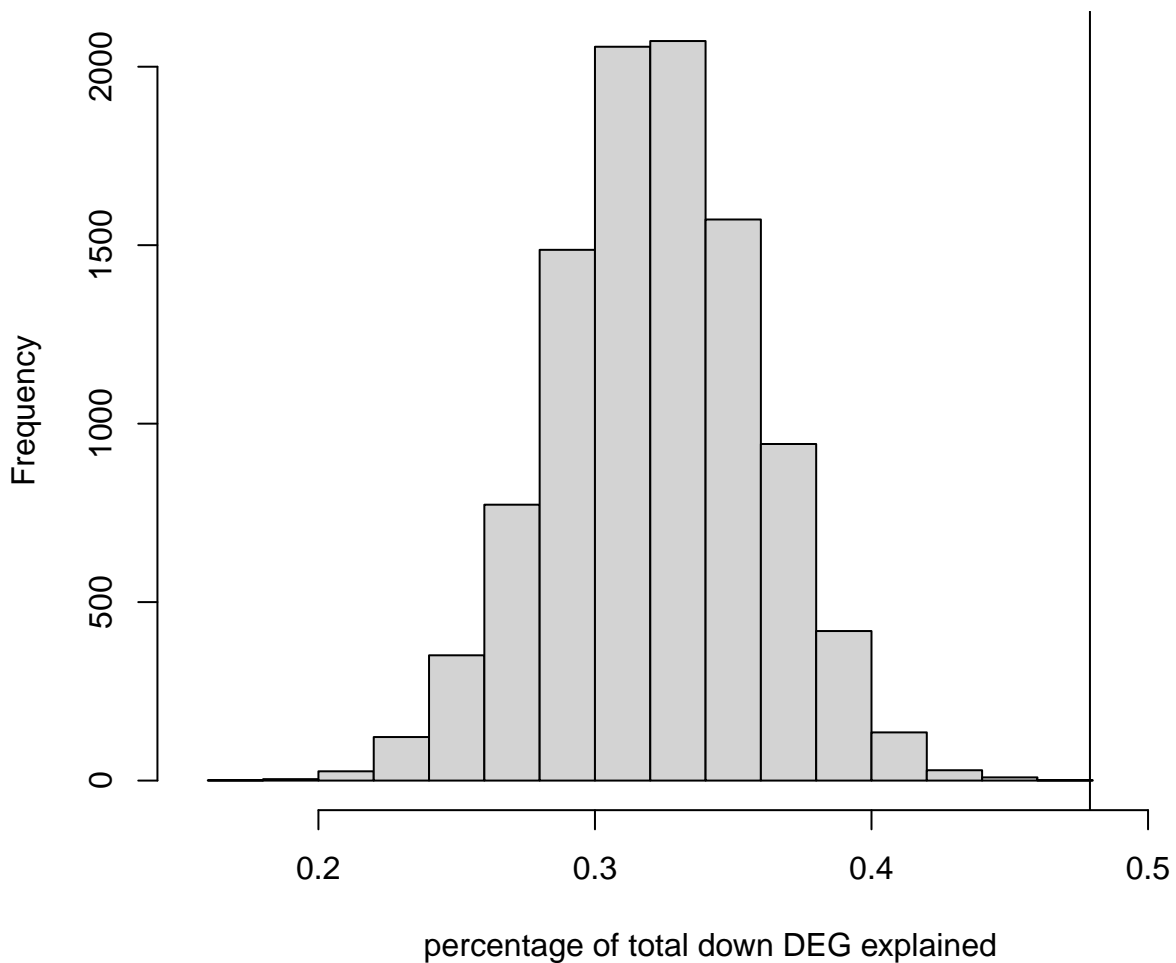
CR model – full data
p < 1e-04



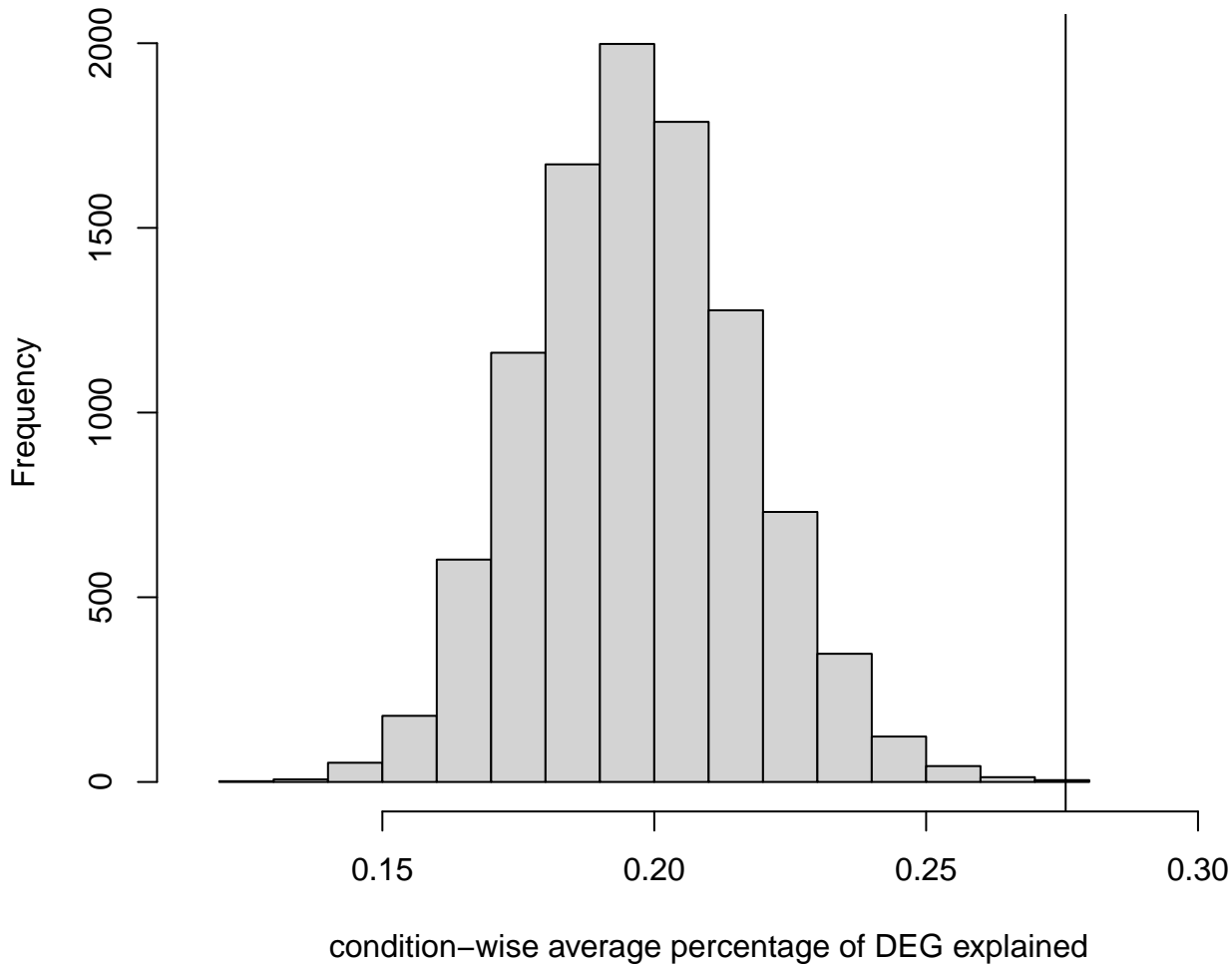
CR model – full data
p < 0.61



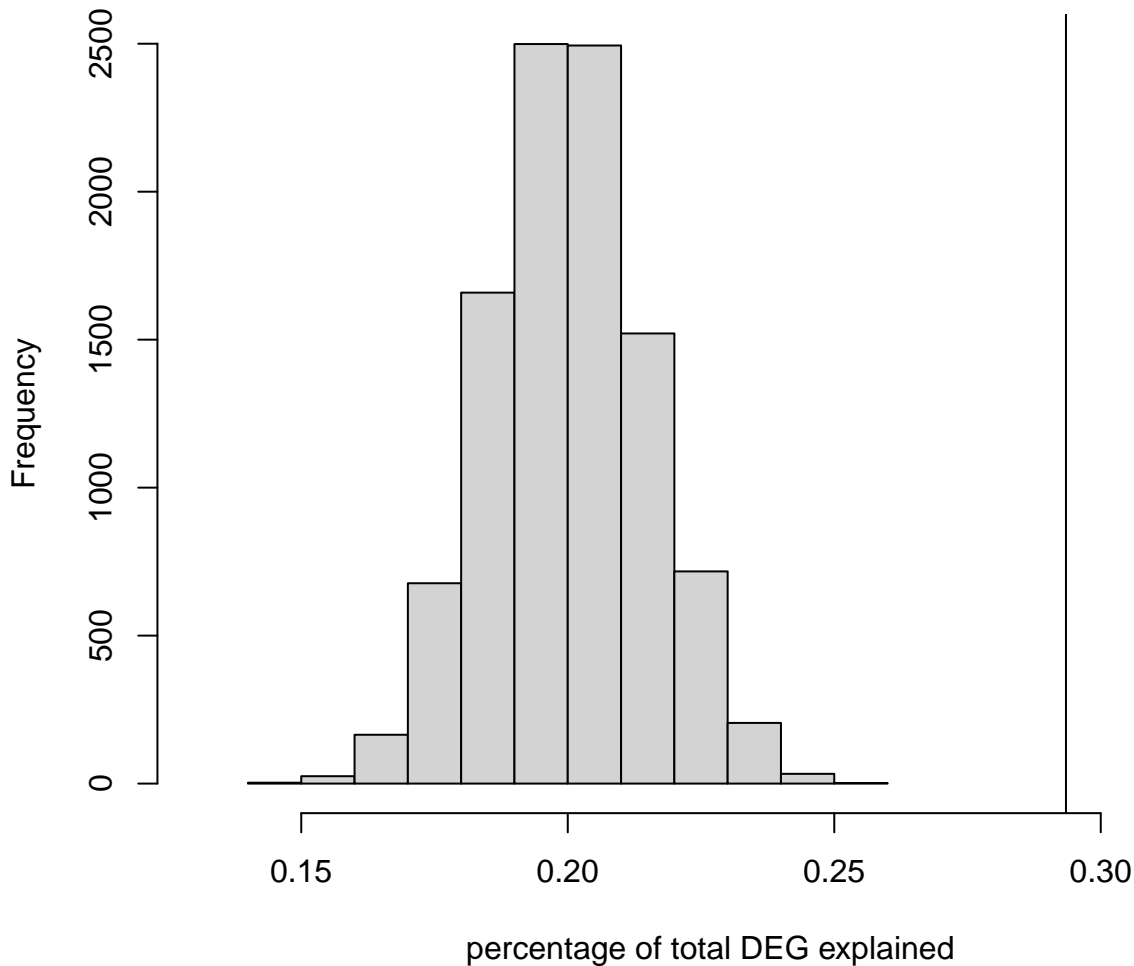
CR model – full data
 $p < 1e-04$



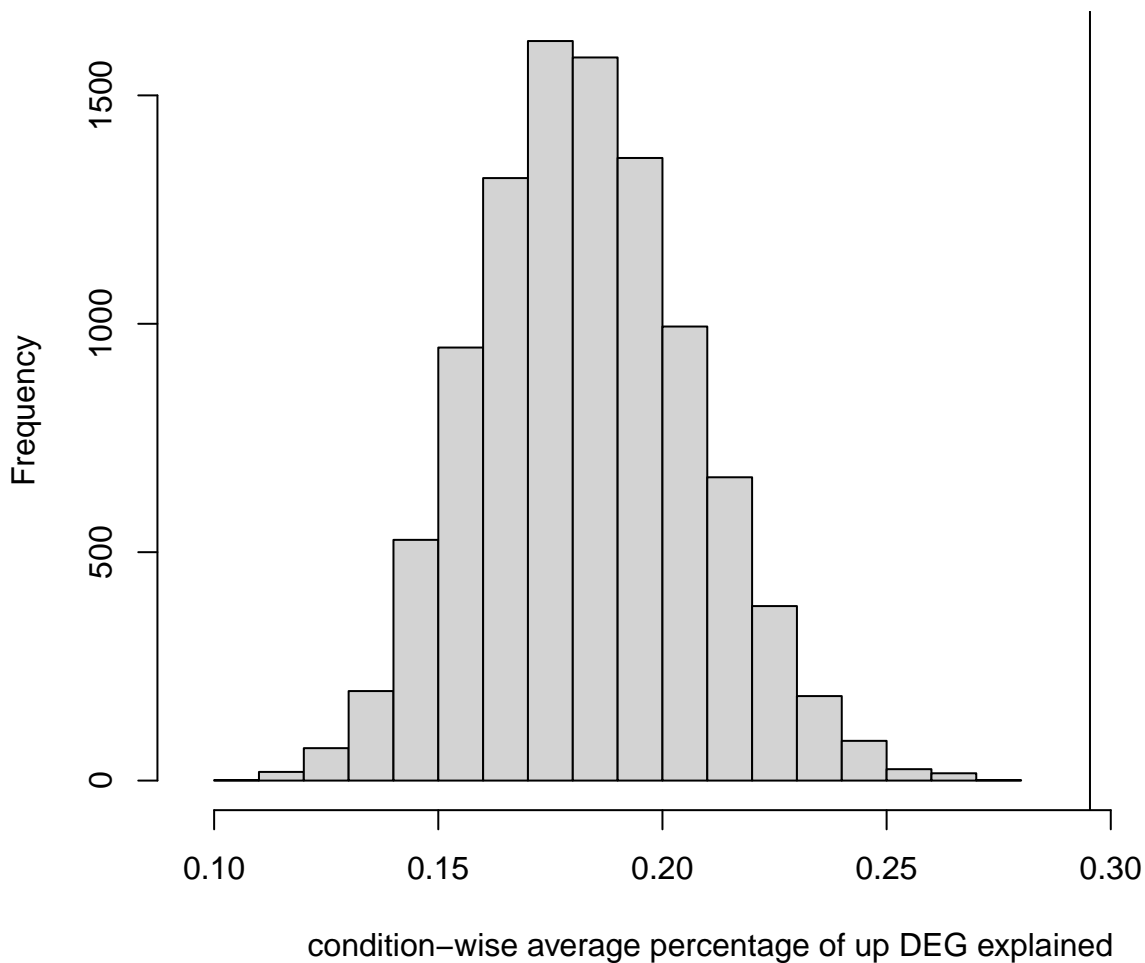
CR model – single data
 $p < 3e-04$



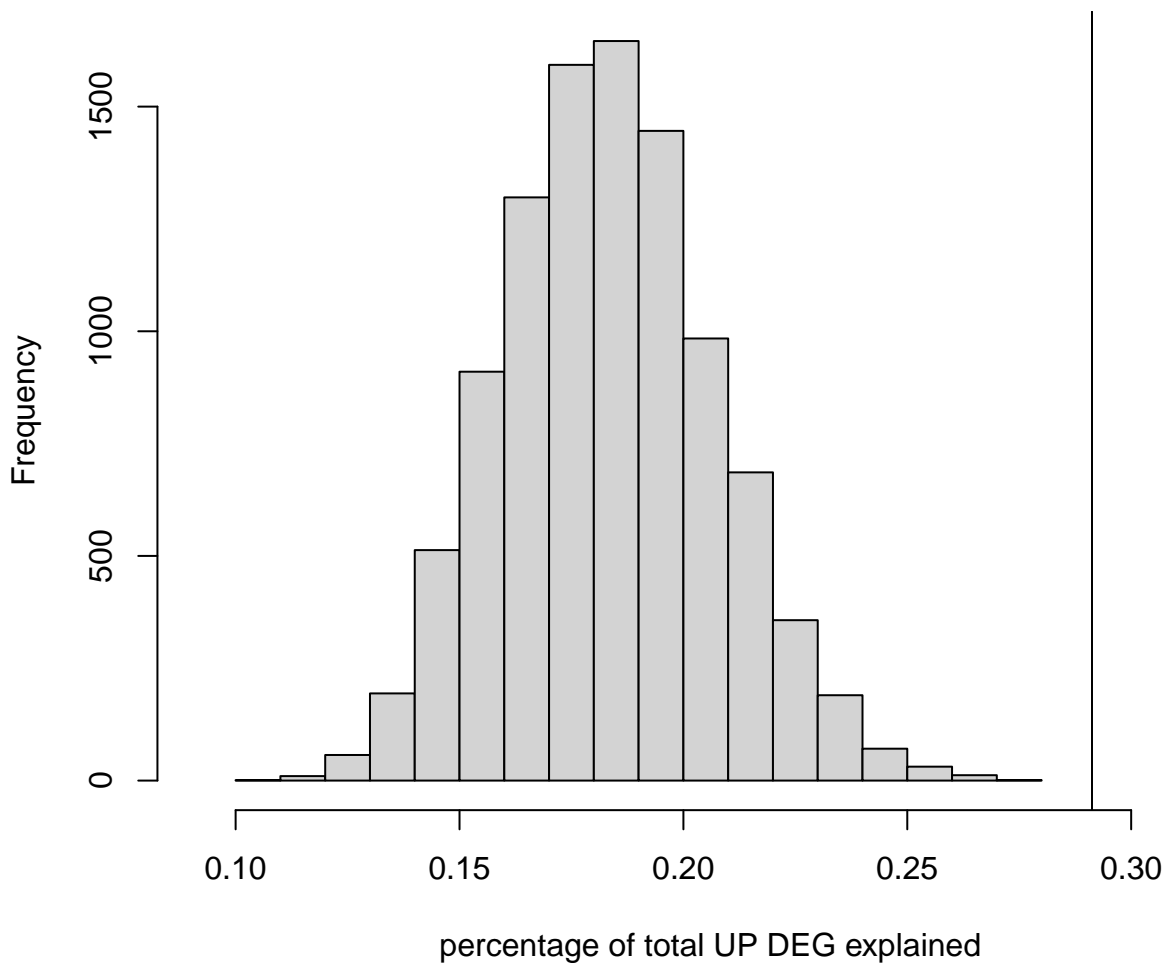
CR model – single data
 $p < 1e-04$



CR model – single data
 $p < 1e-04$

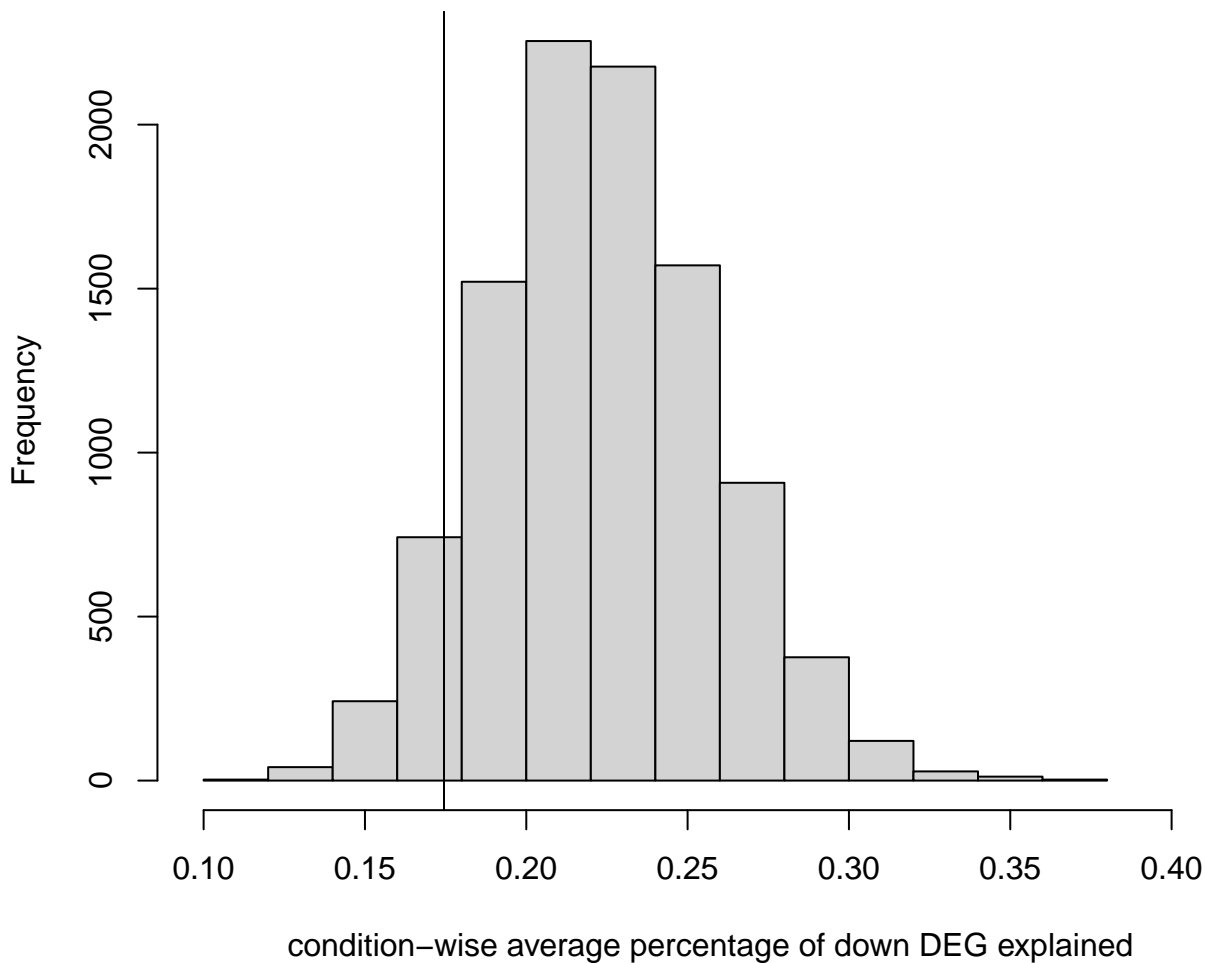


CR model – single data
 $p < 1e-04$

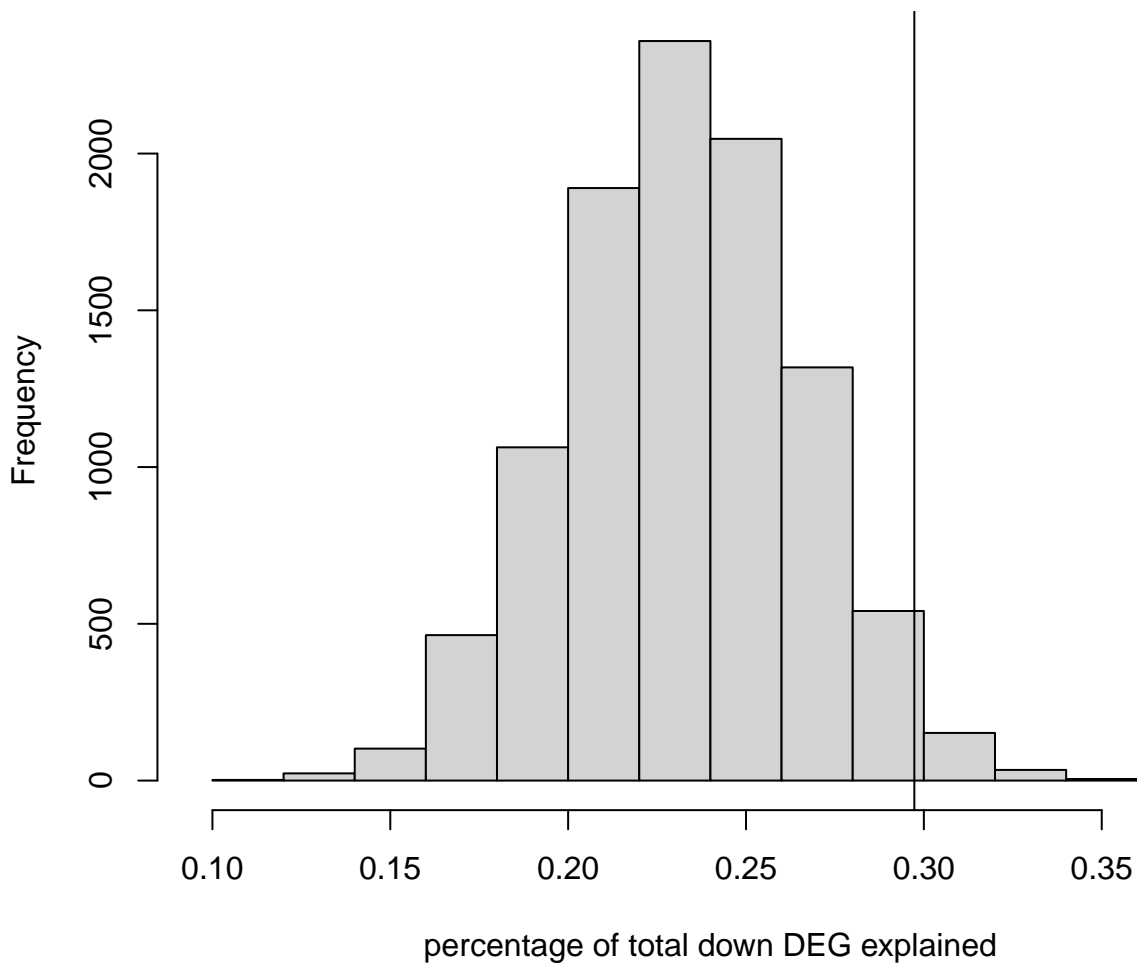


CR model – single data

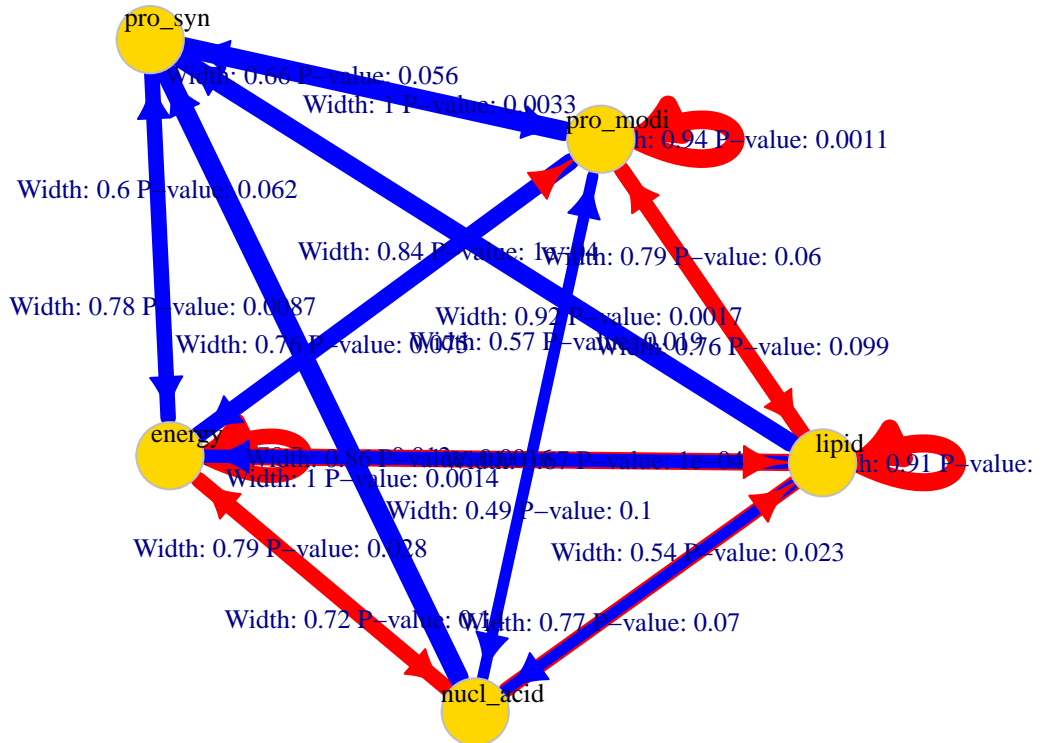
$p < 0.92$



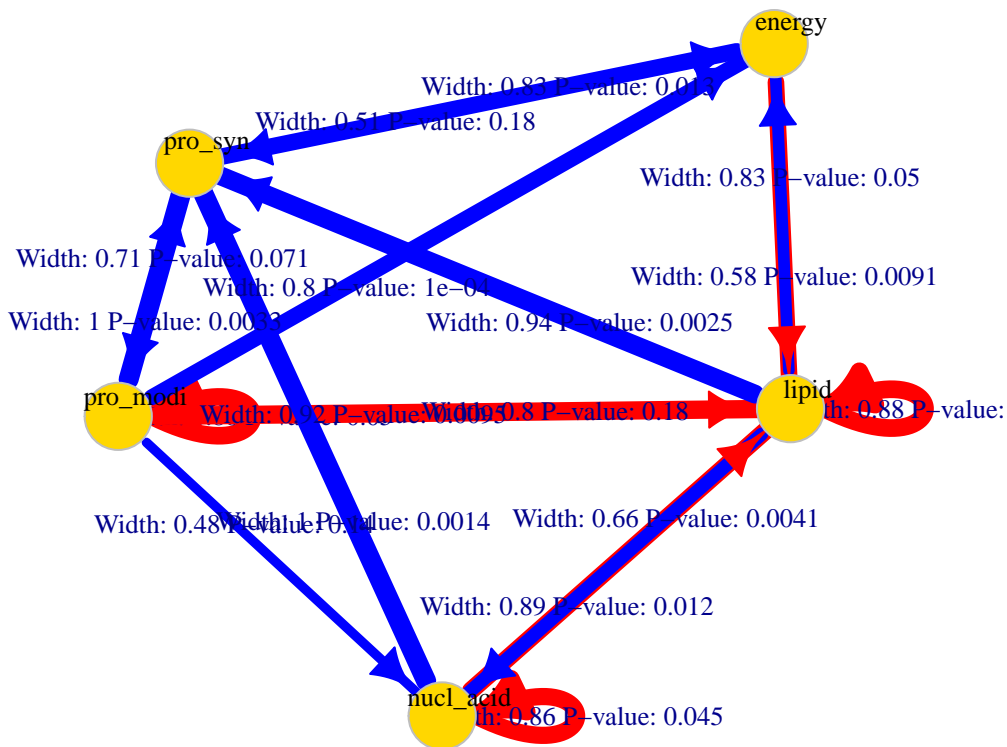
CR model – single data
p < 0.023



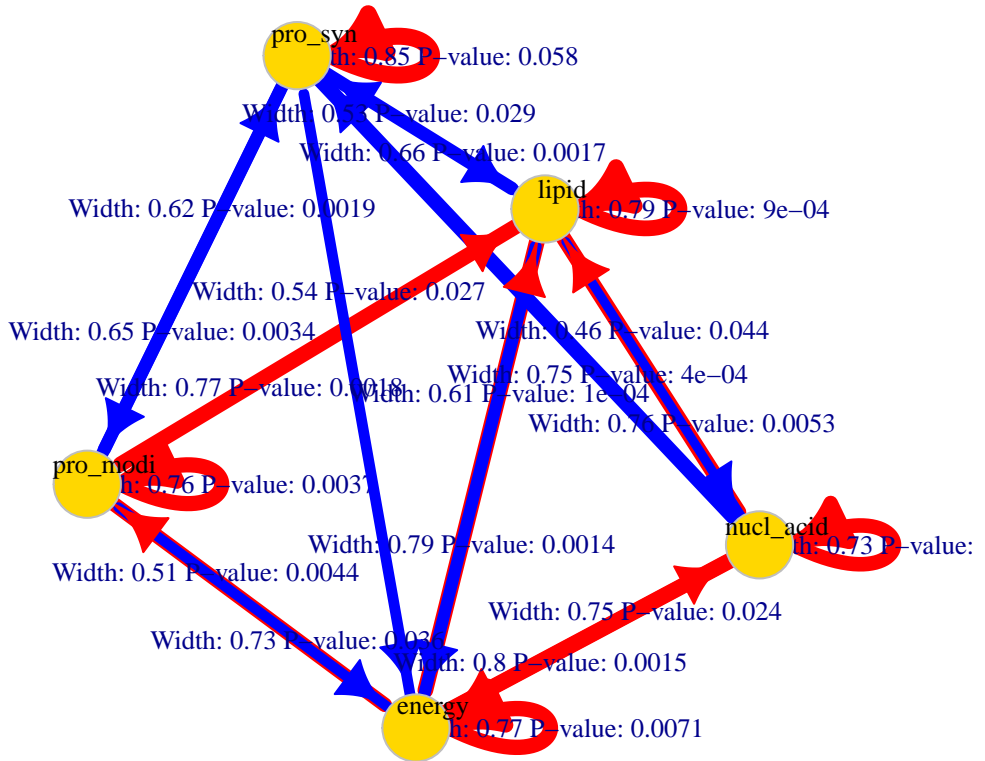
single – raw – p < 0.2



single – normalized – p < 0.2



full – raw – $p < 0.2$



full – normalized – p < 0.2

