

B110-styleguide_v0.1

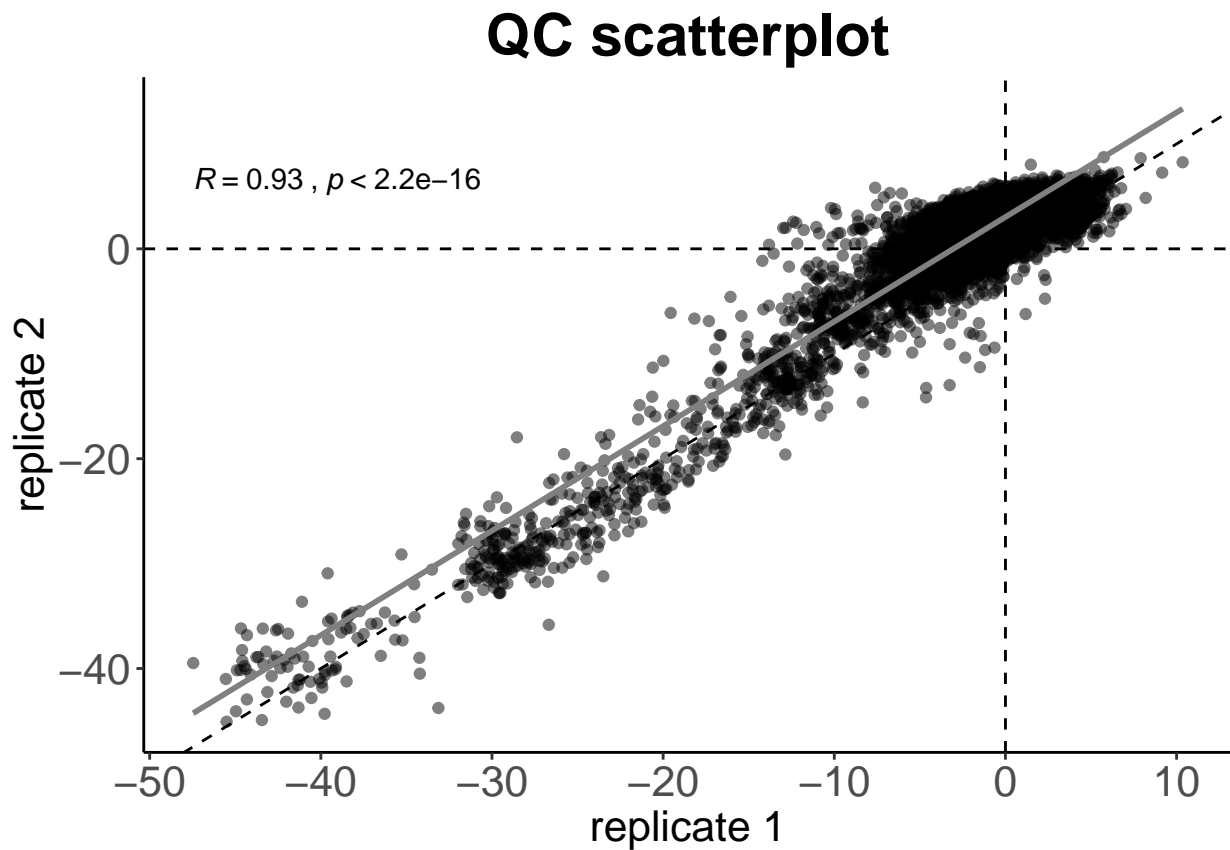
B110 Theme

B110 Colors

```
sgi_blue      = '#5087C8'  
sgi_yellow1   = '#F2EE35'  
sgi_yellow2   = '#FED98E'  
b110_grey     = '#808080'  
b110_grey_light = '#909090'  
b110_transparent_black = alpha('#000000',0.5)
```

Quality control Scatter plot

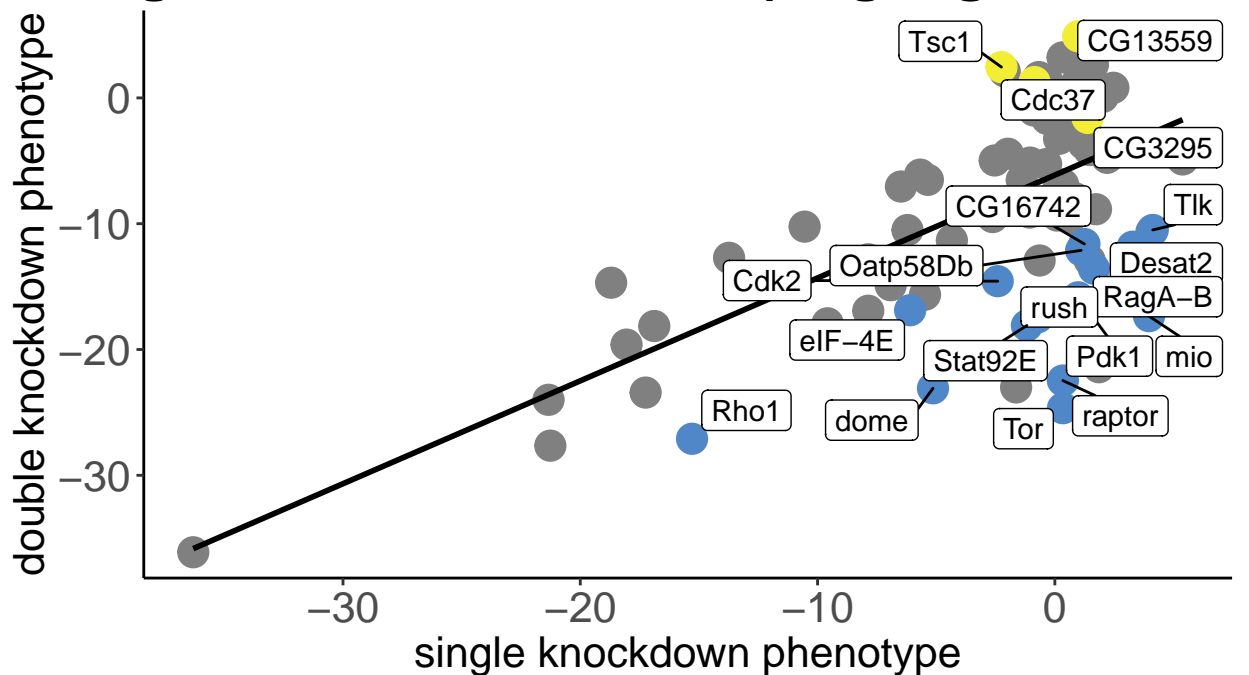
```
## Parsed with column specification:  
## cols(  
##   replicate1 = col_double(),  
##   replicate2 = col_double()  
## )
```



GI Scatter plot (single vs. double mutant)

```
## Parsed with column specification:
## cols(
##   gene1 = col_character(),
##   gene2 = col_character(),
##   doublemut = col_double(),
##   singlemut = col_double(),
##   significant = col_character(),
##   fdr = col_double()
## )
```

single vs. double mutant (target gene = Rho1)

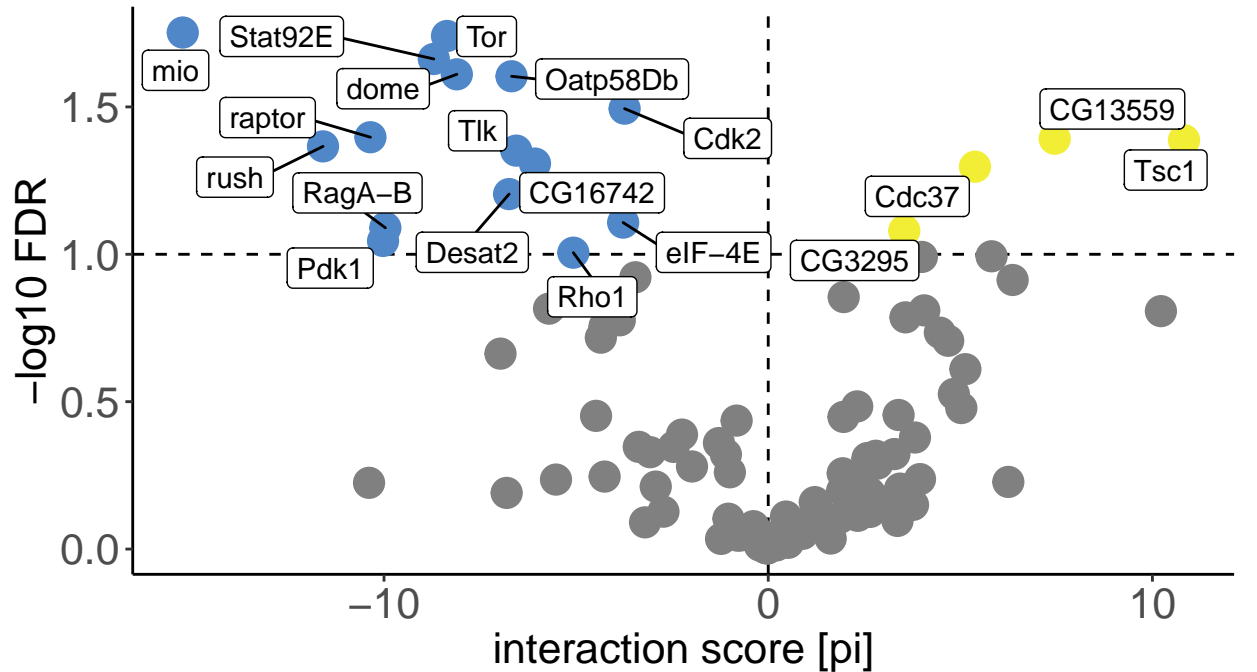


significant ● negative ● non-significant ● positive

GI Volcano plot (pi-score vs. $-\log_{10}$ FDR)

```
## Parsed with column specification:
## cols(
##   gene1 = col_character(),
##   gene2 = col_character(),
##   fdr = col_double(),
##   piscore = col_double(),
##   significant = col_character()
## )
```

GI vocalno (query gene = Rheb)

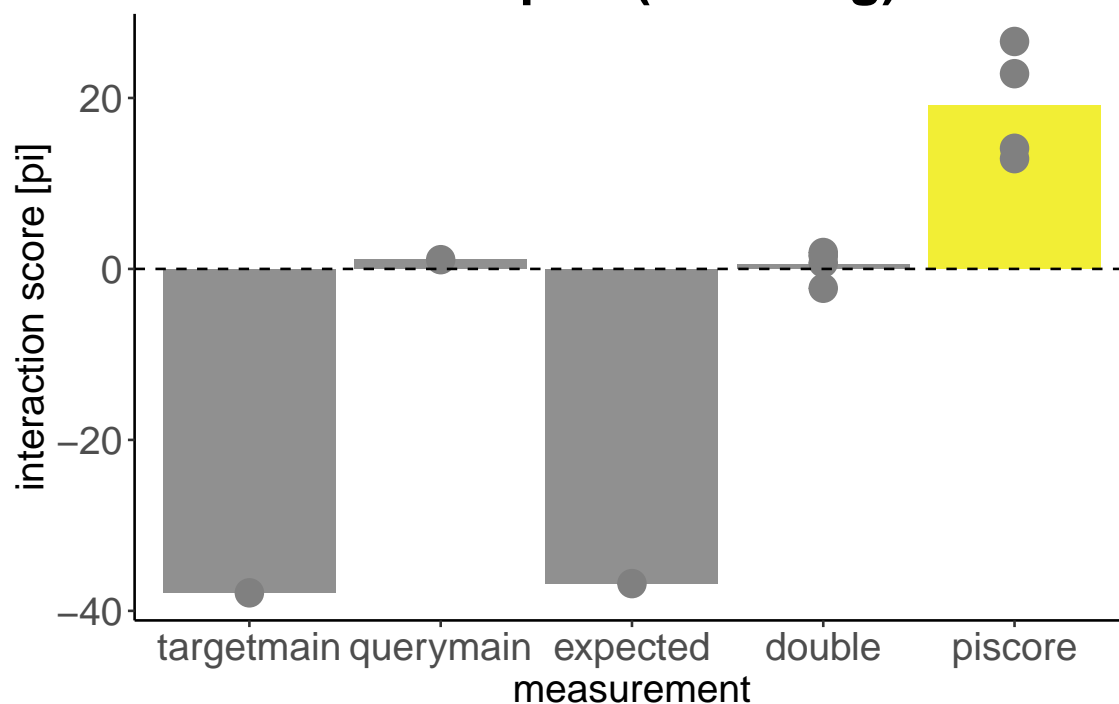


significant ● negative ● non-significant ● positive

GI bar plot (buffering vs synthetic lethal)

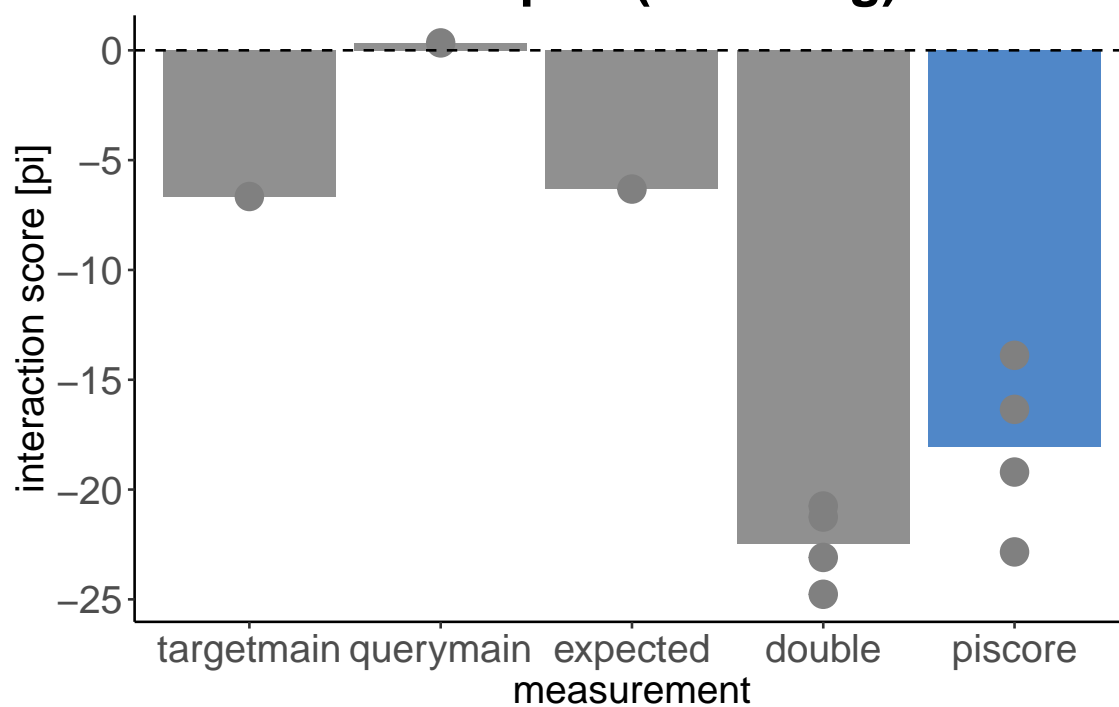
```
## Parsed with column specification:
## cols(
##   target = col_character(),
##   query = col_character(),
##   category = col_character(),
##   value = col_double()
## )
```

GI barplot (buffering)



category targetmain querymain expected double

GI barplot (buffering)



category targetmain querymain expected double