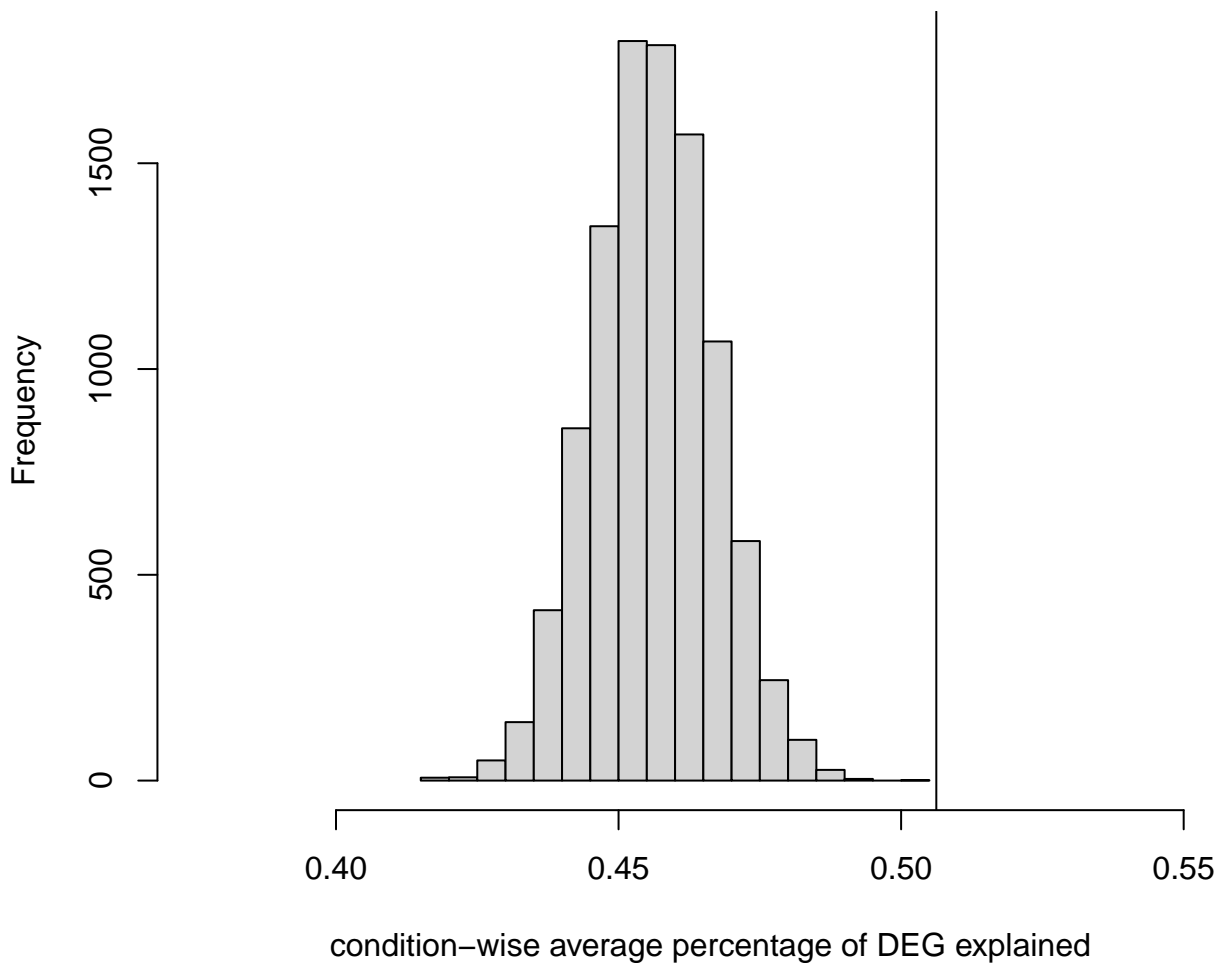
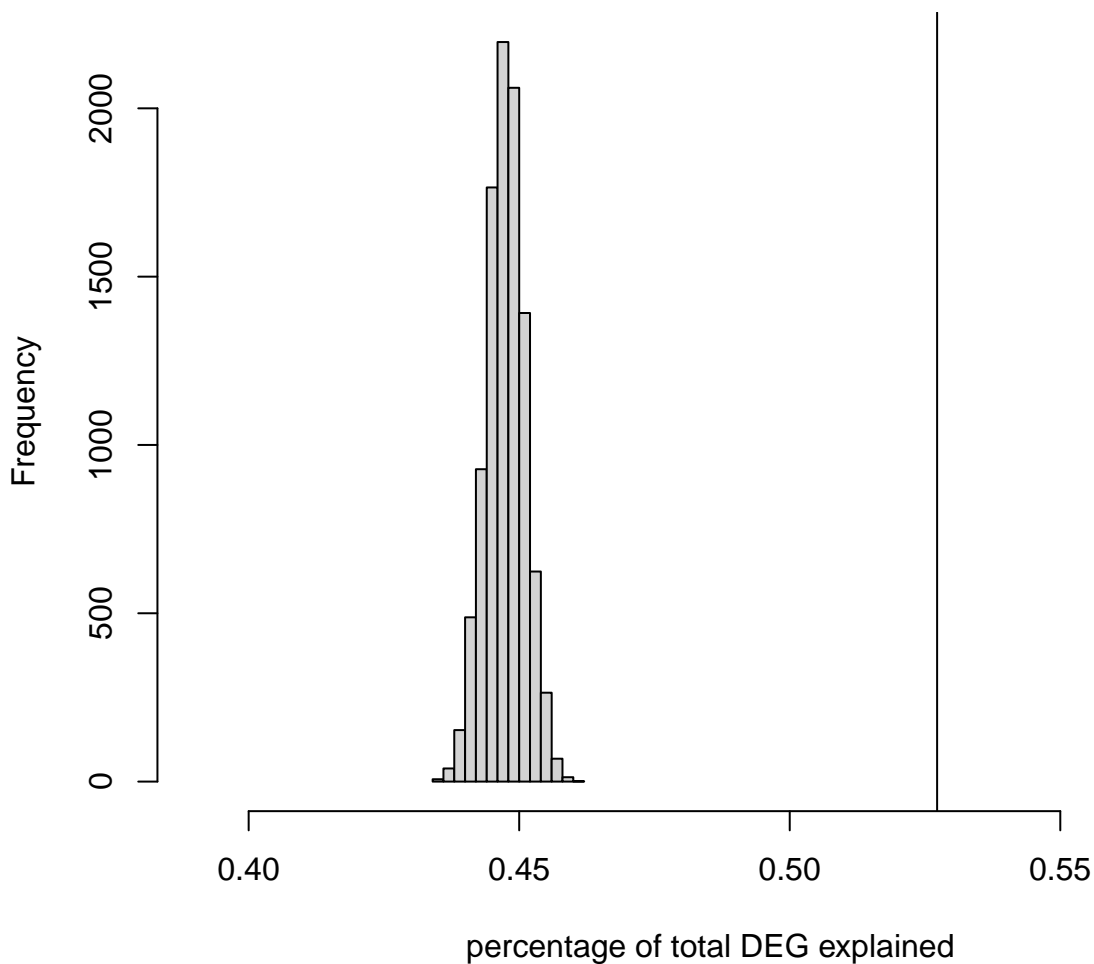


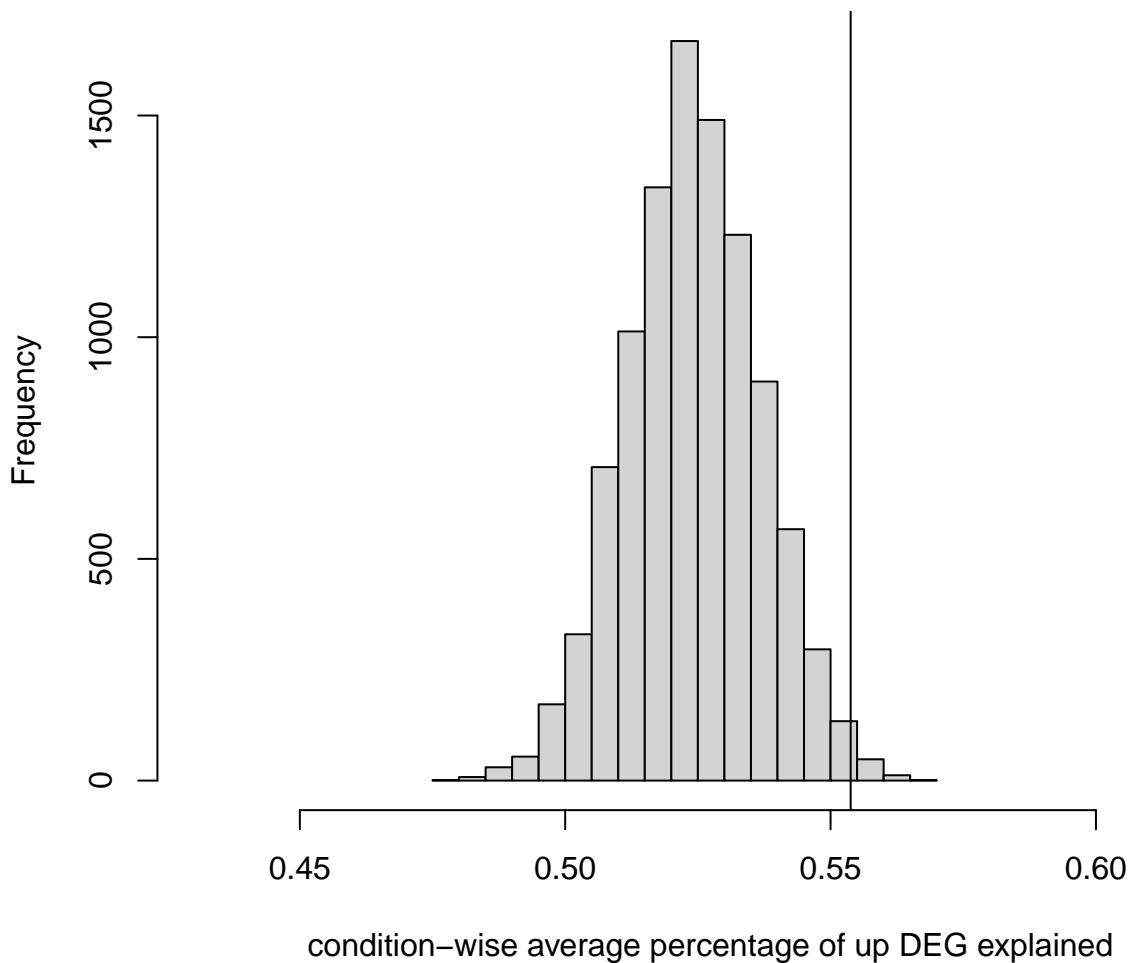
**CR model – full data**  
 **$p < 1e-04$**



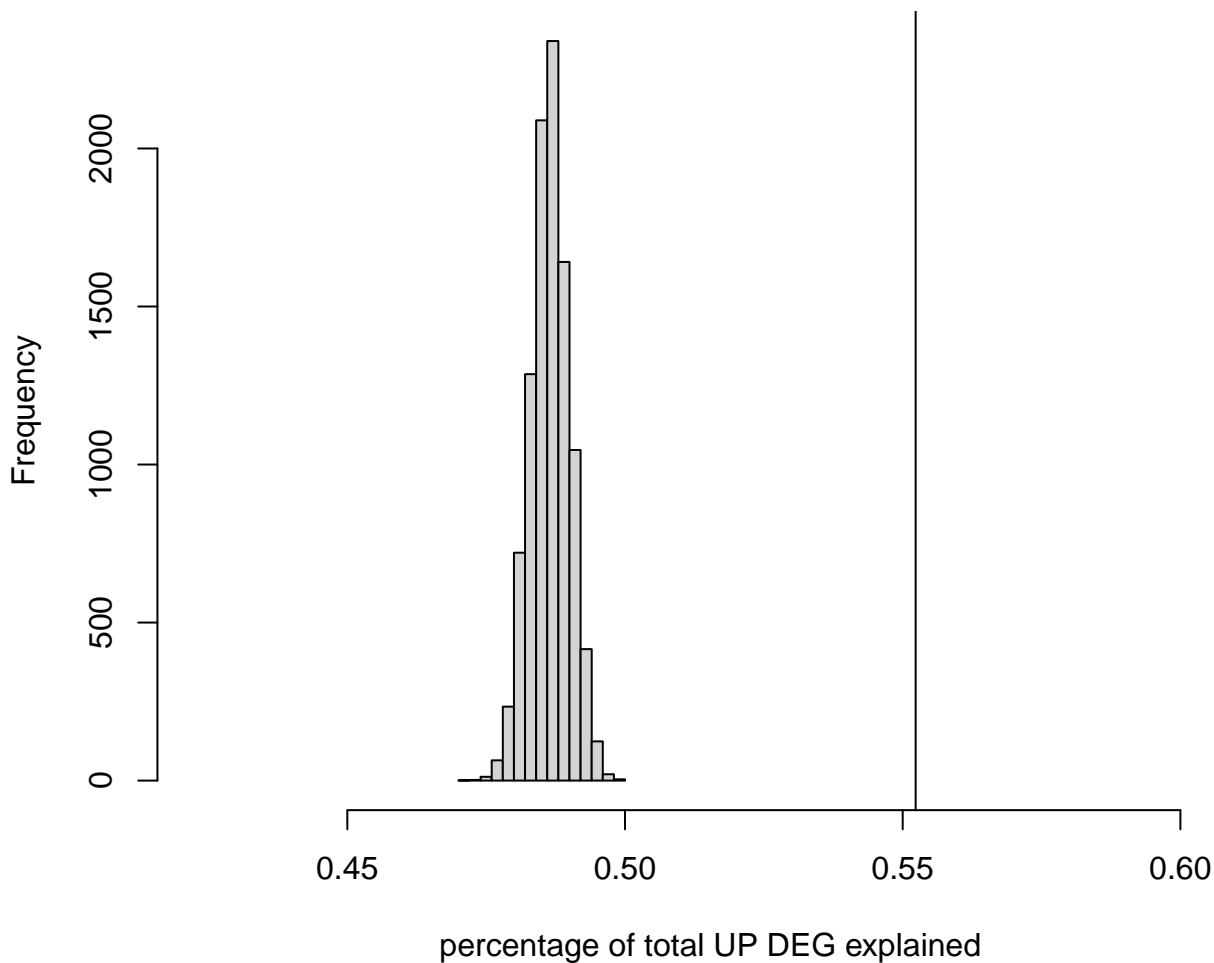
**CR model – full data**  
**p < 1e-04**



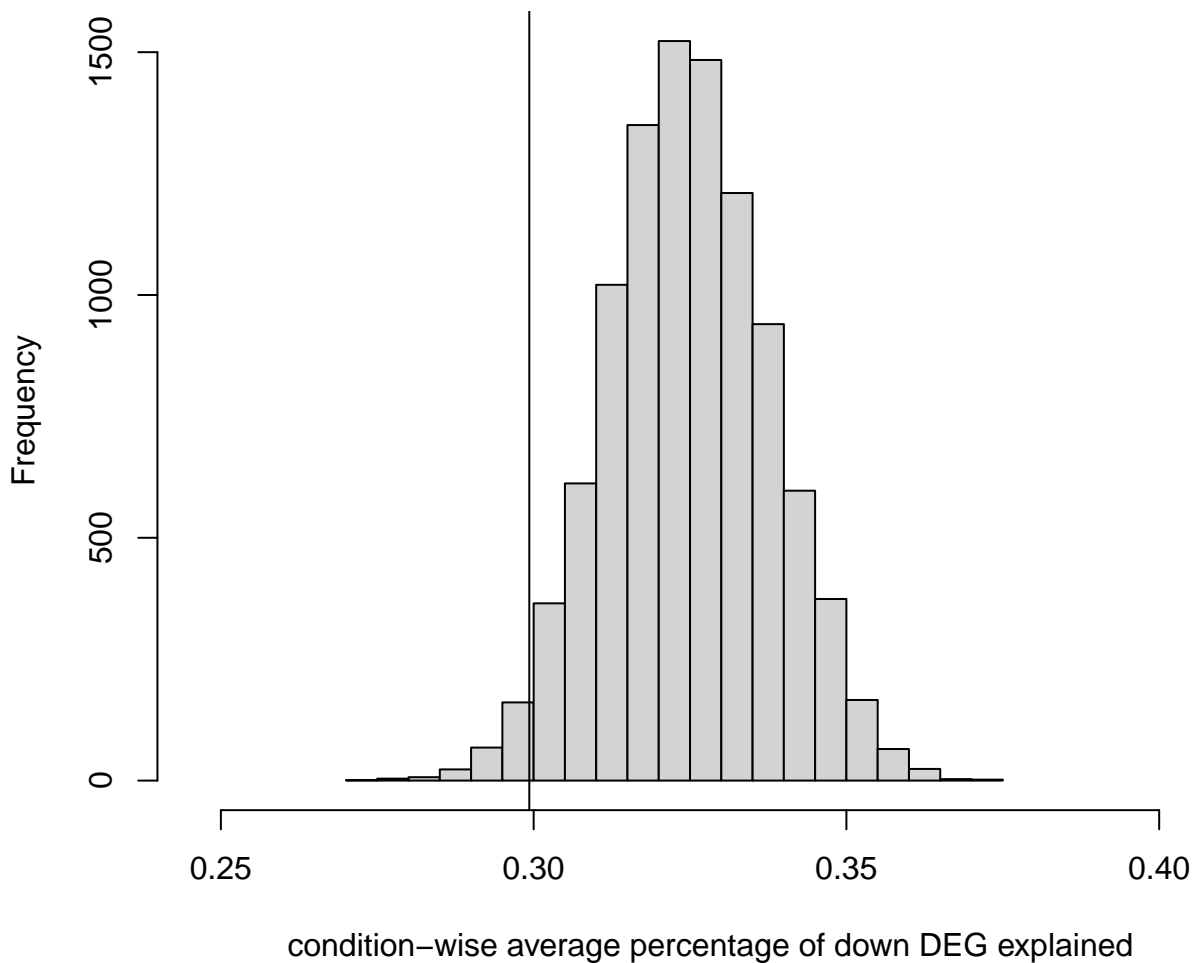
**CR model – full data**  
**p < 0.008**



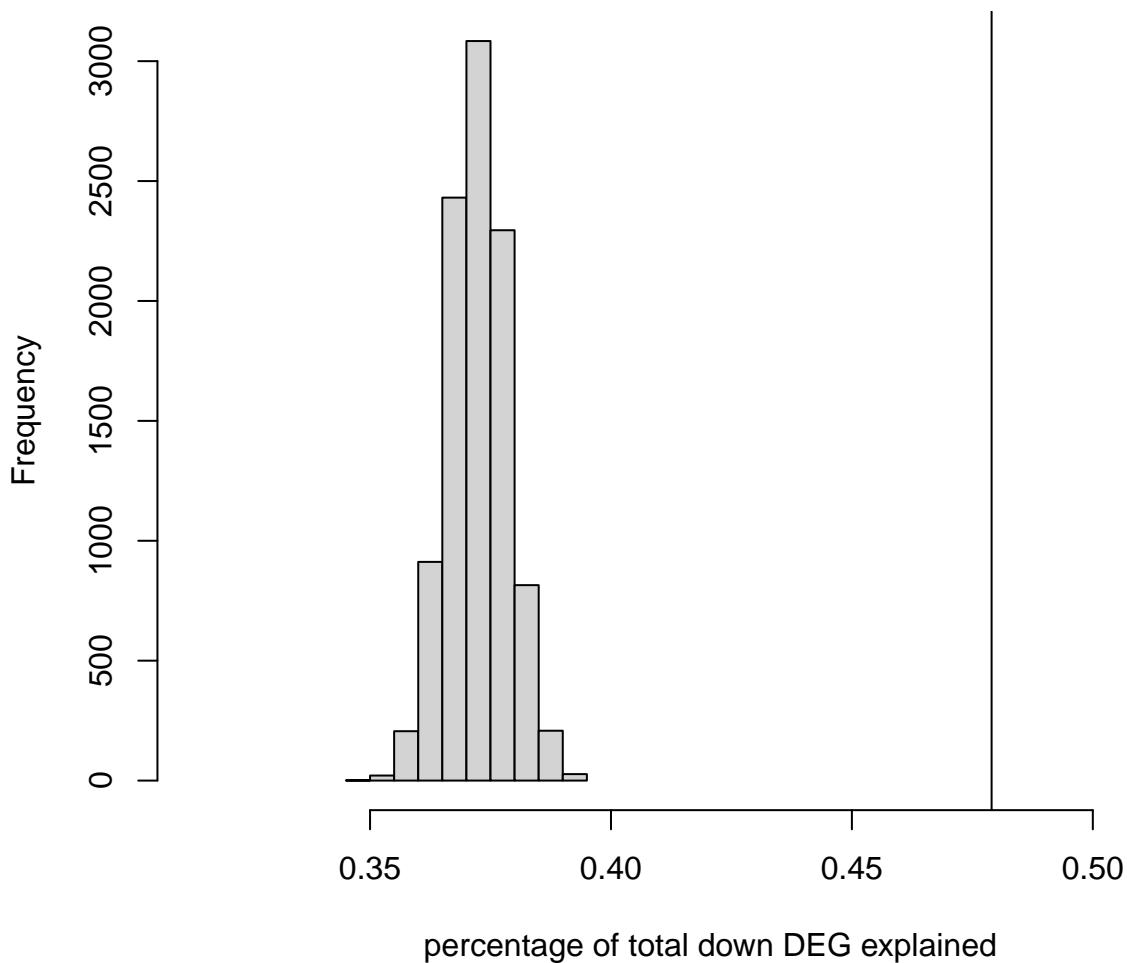
**CR model – full data**  
**p < 1e-04**



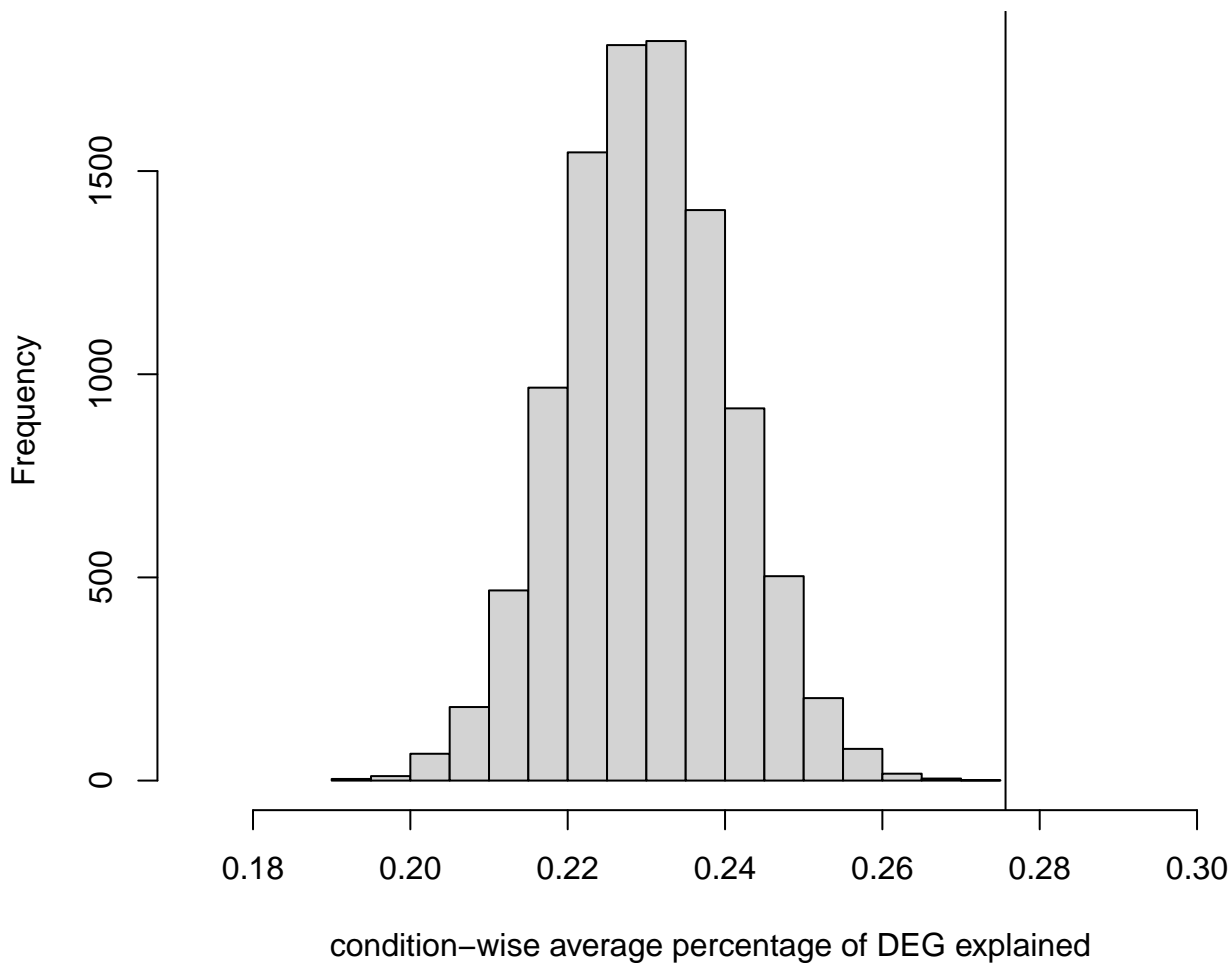
**CR model – full data**  
**p < 0.98**



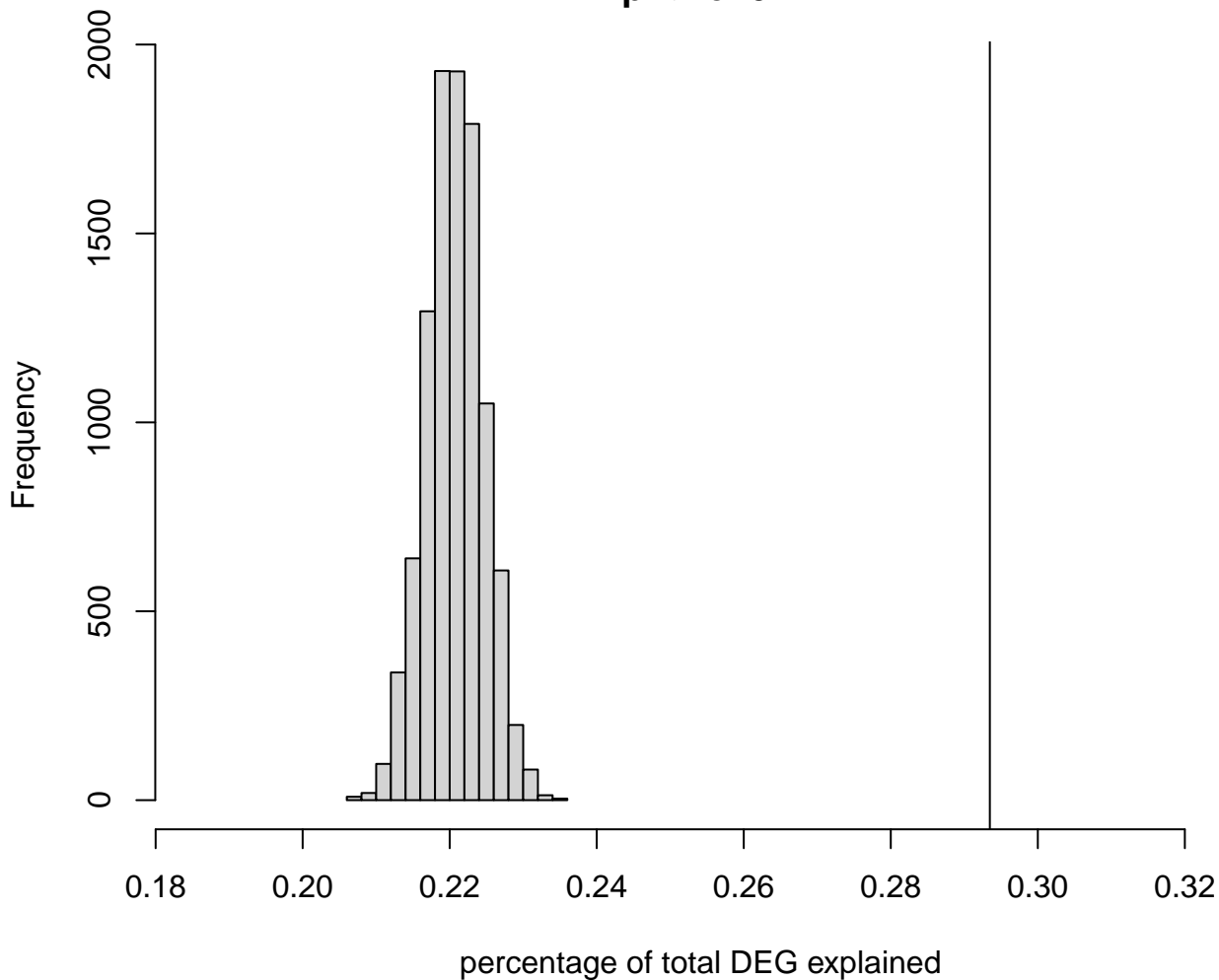
**CR model – full data**  
 **$p < 1e-04$**



**CR model – single data**  
 **$p < 1e-04$**

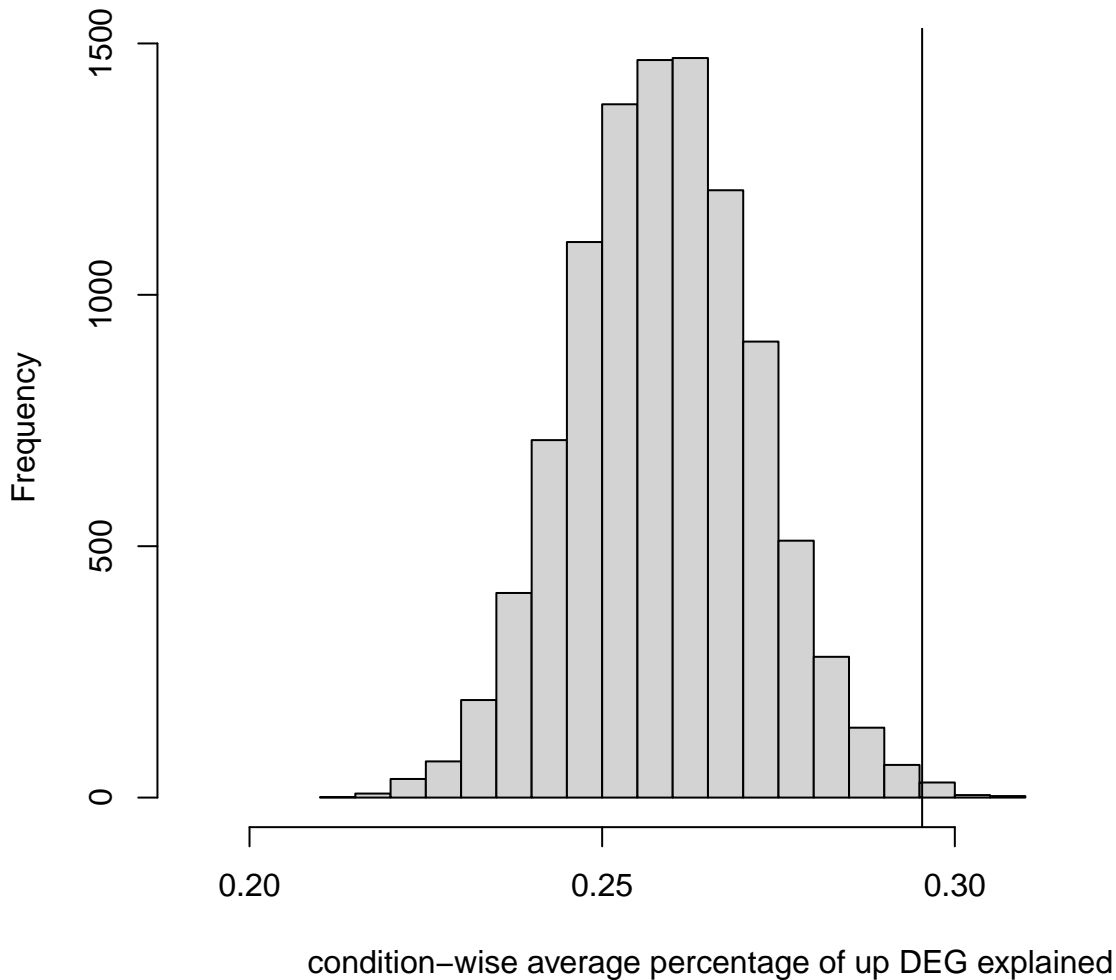


**CR model – single data**  
 **$p < 1e-04$**

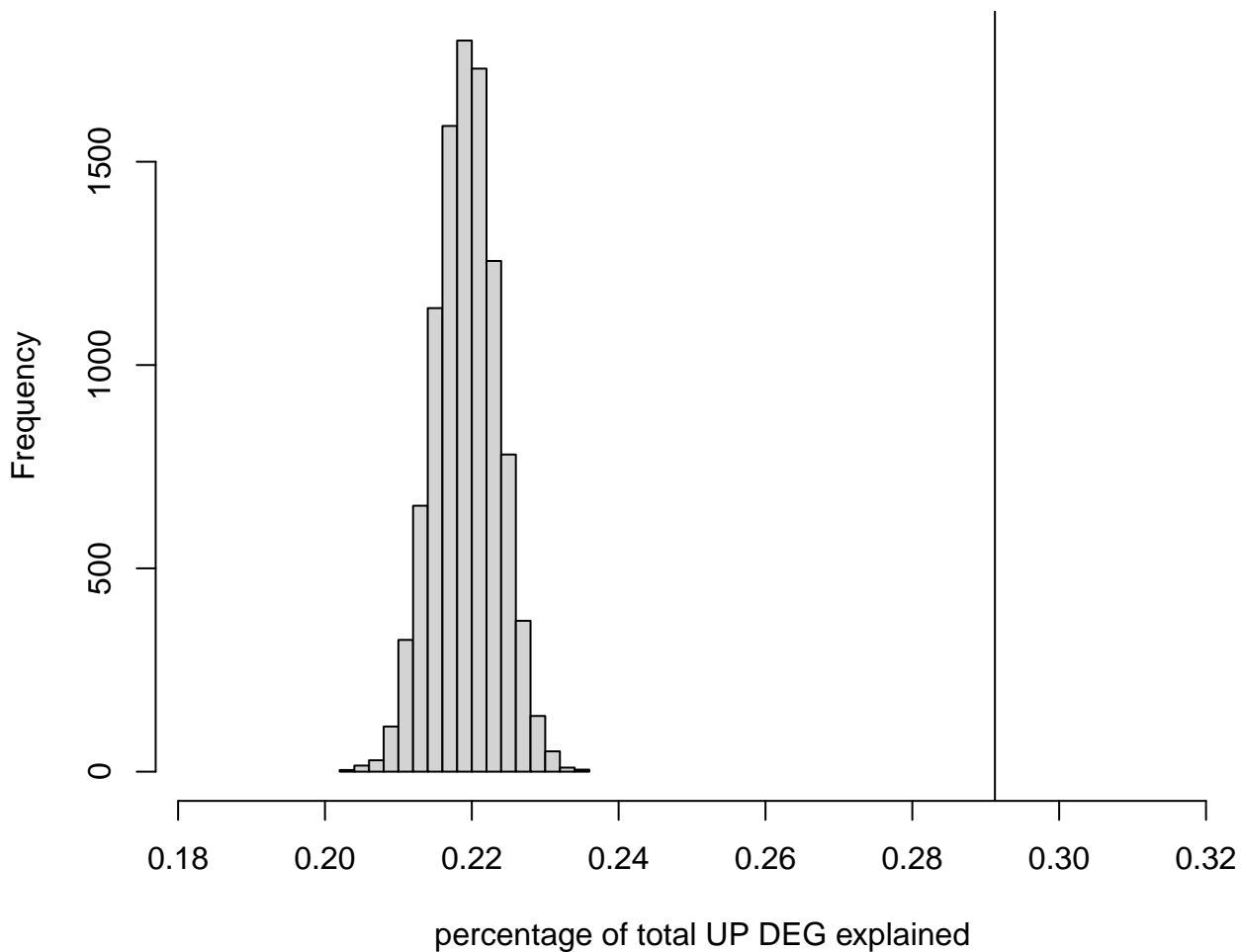




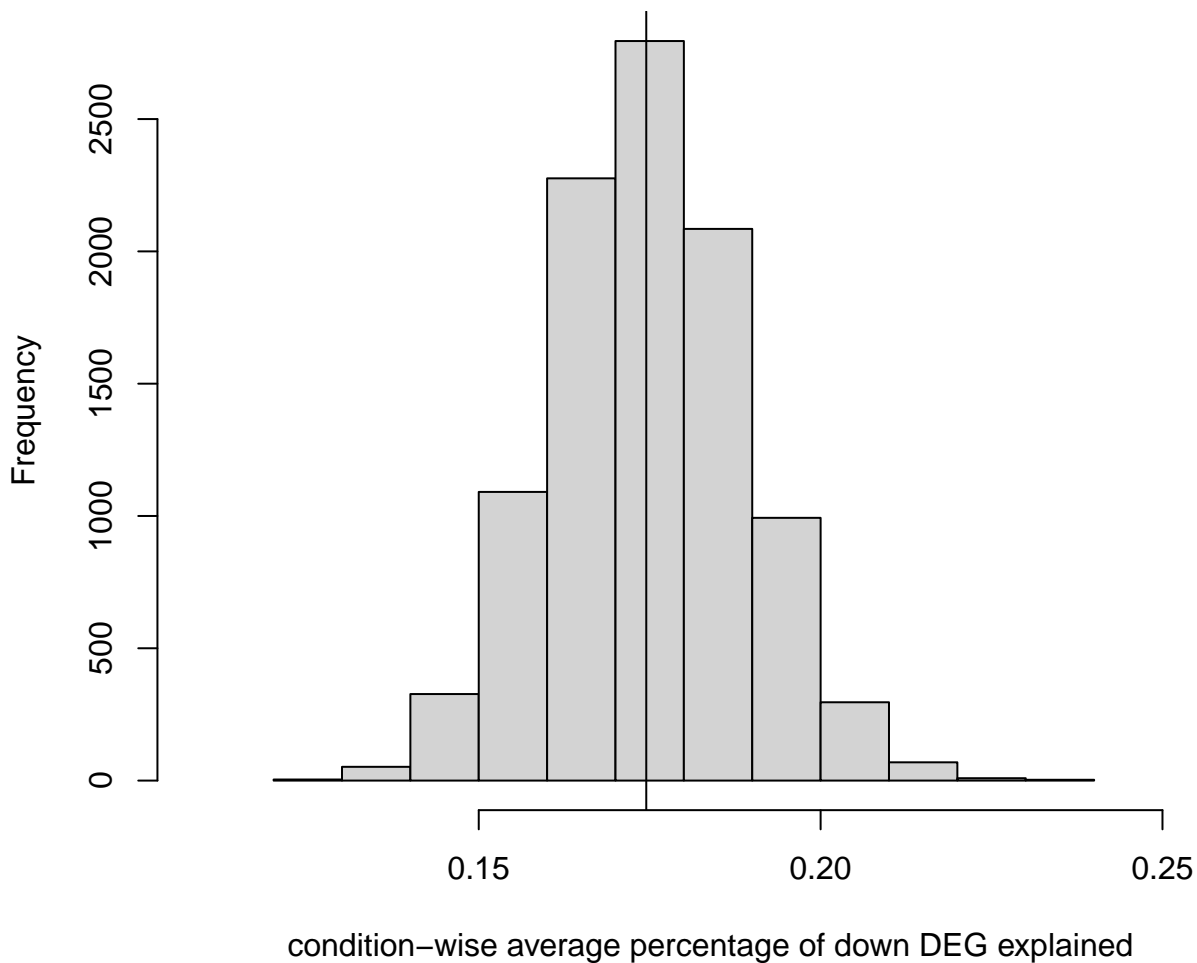
**CR model – single data**  
**p < 0.0033**



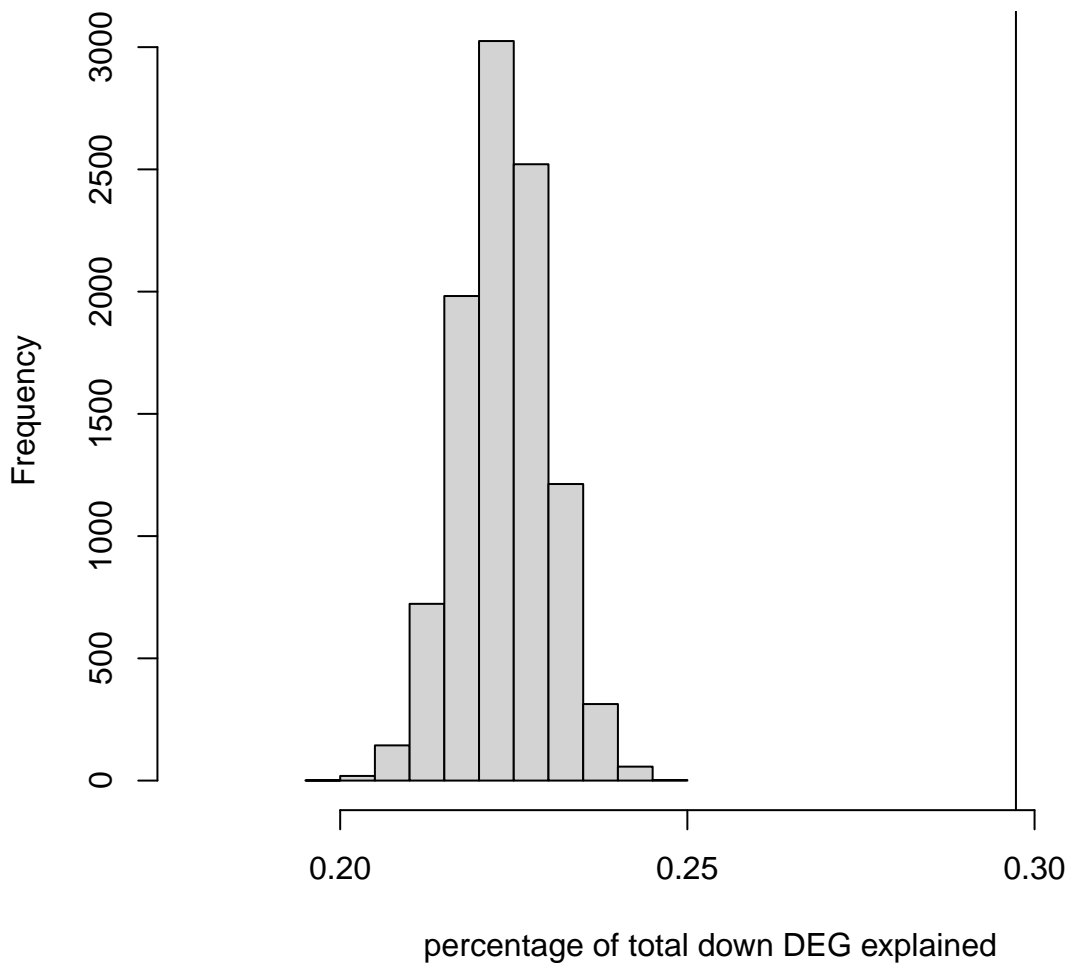
**CR model – single data**  
**p < 1e-04**



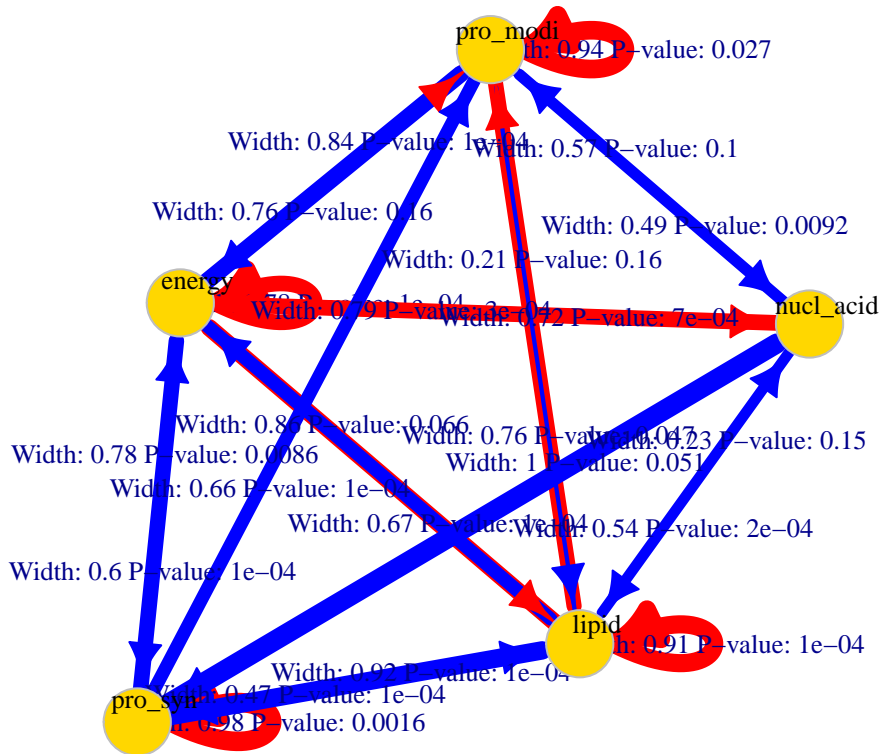
**CR model – single data**  
**p < 0.5**



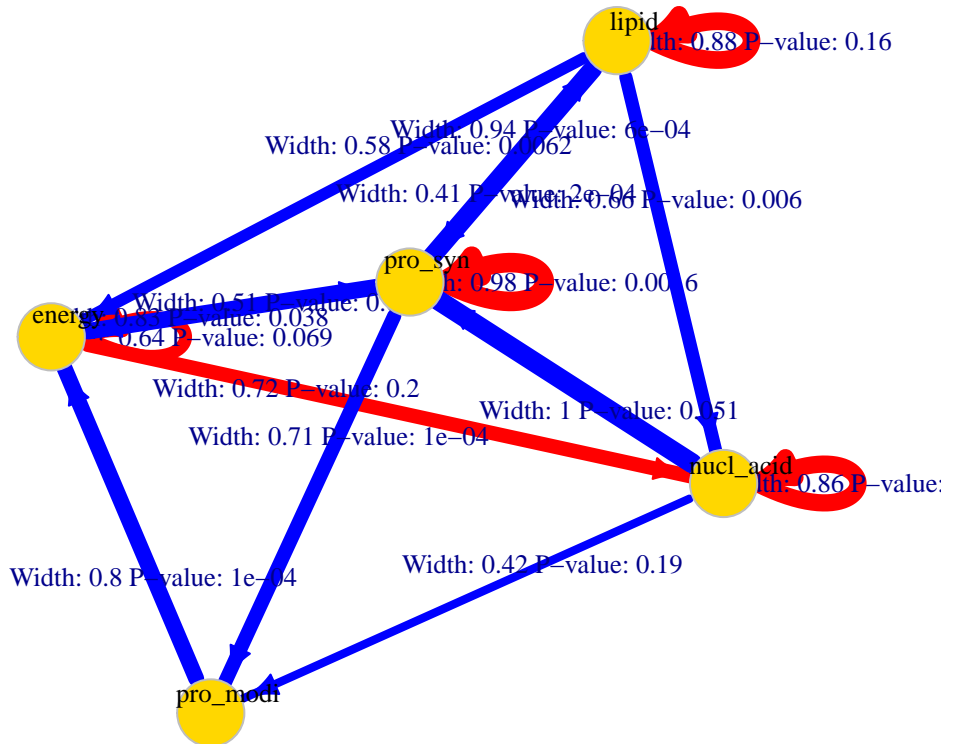
**CR model – single data**  
 **$p < 1e-04$**



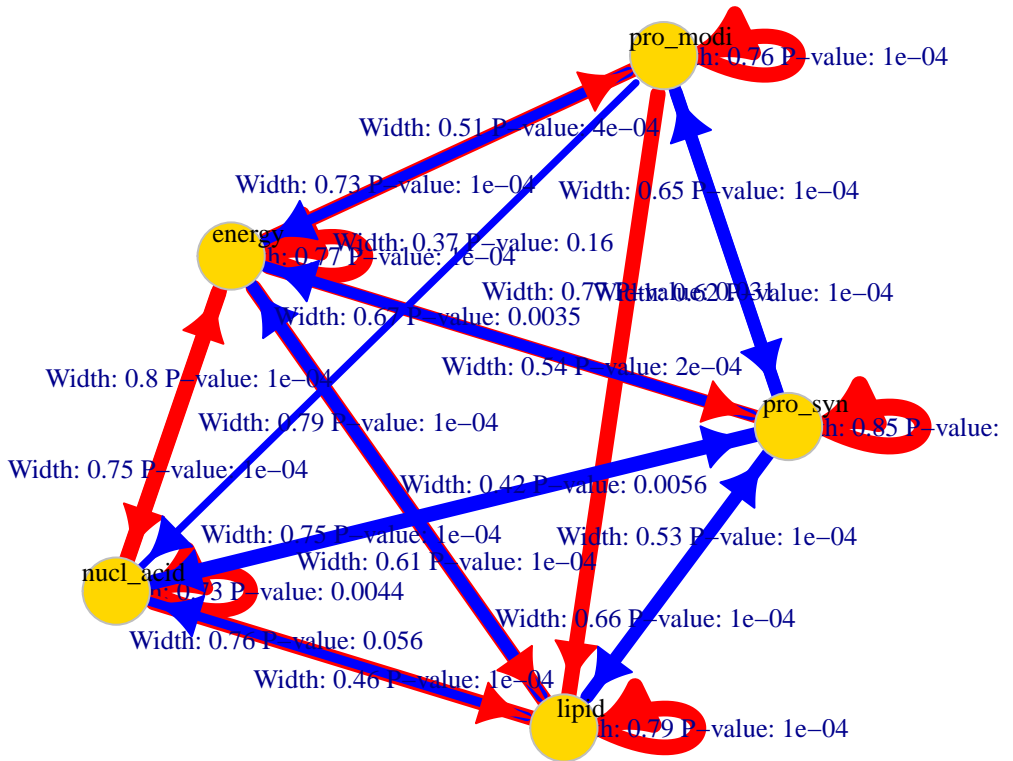
# single – raw – p < 0.2



# single – normalized – $p < 0.2$



**full – raw –  $p < 0.2$**



# full – normalized – p < 0.2

