

Analysis

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Importing the libraries

```
##
## Attaching package: 'dplyr'

## The following objects are masked from 'package:stats':
##
##   filter, lag

## The following objects are masked from 'package:base':
##
##   intersect, setdiff, setequal, union

## Loading required package: Matrix

## Loading required package: metadat

## Loading required package: numDeriv

##
## Loading the 'metafor' package (version 4.4-0). For an
## introduction to the package please type: help(metafor)
```

Reading and Cleaning the dataframe

```
## New names:
## * `` -> `...4`
## * `` -> `...11`
## * `` -> `...20`
## * `` -> `...25`
## * `` -> `...26`
## * `` -> `...34`
## * `` -> `...36`

## [1] "Authors"
## [2] "Year of publication"
## [3] "Origin Country of study"
## [4] "Total sample size"
## [5] "Sample size included"
## [6] "Sex (%female)"
## [7] "Ethnic background (% white)"
## [8] "age range"
## [9] "mean age"
## [10] "Task design (Correlational/Group comparisons/Experimental)"
## [11] "Task"
## [12] "Manipulation"
```

```
## [13] "Trait or phase (state)"
## [14] "Clinical sample"
## [15] "Control Group (yes/no)"
## [16] "Social anxiety measure trait"
## [17] "Social anxiety measure state"
## [18] "Metacognition trait or state"
## [19] "Metacognition measure"
## [20] "Dimensions used (if looked at different dimension make separate rows for each dimension)"
## [21] "Number of items"
## [22] "Statistical anlysis"
## [23] "Relationship between social anxiety and metacognition"
## [24] "d or R"
## [25] "effect size"
## [26] "V"
## [27] "95% CI"
## [28] "Major finding"
## [29] "Risk of bias (selection bias, attrition bias, detection bias and reporting bias)"
```

Select only the columns you're interested in:

Change the names of the columns:

```
## [1] "Authors"                "Sample_size_included"
## [3] "Sex_(%female)"          "Ethnic_background_(%_white)"
## [5] "age_range"              "d_or_R"
## [7] "effect_size"            "V"
## [9] "95%_CI"
```

Select only the effect size available (r or d)

Remove all the rows that present a NA value inside the variance variable

Convert the percentages into prevalence value between 0 and 1:

For the variables ethnic and female, convert some variables from “-” to NA and then convert all the percentages variables

Checking if Sex is numeric:

```
## [1] TRUE
```

Converting the wrong value of effect size (we can see that the correct has the point from the confidence interval)

```
## [1] "1177"
```

```
## [1] "1.177"
```

```
## tibble [43 x 10] (S3: tbl_df/tbl/data.frame)
## $ Authors                : chr [1:43] "Yu, Meng and Lv, Fangyan and Liu, Zicheng and Gao, Dingg
## $ Sample_size_included    : num [1:43] 357 33 40 169 169 169 169 169 62 ...
## $ Sex_(%female)           : num [1:43] 0.75 0.558 0.575 0.787 0.787 0.787 0.787 0.787 0.70
## $ Ethnic_background_(%_white): num [1:43] 0 0 0.9 NA NA NA NA NA NA ...
## $ age_range               : chr [1:43] "11 till 17" "13 - 19" "-" "18 - 24 (most people 111) but
## $ d_or_R                  : Factor w/ 2 levels "d","r": 2 1 1 2 2 2 2 2 2 ...
## $ effect_size             : num [1:43] 0.19 5.22 -0.119 0.587 0.333 0.54 0.338 0.229 0.349 0.52
## $ V                       : num [1:43] 0.004 0.53 0.1 0.006 0.006 0.006 0.006 0.006 0.017
## $ 95%_CI                  : chr [1:43] "[0,0694, 0,315]" "[3.791, 6.658]" "[-0,738, 0,502]" "[0
## $ std                      : num [1:43] 0.0632 0.728 0.3162 0.0775 0.0775 ...
```

handling the CI

Convert the CI that present 3 commas (error) into only 1 (the second)

```
## [1] "[0.0694, 0.315]" "[3.791, 6.658]" "[-0.738, 0.502]" "[0.478, 0.678]"
## [5] "[0.192, 0.461]" "[0.424, 0.639]" "[0.197, 0.465]" "[0.080, 0.367]"
## [9] "[0.209, 0.475]" "[0.311, 0.681]" "[0.018, 0.206]" "[0.413, 0.662]"
## [13] "[0.308, 0.589]" "[0.449, 0.687]" "[0.141, 0.462]" "[0.285, 0.572]"
## [17] "[0.024, 0.364]" "[-0.070, 0.290]" "[0.097, 0.427]" "[0.332, 0.517]"
## [21] "[0.375, 0.552]" "[-0.888, -0.105]" "[-0.581, 0.567]" "[-1.395, -0.088]"
## [25] "[-0.797, 0.214]" "[-0.439, 0.813]" "[-0.262, 0.988]" "[-0.524, 0.717]"
## [29] "[-0.362, 0.882]" "[0.505, 1.848]" "[-0.107, 0.247]" "[0.203, 0.362]"
## [33] "[-0.107, 0.248]" "[0.057, 0.412]" "[-0.259, 0.385]" "[0.471, 1.142]"
## [37] "[-0.147, 0.499]" "[-0.105, 0.985]" "[2.978, 4.681]" "[0.369, 0.487]"
## [41] "[0.782, 0.936]" "[0.048, 0.275]" "[0.373, 0.622]"
```

Define a function to separate the two values of the confidence interval:

```
## lower_CI upper_CI
## 1 0.0694 0.315
## 2 3.7910 6.658
## 3 -0.7380 0.502
## 4 0.4780 0.678
## 5 0.1920 0.461
## 6 0.4240 0.639
## 7 0.1970 0.465
## 8 0.0800 0.367
## 9 0.2090 0.475
## 10 0.3110 0.681
## 11 0.0180 0.206
## 12 0.4130 0.662
## 13 0.3080 0.589
## 14 0.4490 0.687
## 15 0.1410 0.462
## 16 0.2850 0.572
## 17 0.0240 0.364
## 18 -0.0700 0.290
## 19 0.0970 0.427
## 20 0.3320 0.517
## 21 0.3750 0.552
## 22 -0.8880 -0.105
## 23 -0.5810 0.567
## 24 -1.3950 -0.088
## 25 -0.7970 0.214
## 26 -0.4390 0.813
## 27 -0.2620 0.988
## 28 -0.5240 0.717
## 29 -0.3620 0.882
## 30 0.5050 1.848
## 31 -0.1070 0.247
## 32 0.2030 0.362
## 33 -0.1070 0.248
## 34 0.0570 0.412
## 35 -0.2590 0.385
## 36 0.4710 1.142
## 37 -0.1470 0.499
```

```
## 38 -0.1050    0.985
## 39  2.9780    4.681
## 40  0.3690    0.487
## 41  0.7820    0.936
## 42  0.0480    0.275
## 43  0.3730    0.622
```

Using the formula for calculating the SE from the CI, but

Using the formula for calculating the SE from the CI for both lower and upper values:

```
## [1] 0.06153174 0.72909503 0.31582213 0.05561327 0.07194010 0.05918476
## [7] 0.07194010 0.07602181 0.07142988 0.10663461 0.04847028 0.06989924
## [13] 0.07755245 0.06683796 0.08622607 0.07908309 0.08979757 0.09183842
## [19] 0.08826693 0.04949071 0.04744985 0.12653294 0.29133188 0.39031329
## [25] 0.19745261 0.32143448 0.31888341 0.31327106 0.31735277 0.34286344
## [31] 0.09030778 -0.00153064 0.09030778 0.08826693 0.16275809 0.16786023
## [37] 0.16173767 0.27806633 0.43470187 0.03112302 0.05000092 0.05714391
## [43] 0.04438857

## [1] 0.06377668 0.73368695 0.31684256 0.04642942 0.06530732 0.05051113
## [7] 0.06479711 0.07040946 0.06428690 0.08214437 0.04744985 0.05714391
## [13] 0.06581754 0.05459284 0.07755245 0.06734818 0.08367501 0.09183842
## [19] 0.08010351 0.04489878 0.04285793 0.27296420 0.29439316 0.27653569
## [25] 0.31837320 0.31735277 0.31888341 0.31990384 0.31735277 0.34235323
## [31] 0.09030778 0.08265458 0.09081800 0.09285885 0.16581937 0.17449300
## [37] 0.16786023 0.27806633 0.43419165 0.02908217 0.02857195 0.05867455
## [43] 0.08265458
```

Comparing it to the standard deviation of the estimate is identical: therefore you used the sd for calculating the CI, but this is indeed wrong!

```
## [1] 0.06324555 0.72801099 0.31622777 0.07745967 0.07745967 0.07745967
## [7] 0.07745967 0.07745967 0.07745967 0.13038405 0.04472136 0.08944272
## [13] 0.08944272 0.08944272 0.08944272 0.08944272 0.08944272 0.08944272
## [19] 0.08944272 0.05477226 0.05477226 0.33316662 0.33316662 0.33316662
## [25] 0.33316662 0.31780497 0.31937439 0.31622777 0.31780497 0.34205263
## [31] 0.08944272 0.08944272 0.08944272 0.08944272 0.16431677 0.17029386
## [37] 0.16431677 0.27748874 0.43474130 0.03162278 0.16733201 0.05477226
## [43] 0.06324555
```

You need to use the $SE = \sqrt{sd/n}$

Rewriting the CI correctly!

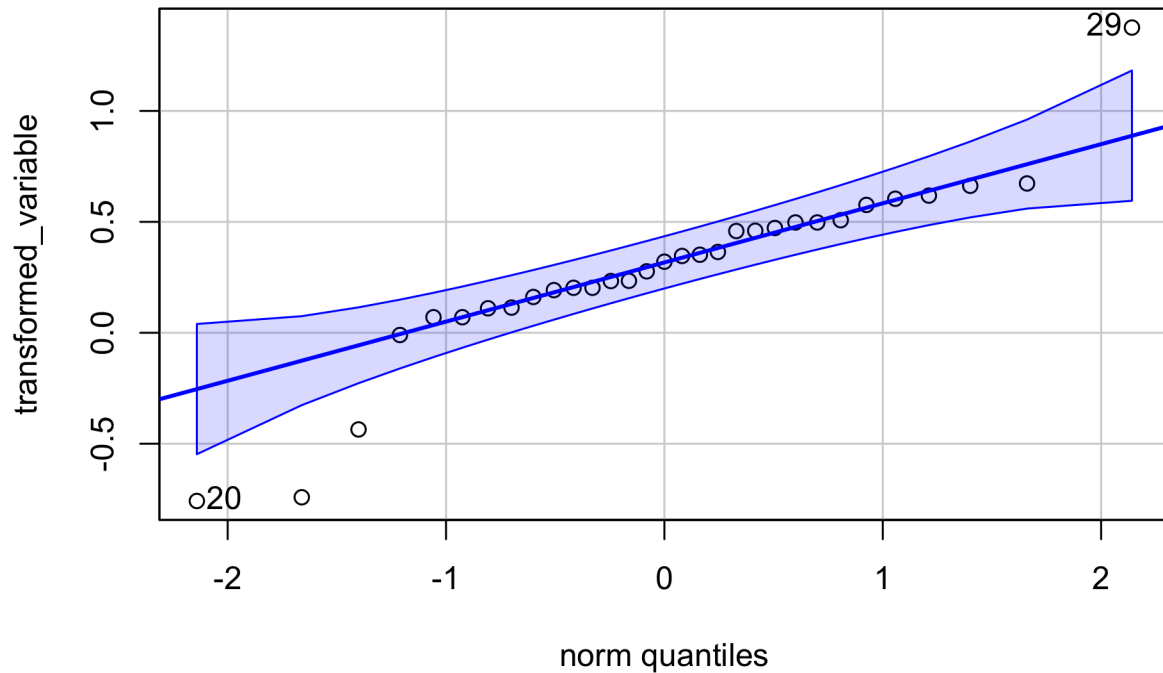
```
## [1] "Authors" "Sample_size_included"
## [3] "Sex_(%female)" "Ethnic_background_(%_white)"
## [5] "age_range" "d_or_R"
## [7] "effect_size" "V"
## [9] "95%_CI" "std"
## [11] "lower_CI" "upper_CI"
## [13] "SE"
```

Considering only the r

Selecting some columns and renaming some of them:

Meta-Analysis for the r-correlation

```
## [1] "Smith, et al." "Doe, et al." "Johnson, et al."
```



```
## [1] 29 20
```

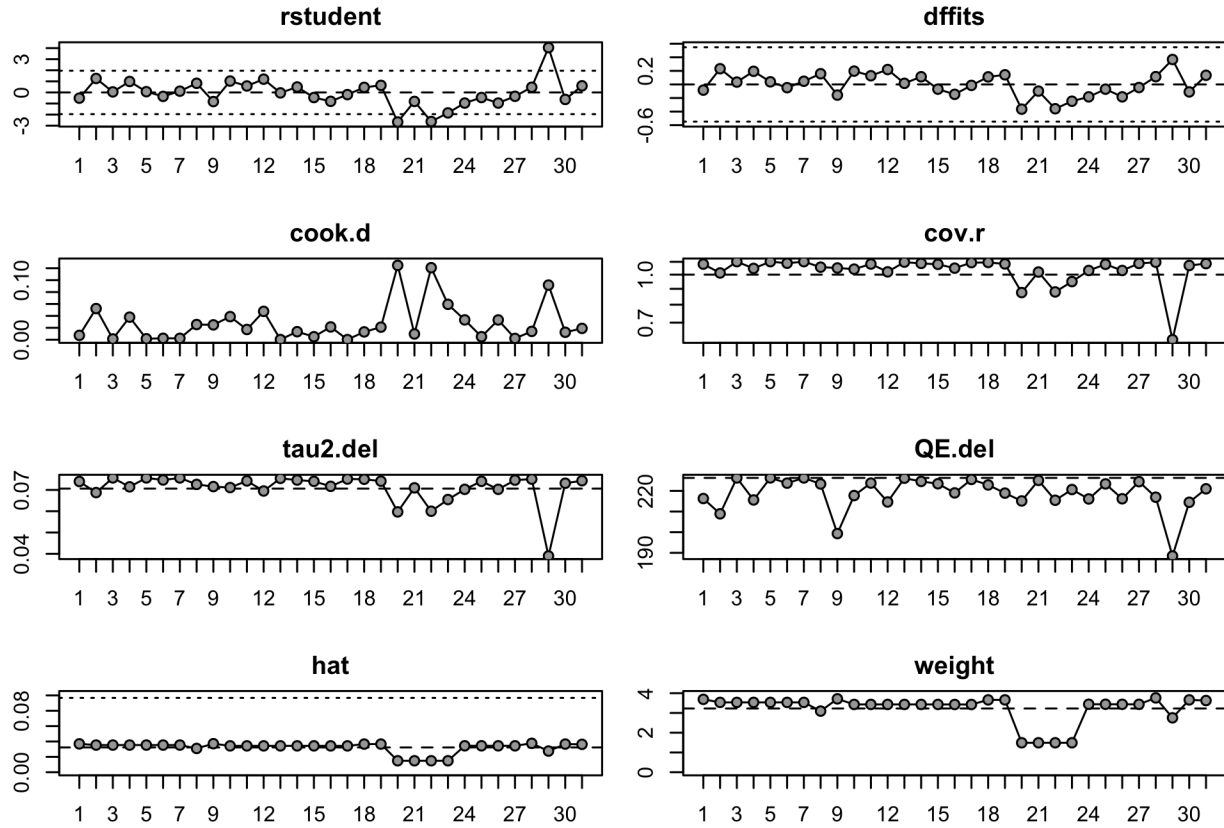
Notice that there are 4 outliers!

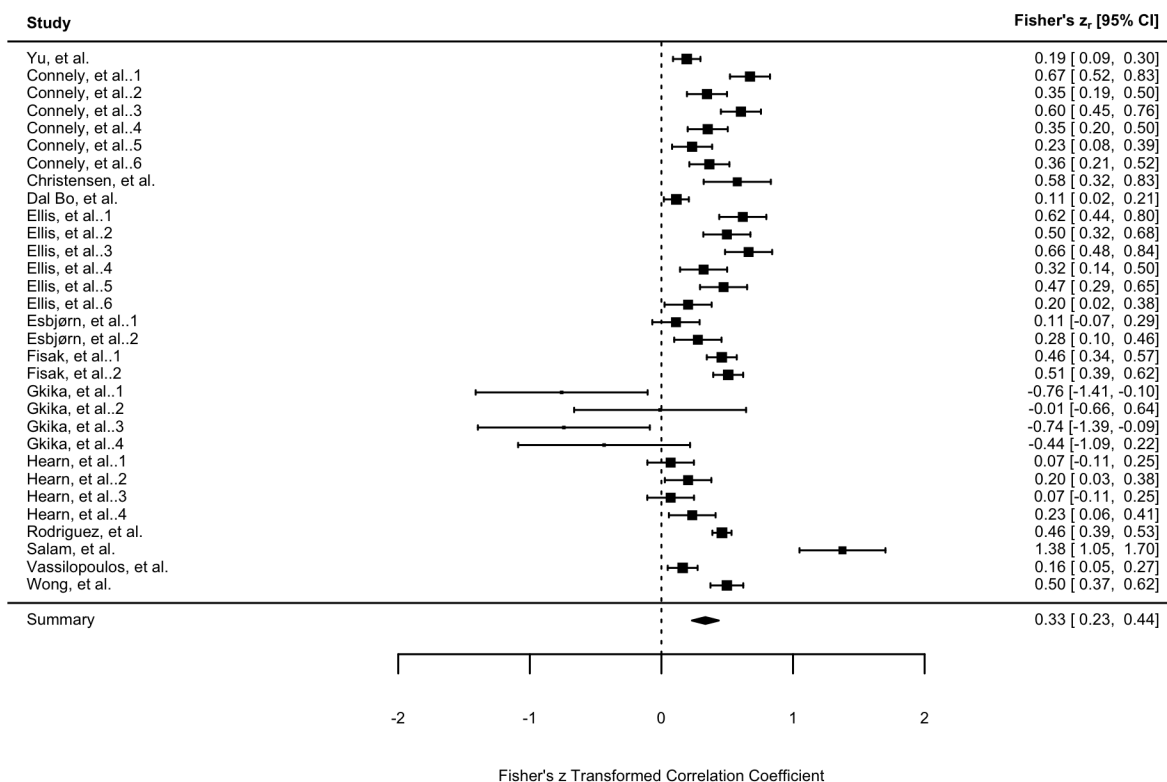
```
##
## Random-Effects Model (k = 31; tau^2 estimator: REML)
##
## tau^2 (estimated amount of total heterogeneity): 0.0706 (SE = 0.0212)
## tau (square root of estimated tau^2 value):      0.2657
## I^2 (total heterogeneity / total variability):    92.07%
## H^2 (total variability / sampling variability):    12.61
##
## Test for Heterogeneity:
## Q(df = 30) = 226.2882, p-val < .0001
##
## Model Results:
##
## estimate      se      zval      pval      ci.lb      ci.ub
## 0.3339 0.0520 6.4152 <.0001 0.2319 0.4359 ***
##
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Usually reported the Q statistics: binary test

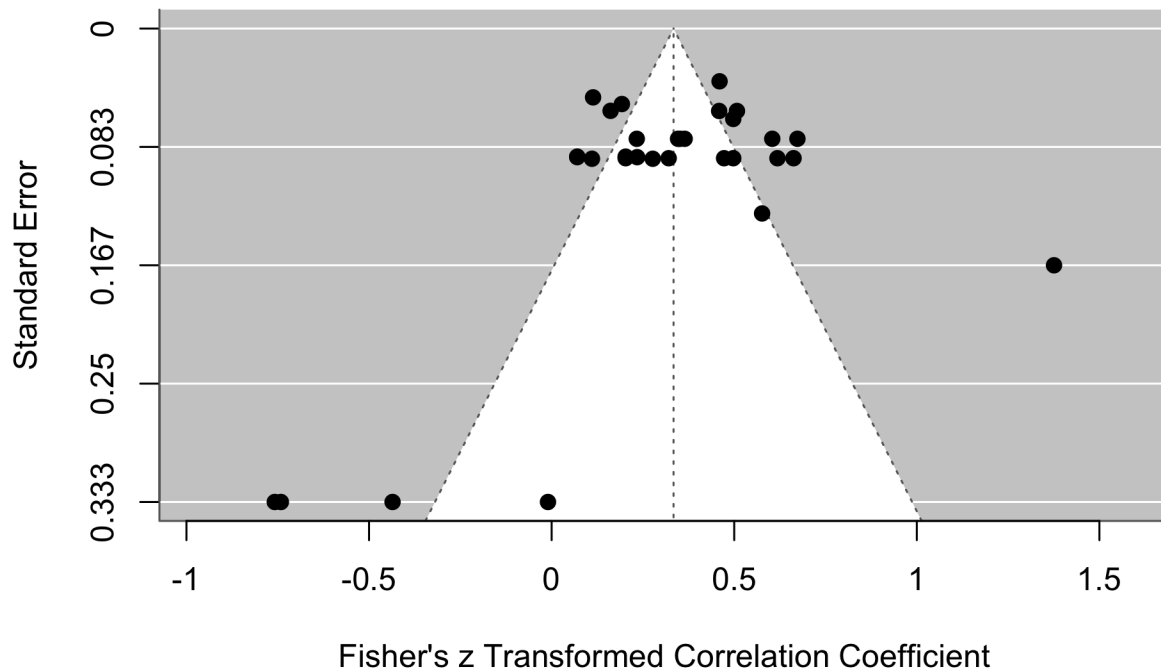
[illegible]

No influential studies, since no “*” in the inf column





Study Bias:



```
##
## Regression Test for Funnel Plot Asymmetry
##
## Model:      mixed-effects meta-regression model
## Predictor: standard error
##
## Test for Funnel Plot Asymmetry: z = -2.9154, p = 0.0036
## Limit Estimate (as sei -> 0):  b = 0.5591 (CI: 0.3786, 0.7396)
```

There is some Funnel Plot asymmetry, also proven by this formal test

Moderator analysis on r

```
## [1] "Authors"      "ni"            "type"          "effect_size"
## [5] "V"            "std"           "lower_CI"      "upper_CI"
## [9] "SE"           "female_perc"   "white_ethnicity" "yi"
## [13] "vi"

## Warning: 1 study with NAs omitted from model fitting.

##
## Mixed-Effects Model (k = 30; tau^2 estimator: REML)
##
## tau^2 (estimated amount of residual heterogeneity): 0.0435 (SE = 0.0142)
## tau (square root of estimated tau^2 value): 0.2086
## I^2 (residual heterogeneity / unaccounted variability): 88.01%
## H^2 (unaccounted variability / sampling variability): 8.34
## R^2 (amount of heterogeneity accounted for): 0.00%
```



```

##
## Test for Residual Heterogeneity:
## QE(df = 28) = 184.1213, p-val < .0001
##
## Test of Moderators (coefficient 2):
## QM(df = 1) = 4.3411, p-val = 0.0372
##
## Model Results:
##
##           estimate      se      zval      pval      ci.lb      ci.ub
## intrcpt          0.7830  0.2283   3.4300  0.0006   0.3356   1.2304 ***
## female_perc     -0.6836  0.3281  -2.0835  0.0372  -1.3267  -0.0405  *
##
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

## Warning: 13 studies with NAs omitted from model fitting.

##
## Mixed-Effects Model (k = 18; tau^2 estimator: REML)
##
## tau^2 (estimated amount of residual heterogeneity):      0.0527 (SE = 0.0212)
## tau (square root of estimated tau^2 value):              0.2296
## I^2 (residual heterogeneity / unaccounted variability): 90.08%
## H^2 (unaccounted variability / sampling variability):    10.08
## R^2 (amount of heterogeneity accounted for):              19.55%
##
## Test for Residual Heterogeneity:
## QE(df = 16) = 124.0092, p-val < .0001
##
## Test of Moderators (coefficient 2):
## QM(df = 1) = 5.7961, p-val = 0.0161
##
## Model Results:
##
##           estimate      se      zval      pval      ci.lb      ci.ub
## intrcpt          0.7000  0.1491   4.6949 <.0001   0.4078   0.9922 ***
## white_ethnicity  -0.4438  0.1843  -2.4075  0.0161  -0.8050  -0.0825  *
##
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

```

We get a significant result for white_ethnicity prevalence (maybe we need to transform this variable too!).

Removing Outliers only r

Removing the rows of the author Gkika and Salam

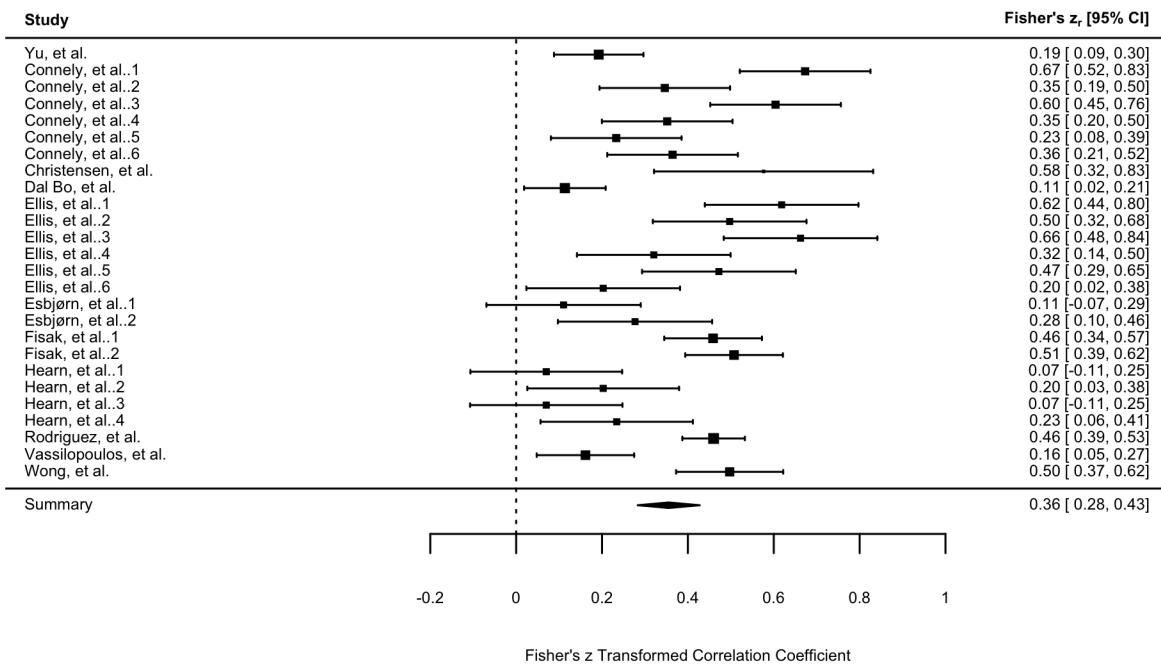
```

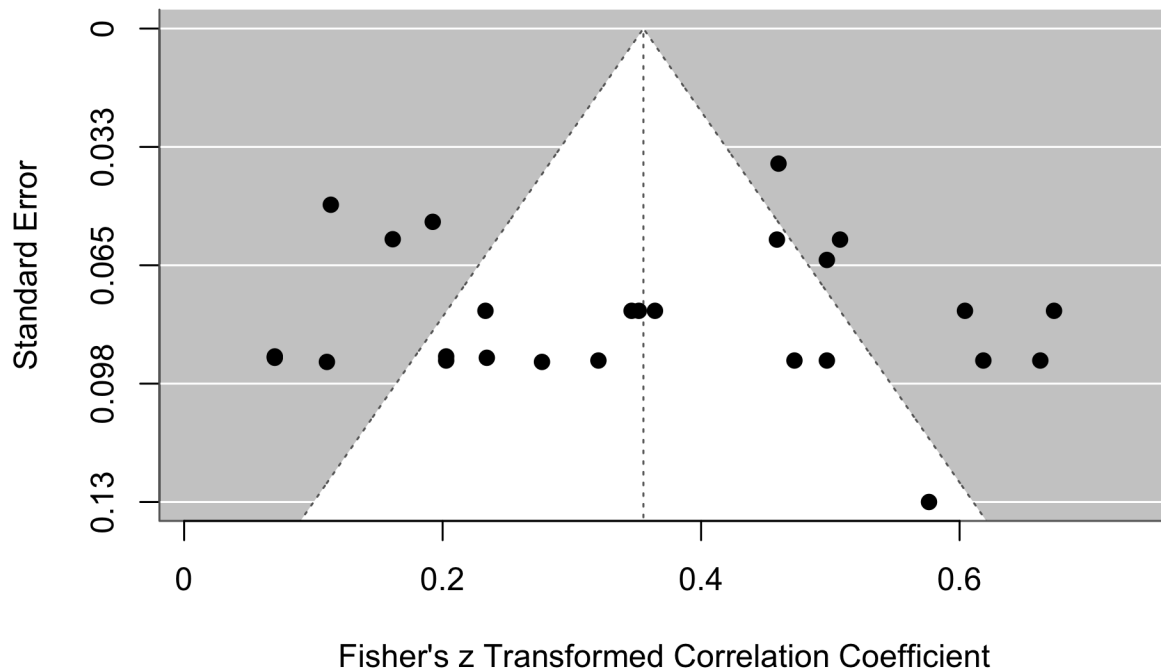
## [1] TRUE TRUE TRUE TRUE TRUE TRUE TRUE TRUE TRUE TRUE TRUE TRUE
## [13] TRUE TRUE TRUE TRUE TRUE TRUE TRUE TRUE FALSE FALSE FALSE FALSE TRUE
## [25] TRUE TRUE TRUE TRUE FALSE TRUE TRUE

##
## Random-Effects Model (k = 26; tau^2 estimator: REML)
##
## tau^2 (estimated amount of total heterogeneity): 0.0289 (SE = 0.0100)
## tau (square root of estimated tau^2 value):      0.1701

```

```
## I^2 (total heterogeneity / total variability): 84.89%
## H^2 (total variability / sampling variability): 6.62
##
## Test for Heterogeneity:
## Q(df = 25) = 159.9674, p-val < .0001
##
## Model Results:
##
## estimate      se      zval      pval      ci.lb      ci.ub
## 0.3554 0.0369 9.6379 <.0001 0.2831 0.4276 ***
##
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```





```
##
## Regression Test for Funnel Plot Asymmetry
##
## Model:      mixed-effects meta-regression model
## Predictor: standard error
##
## Test for Funnel Plot Asymmetry: z = 0.3717, p = 0.7101
## Limit Estimate (as sei -> 0):  b = 0.2975 (CI: -0.0165, 0.6115)
```

Accounting for non-independence of the effect sizes from same study only in r

Aooo

```
## # A tibble: 6 x 11
##   Authors          ni type effect_size      V      std lower_CI upper_CI      SE
##   <chr>          <dbl> <fct>      <dbl> <dbl> <dbl>      <dbl>      <dbl>      <dbl>
## 1 Yu, Meng and L~  357 r          0.19  0.004 0.0632    0.183    0.197 0.00335
## 2 Connely, K.     169 r          0.587 0.006 0.0775    0.575    0.599 0.00596
## 3 Connely, K.     169 r          0.333 0.006 0.0775    0.321    0.345 0.00596
## 4 Connely, K.     169 r          0.54   0.006 0.0775    0.528    0.552 0.00596
## 5 Connely, K.     169 r          0.338 0.006 0.0775    0.326    0.350 0.00596
## 6 Connely, K.     169 r          0.229 0.006 0.0775    0.217    0.241 0.00596
## # i 2 more variables: female_perc <dbl>, white_ethnicity <dbl>

## RVE: Hierarchical Effects Model with Small-Sample Corrections
##
## Model: yi ~ 1
```

```
##
## Number of clusters = 14
## Number of outcomes = 31 (min = 1 , mean = 2.21 , median = 1.5 , max = 6 )
## Omega.sq = 0
## Tau.sq = 0.04231477
##
##           Estimate StdErr t-value  dfs P(|t|>) 95% CI.L 95% CI.U Sig
## 1 X.Intercept.    0.343 0.0656    5.23 7.37 0.00103    0.189    0.496 ***
## ---
## Signif. codes: < .01 *** < .05 ** < .10 *
## ---
## Note: If df < 4, do not trust the results
```

Considering only d

```
## [1] "Authors"      "ni"            "type"          "effect_size"
## [5] "V"            "std"          "lower_CI"      "upper_CI"
## [9] "SE"           "female_perc"  "white_ethnicity"
```

Full meta-analysis

Converting r to d, standardize and perform the analysis

```
## # A tibble: 31 x 2
##   effect_size SE
##   <dbl> <dbl>
## 1    0.19 0.00335
## 2    0.587 0.00596
## 3    0.333 0.00596
## 4    0.54 0.00596
## 5    0.338 0.00596
## 6    0.229 0.00596
## 7    0.349 0.00596
## 8    0.52 0.0166
## 9    0.113 0.00216
## 10    0.55 0.00806
## # i 21 more rows

## # A tibble: 31 x 2
##   effect_size_d SE
##   <dbl> <dbl>
## 1    0.387 0.508
## 2    1.45 1.05
## 3    0.706 0.663
## 4    1.28 0.932
## 5    0.718 0.667
## 6    0.471 0.602
## 7    0.745 0.675
## 8    1.22 1.15
## 9    0.227 0.440
## 10    1.32 1.03
## # i 21 more rows
```

Now all our effect sizes are d!

metafor analysis

```
## # A tibble: 43 x 2
##   effect_size SE
##   <dbl> <dbl>
## 1      0.387 0.508
## 2      5.22 0.127
## 3     -0.119 0.05
## 4      1.45 1.05
## 5      0.706 0.663
## 6      1.28 0.932
## 7      0.718 0.667
## 8      0.471 0.602
## 9      0.745 0.675
## 10     1.22 1.15
## # i 33 more rows
```

Converting our effect sizes into the SMD, need to check that this is indeed true

```
##
## Random-Effects Model (k = 43; tau^2 estimator: DL)
##
## tau^2 (estimated amount of total heterogeneity): 0.6593 (SE = 0.4414)
## tau (square root of estimated tau^2 value):      0.8120
## I^2 (total heterogeneity / total variability):   99.62%
## H^2 (total variability / sampling variability):   262.98
##
## Test for Heterogeneity:
## Q(df = 42) = 11045.2953, p-val < .0001
##
## Model Results:
##
## estimate      se      zval      pval      ci.lb      ci.ub
##    0.7912    0.1544    5.1254    <.0001    0.4886    1.0937    ***
##
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

