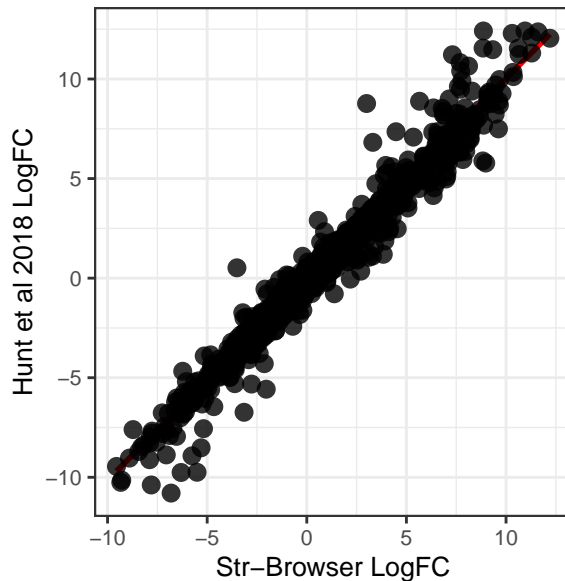


# *S. venezuelensis*: Str-Browser vs Hunt Data

Batch FLF\_PF; comparison = PF vs FLF



points = genes; red line/shading = linear regression  
w/ 95% confidence regions (formula =  $y \sim x$ ).  
Adj R-squared = 0.966