**Moderator analyses**

Always fit to outcome-level data.

For outcomes that **did** have variances (diff, FEest):

* rma.mv (nested random intercepts of eID within pID) with robust SEs under the CHE working model (multilevel random effects model with constant sampling correlation working model)

For outcomes that **did not have variances** ("pw.ratio", "pw.PIRepInside", "pw.PIRepInside.sens", "pw.Porig", "pw.PorigSens", "pw.PsigAgree1"):

* lmer with same specification as above and robust SEs
* so should work even though some outcomes probably have weird residuals

Scatterplot

* At outcome level, omits 2 pairs with extreme originals (SMD >80)

Need in de dataset:

* origES3, etc.
* animal vs. not animal