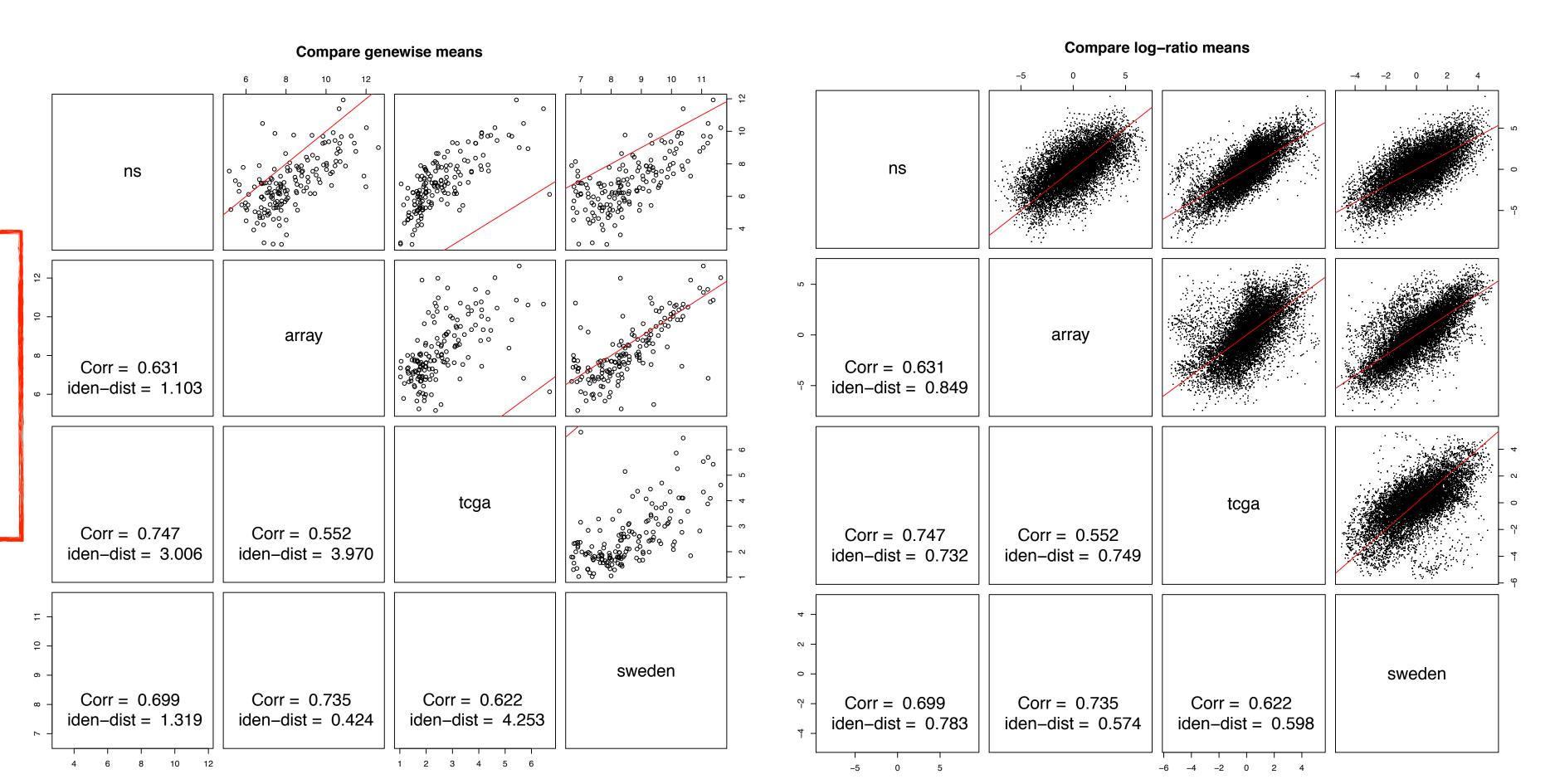
## The solution is trivial.

We have "standardised features" within every patient to build models

Log-ratio
Take the differences between log(gene A) and log(gene B)



## The solution is trivial?

Some data shifts still remain

Lasso offers feature selection but Ridge offers stability in prediction

