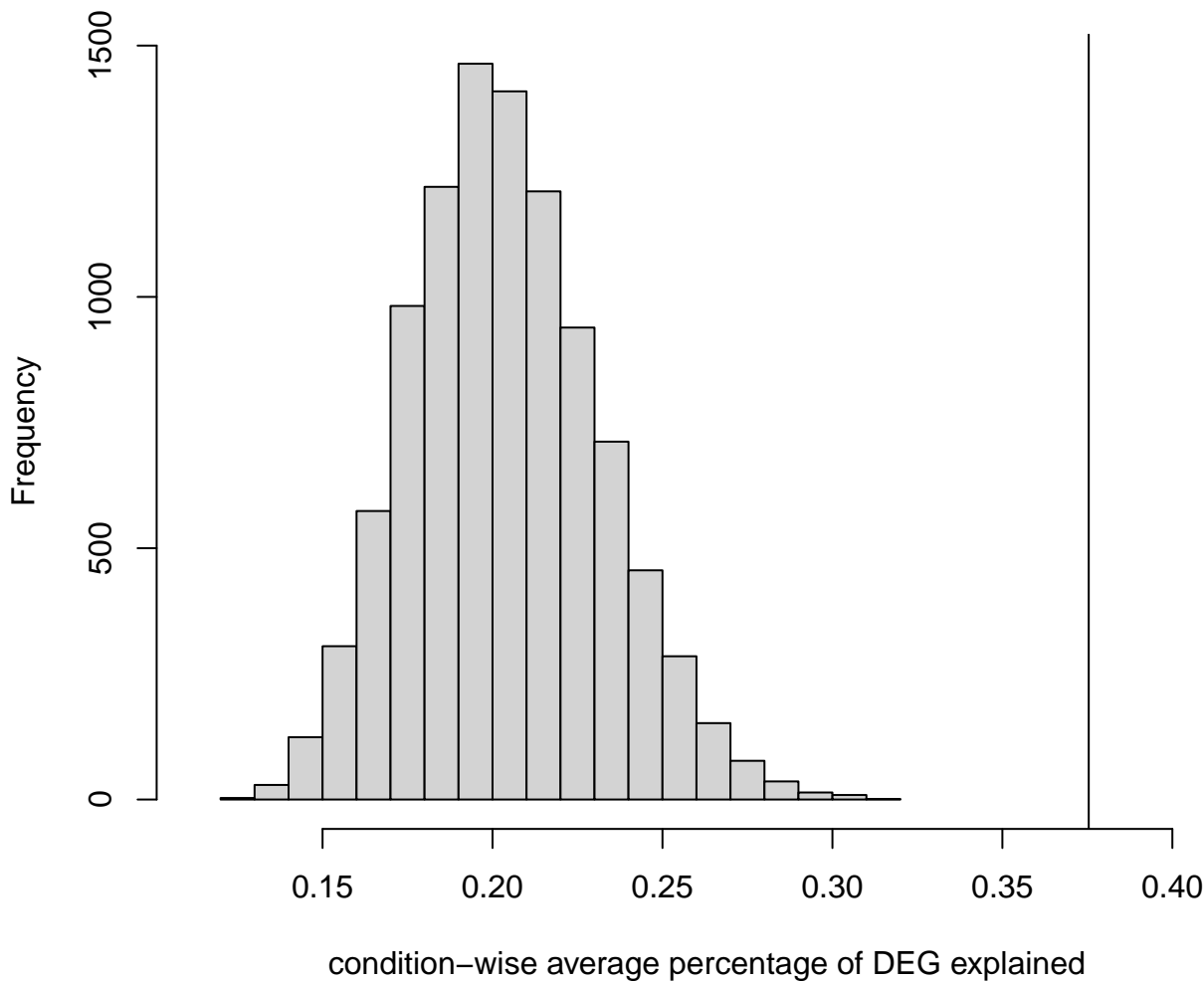
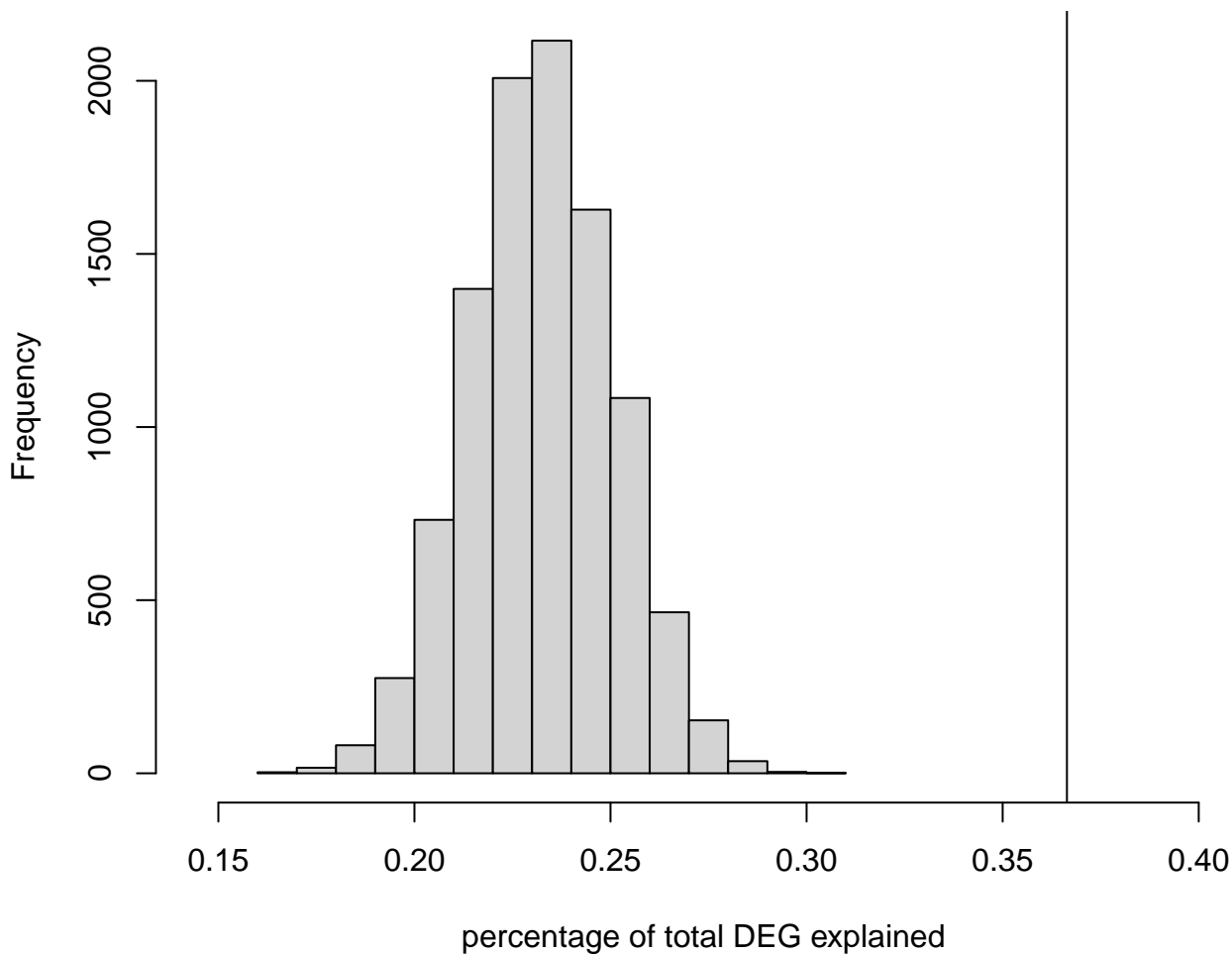


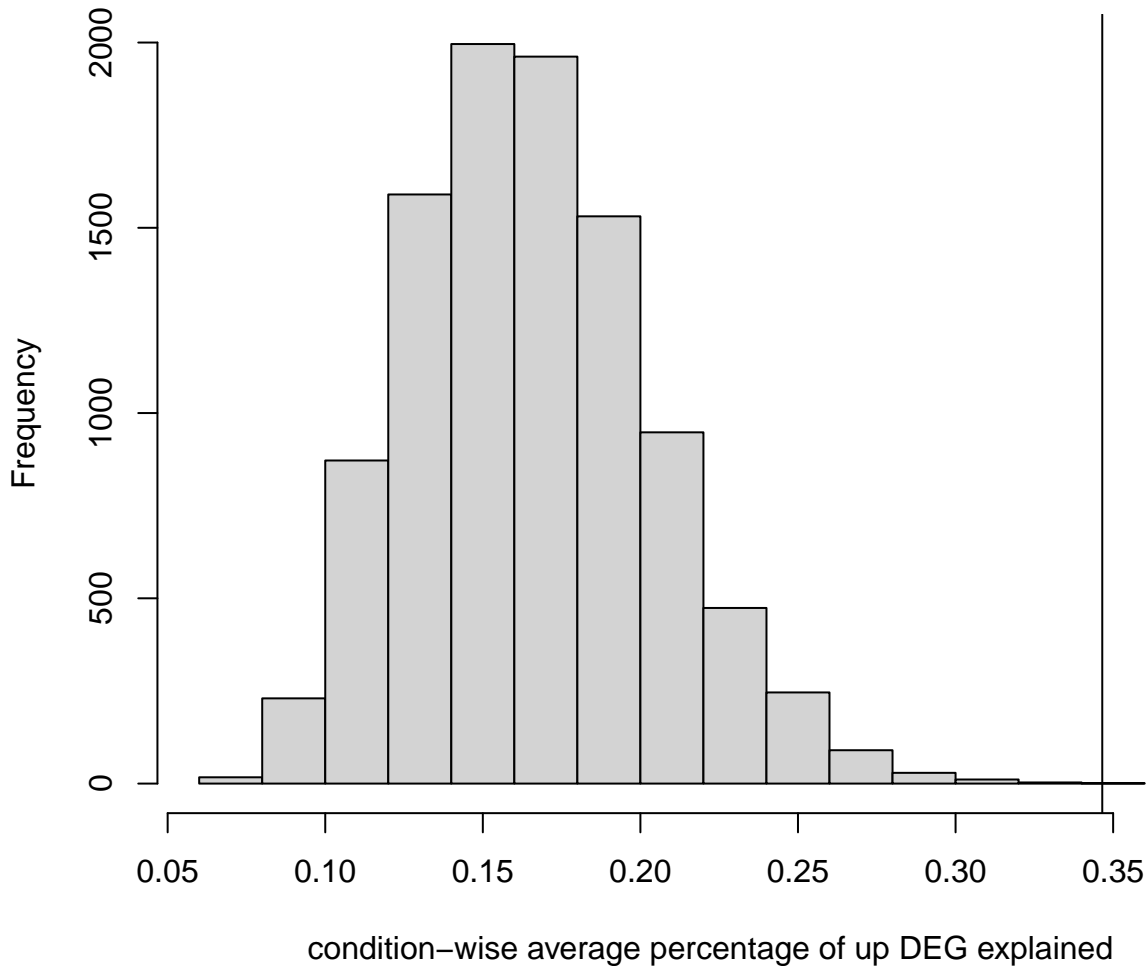
CR model – full data
p < 1e-04



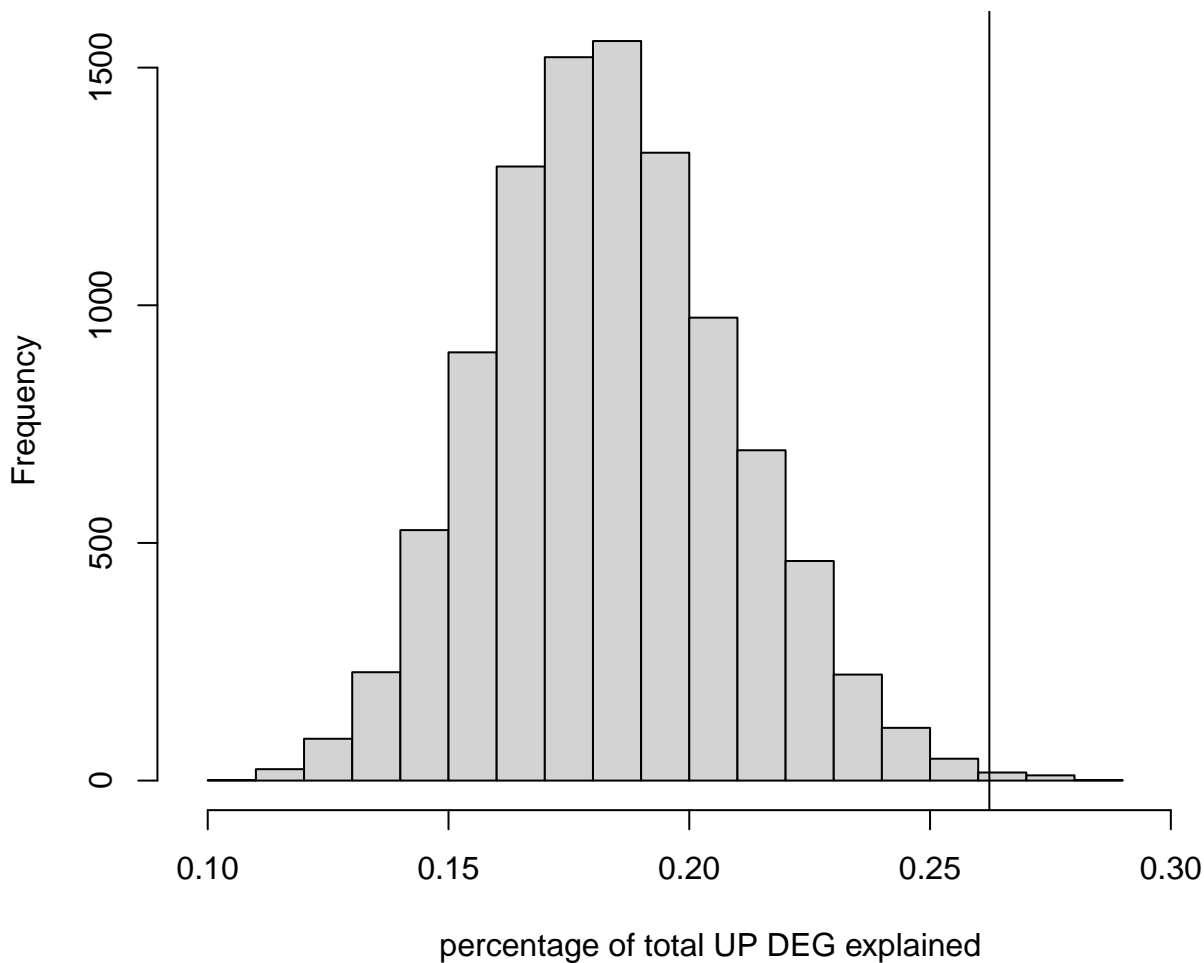
CR model – full data
p < 1e-04



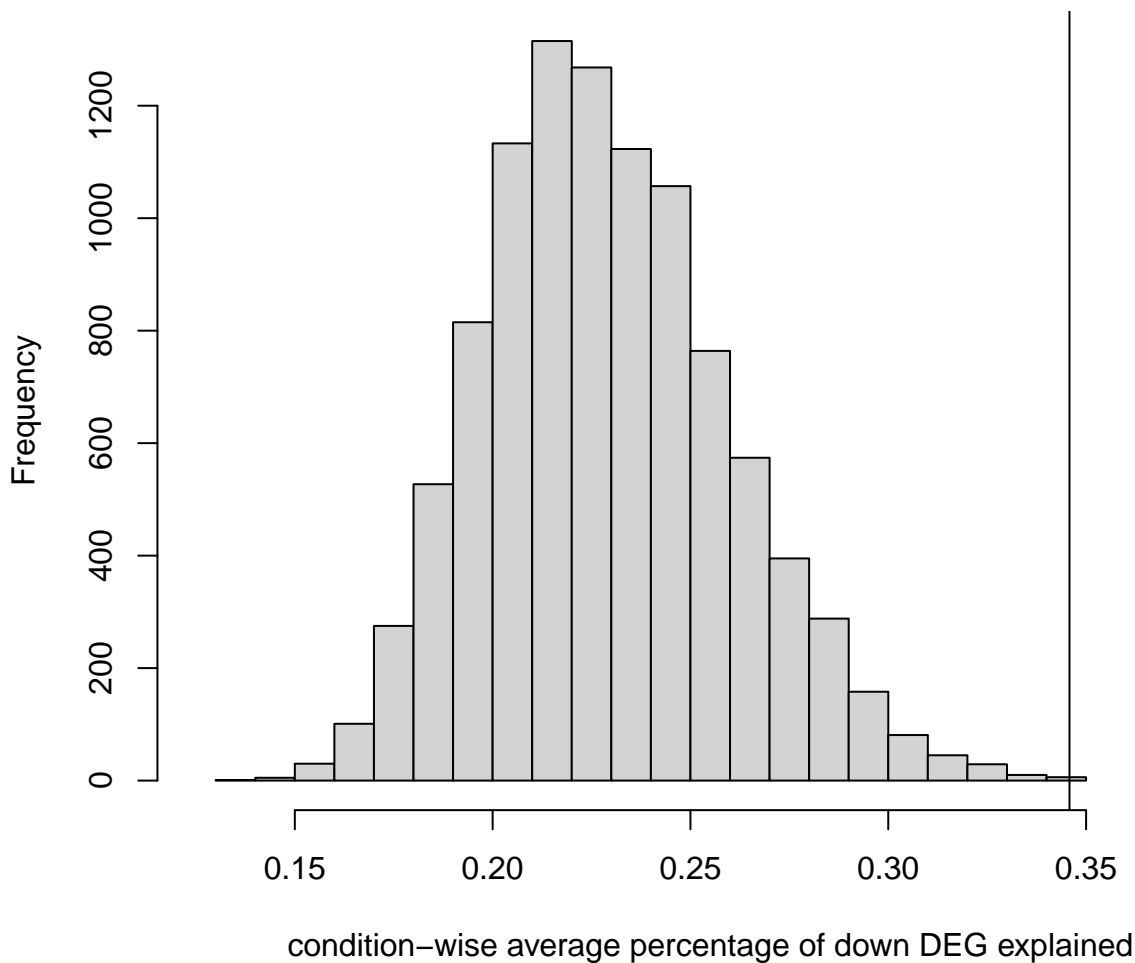
CR model – full data
 $p < 2e-04$



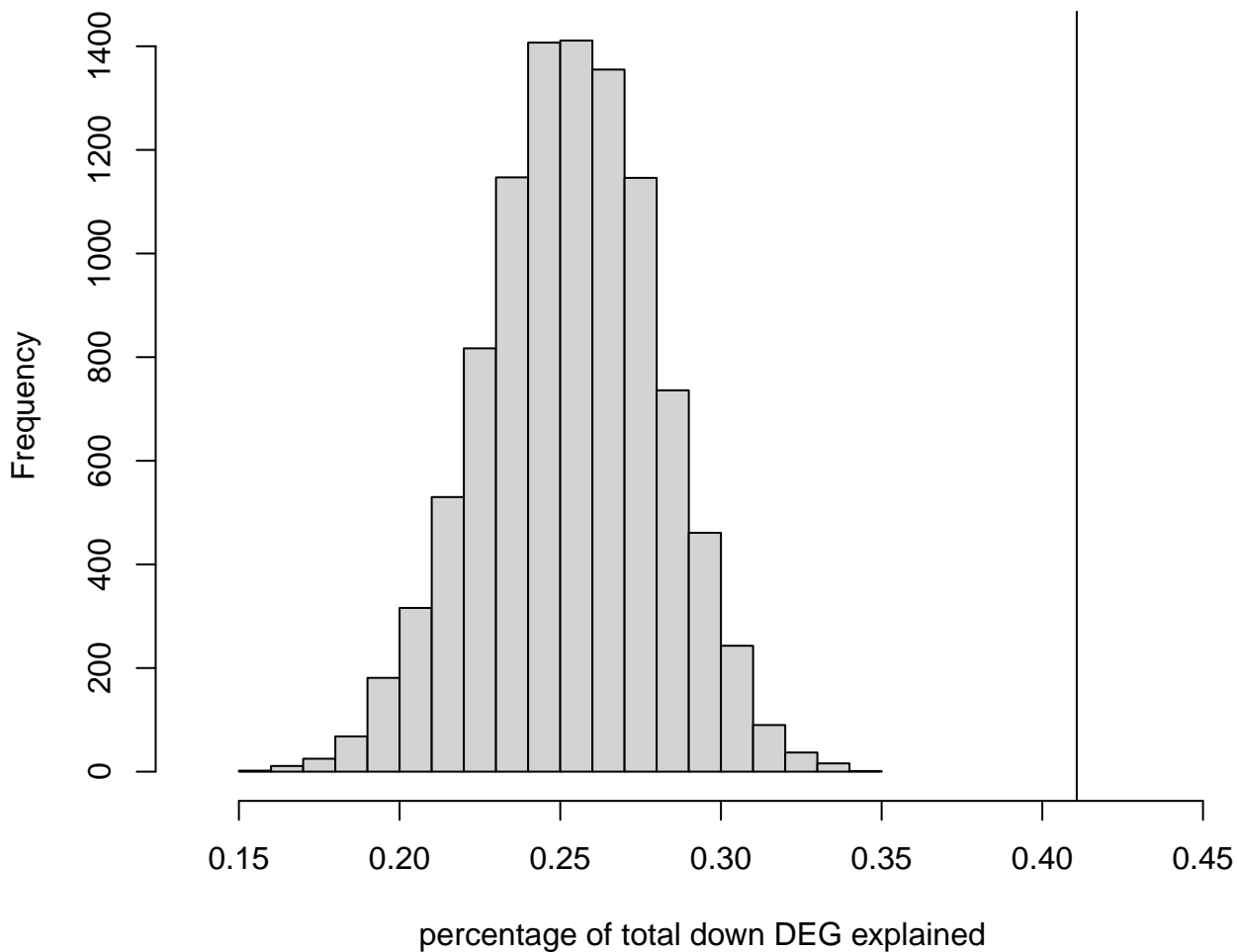
CR model – full data
p < 0.0025



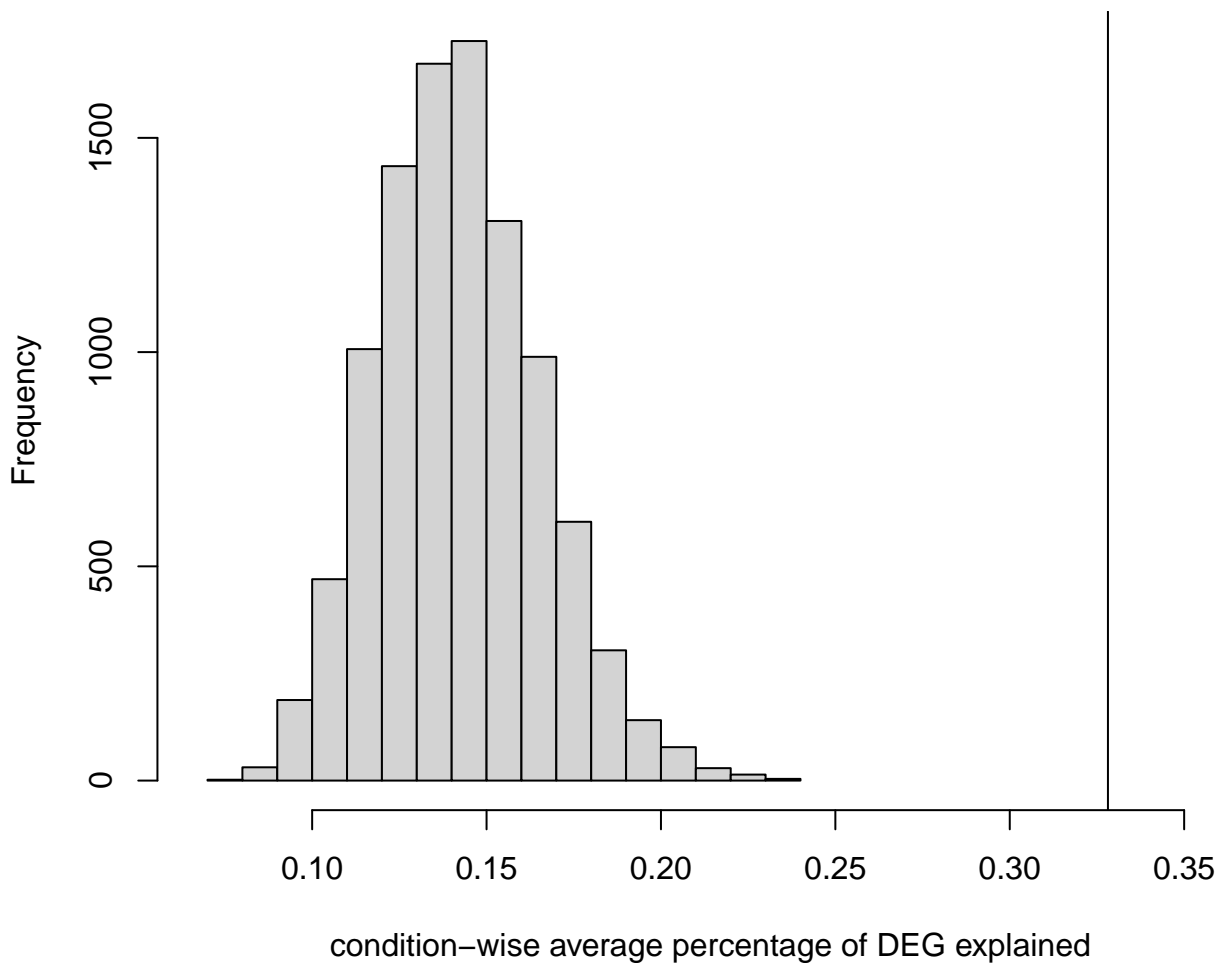
CR model – full data
 $p < 2e-04$



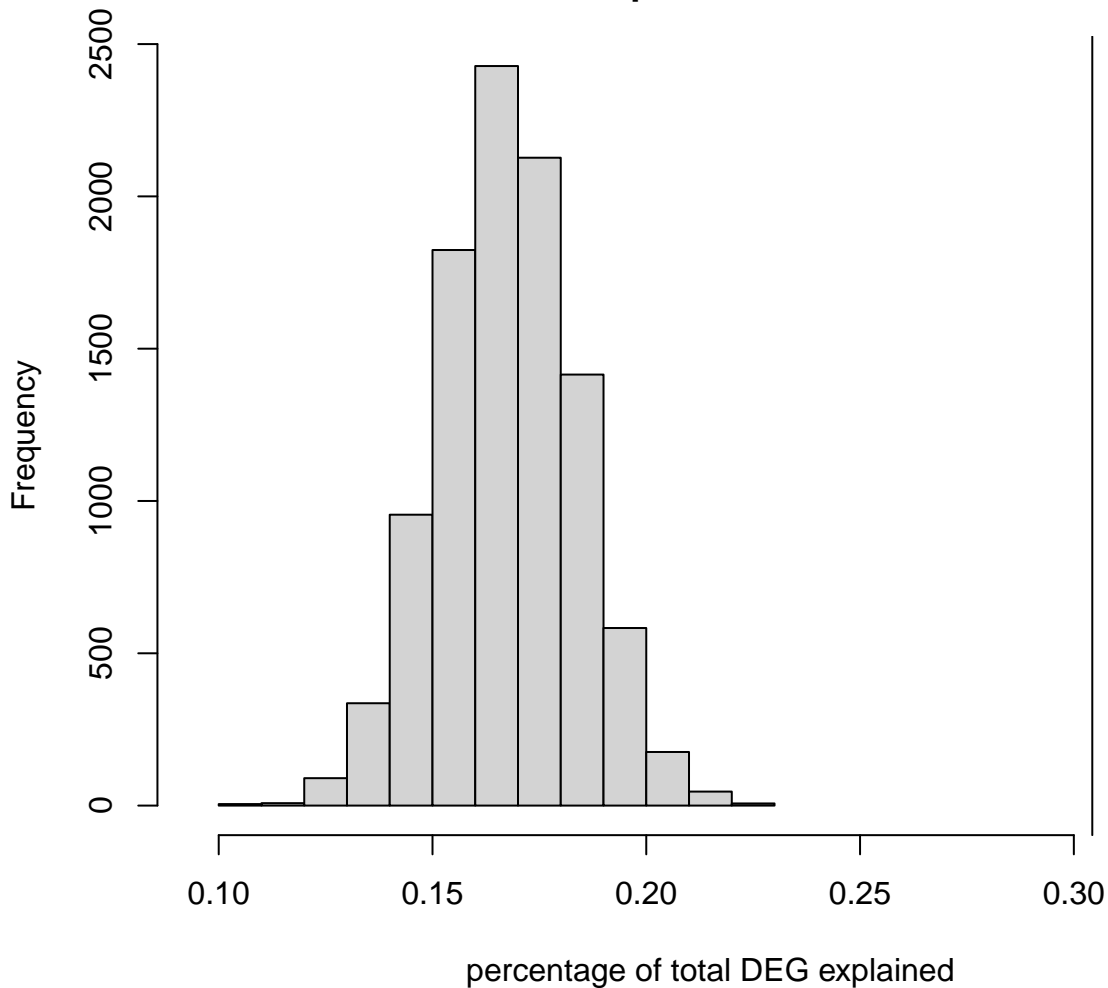
CR model – full data
p < 1e-04



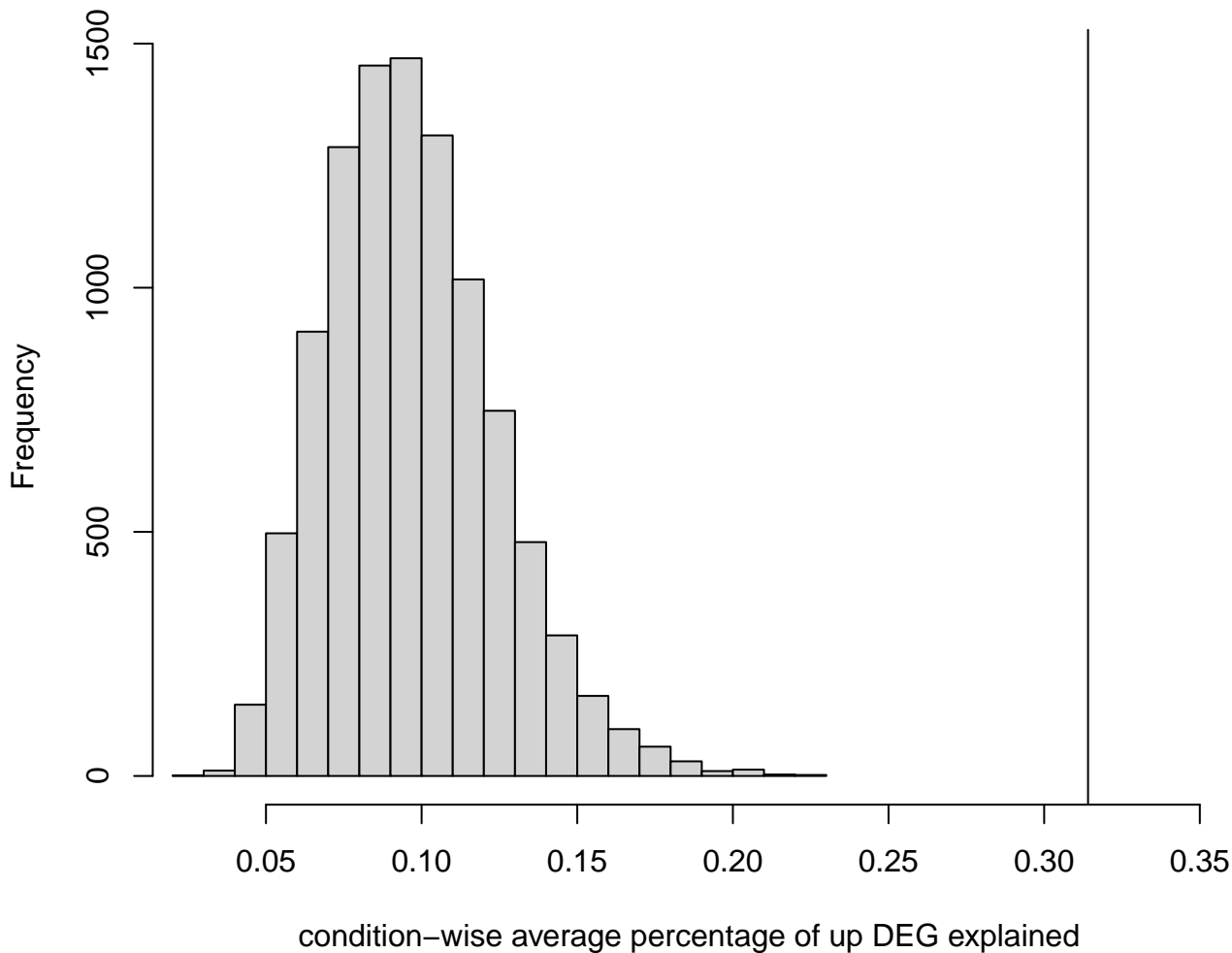
CR model – single data
 $p < 1e-04$



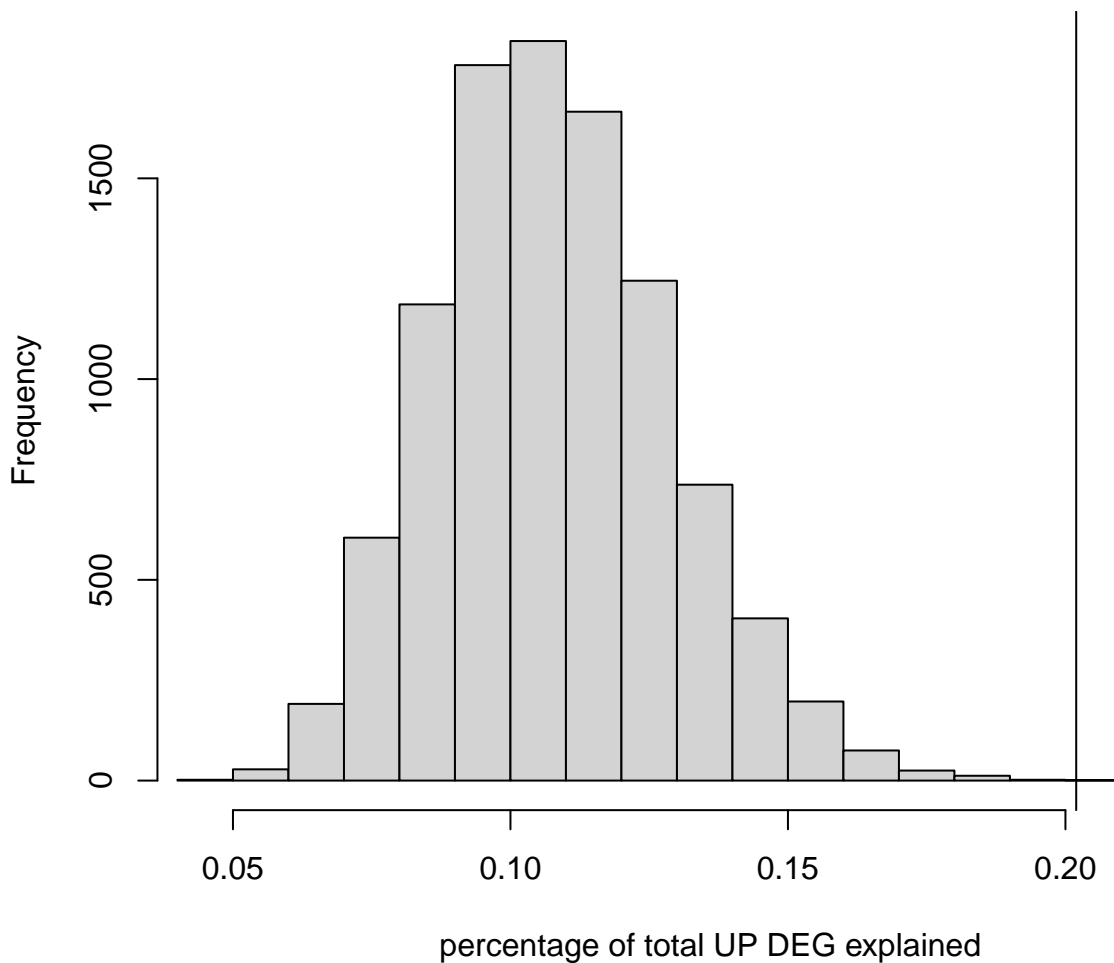
CR model – single data
p < 1e-04



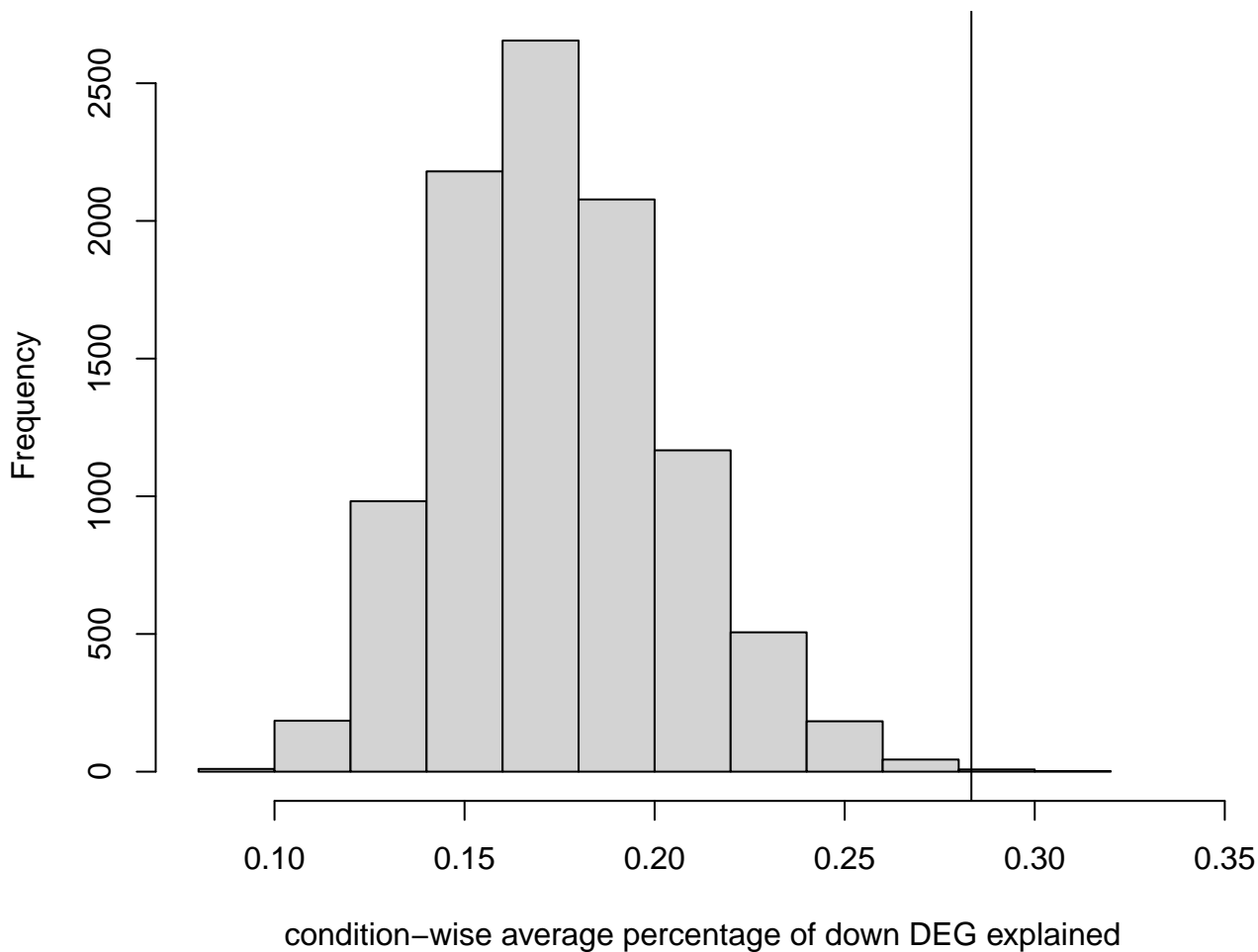
CR model – single data
 $p < 1e-04$



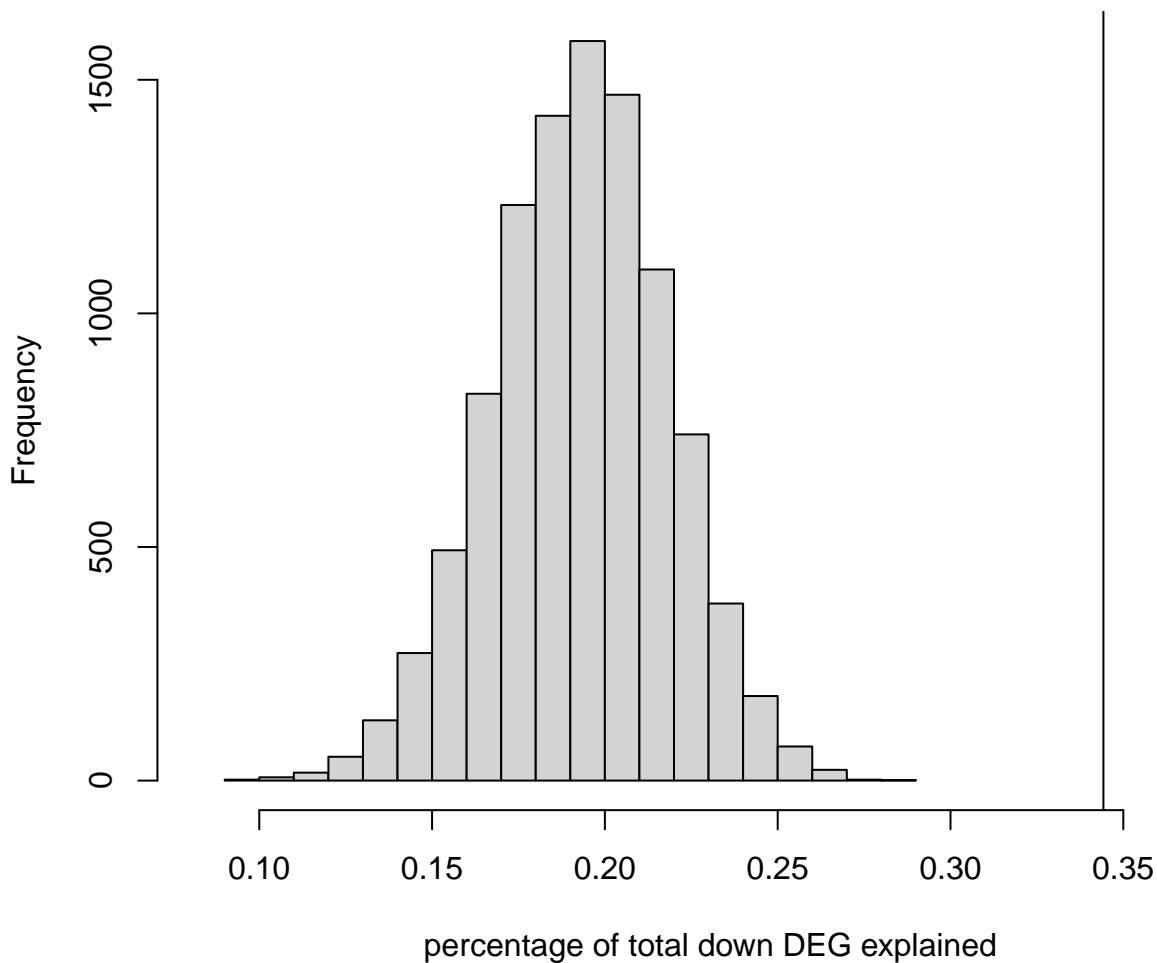
CR model – single data
 $p < 2e-04$



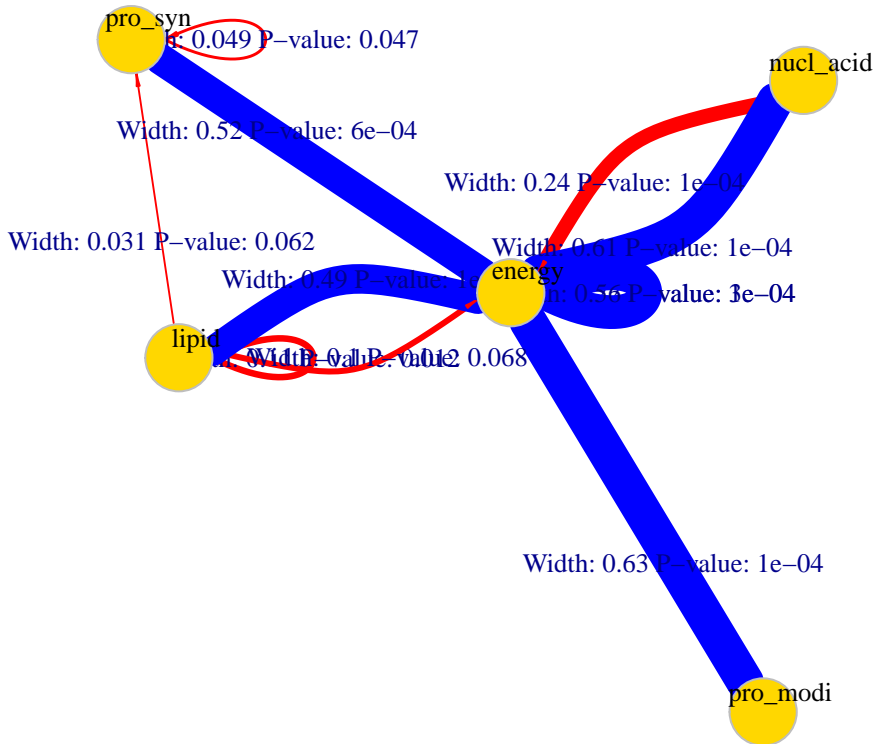
CR model – single data
p < 0.001



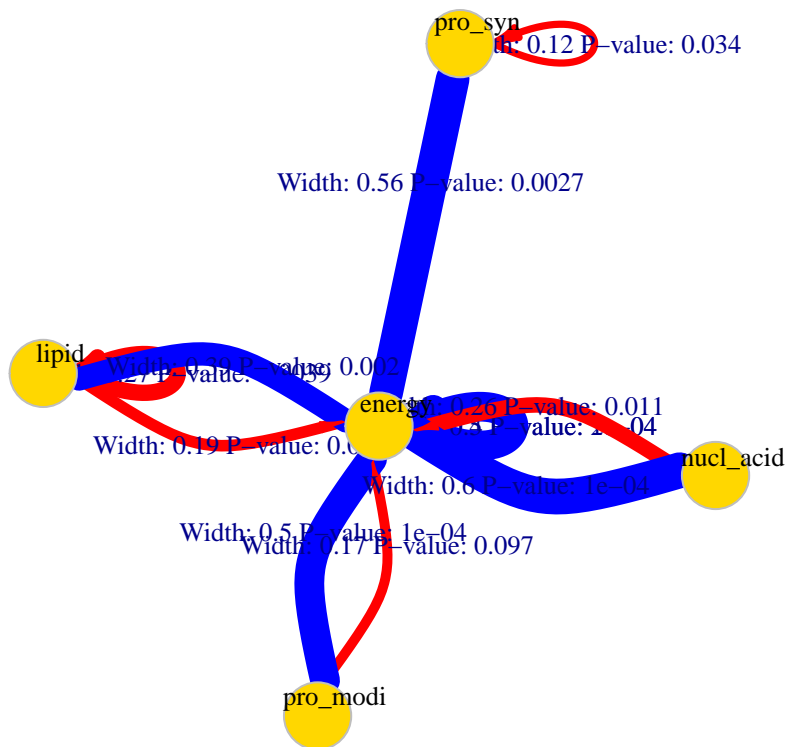
CR model – single data
 $p < 1e-04$



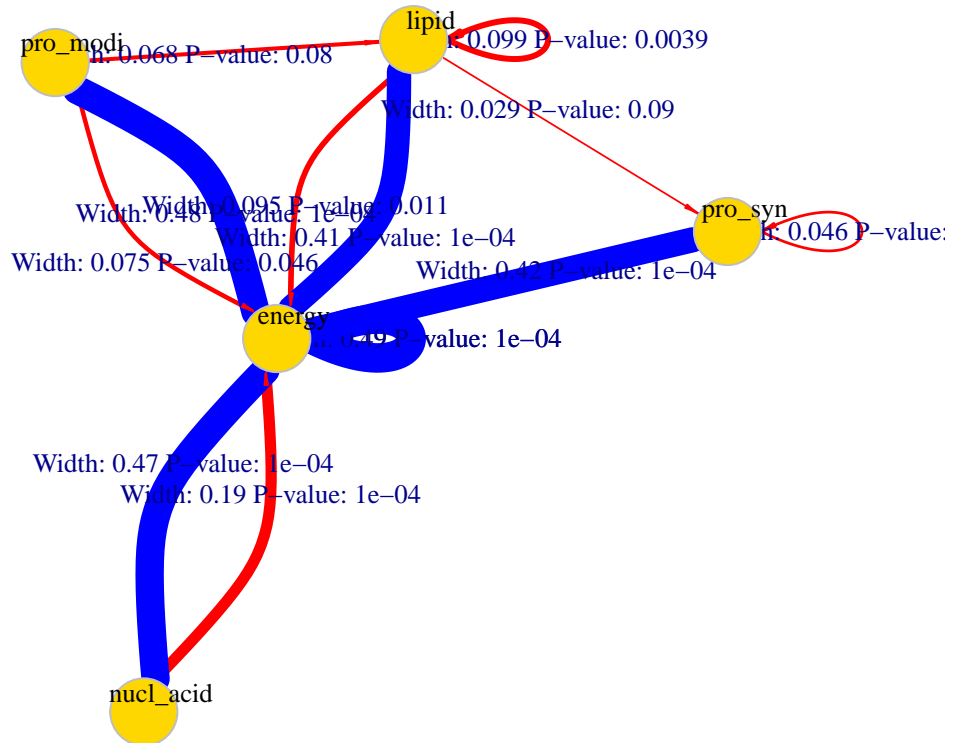
single – raw – $p < 0.2$



single – normalized – $p < 0.2$



full – raw – p < 0.2



full – normalized – $p < 0.2$

