

Meta-Analysis_Kinesiophobia_Physical_Activity

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R Markdown file set-up and packages

Packages required: 1) dmetar 2) tidyverse 3) meta 4) metafor 5) metasens 6) esc

Meta-analysis method

We pooled Pearson product-moment correlations from eligible studies to examine the relationship between kinesophobia and physical activity. Correlations were pooled using the generic inverse pooling method via the ‘metacor’ function in the {meta} R package. This function automatically performs a necessary Fisher’s z-transformation on the original, untransformed correlations prior to pooling. The ‘metacor’ function also reconverts the pooled association back to its original form for ease of interpretation. Correlation estimates were nested within studies using the ‘cluster’ argument to account for the dependencies between these estimates, resulting in a three-level meta-analysis (level 1: participants, level 2: correlation estimates, level 3: studies). The distribution of variance across levels was assessed using the multilevel version of I2. The performance of the 2-level and 3-level meta-analyses was assessed and compared using the {metafor} R package.

We anticipated considerable between-study heterogeneity, and therefore used a random-effects model to pool correlations. The restricted maximum likelihood (RML) estimator (Viechtbauer, 2005) was used to calculate the heterogeneity variance Tau2. In addition to Tau2, to quantify between-study heterogeneity, we report the I2 statistic, which provides the percentage of variability in the correlations that is not caused by sampling error⁵⁶. The I2 statistic was interpreted as follows: 0-40%, may not be important; 30-60%, may represent moderate heterogeneity; 50-90%, may represent substantial heterogeneity; and 75-100%, may represent considerable heterogeneity. To reduce the risk of false positives, we used a Knapp-Hartung adjustment (Knapp and Hartung, 2003) to calculate the confidence interval around the pooled association. We also report the prediction interval, which provides a range within which we can expect the associations of future studies to fall based on the current evidence. The pooled correlation was interpreted using Cohen’s conventions (Cohen, 1988): $r = -0.10$, small negative correlation; $r = -0.30$, moderate negative correlation; $r = -0.50$, large negative correlation.

Publication bias was assessed using a funnel plot, which is a scatter plot of the studies’ effect size expressed as the Fisher’s z transformed correlation on the x-axis against a measure of their standard error (which is indicative of precision of the study’s effect size) on the y-axis. When there is no publication bias, the data points in a funnel plot should form a roughly symmetrical, upside-down funnel. Studies in the top part of the plot, which have lower standard errors, are expected to lie closely together, and not far away from the pooled effect size. In the lower part of the plot, studies have higher standard errors, the funnel “opens up”, and effect sizes are expected to scatter more heavily to the left and right of the pooled effect. Egger’s regression (Egger et al., 1997) can be used to formally test funnel plot’s asymmetry. However, since there is no direct function to conduct Egger’s test for 3-level models, we calculated it by using the standard errors of the effect size estimates as a predictor in the meta-regression⁶⁰.

P-curve analysis (Simonsohn et al., 2014) was conducted to assess whether the distribution of the statistically significant results was consistent with what would be expected if only true effects were present. When the

null hypothesis is true (i.e., there is no true effect), p-values are assumed to follow a uniform distribution: highly significant effects (e.g., $p = 0.01$) are as likely as barely significant effects (e.g., $p = 0.049$). However, when the null hypothesis is false (i.e., there is a true effect in our data), p-values are assumed to follow a right-skewed distribution: highly significant effects are more likely than barely significant effects. A left-skewed distribution would suggest that some studies used statistical tests to find significant results in ways that may not be reproducible or generalizable (i.e., p-hacking).

A secondary meta-analysis was conducted using the same approach, but based on Spearman's rho values, to further test the relationship between kinesiophobia and device-based physical activity.

Subgroup analyses were conducted to examine the differences in correlations between studies including participants with different health conditions and using different types of physical activity measures (i.e., device-based versus self-reported), physical activity measurement instruments (i.e., type of questionnaires, type of devices), physical activity outcomes, and kinesiophobia measures.

Exploratory meta-regressions were conducted to examine if the average age of participants, the proportion of women, and pain in a study predicted the reported correlation between kinesiophobia and physical activity. Pain was normalized to a 0-100 scale to make the data comparable across pain scales. A sensitivity analysis was conducted to examine whether the quality of the studies affected the results.

All analyses were performed in RStudio integrated development environment (IDE) (2023.06.1+524, "Mountain Hydrangea" release) for R software environment (R Core Team, 2023) using the 'meta' (Schwarzer, 2023), 'metasens' (Schwarzer et al., 2023), and 'metafor' (Vietchbauer, 2010, 2023) R packages.

Meta-analysis: primary analysis

Import Pearson r data

Data file glimpse (not included in PDF)

Primary analysis using metafor to compare models

```
m.cor <- rma.mv(yi = z, # pool correlation coefficients Fishers Z transformation https://bookdown.org/m
               V = var.z, # sampling variance of Fisher's Z values: 1/(N-3), where N = the sample size
               slab = author,
               data = Kinphob_r,
               random = ~ 1 | author/cor_id,
               test = "t",
               method = "REML")
summary(m.cor)
```

```
##
## Multivariate Meta-Analysis Model (k = 83; method: REML)
##
##   logLik  Deviance      AIC      BIC      AICc
## -11.1832  22.3664   28.3664   35.5865   28.6741
##
## Variance Components:
##
##           estim      sqrt  nlvls  fixed      factor
## sigma^2.1  0.0576  0.2399     63     no      author
## sigma^2.2  0.0058  0.0762     83     no  author/cor_id
##
## Test for Heterogeneity:
## Q(df = 82) = 564.7688, p-val < .0001
```

```
##
## Model Results:
##
## estimate      se      tval  df    pval    ci.lb    ci.ub
## -0.1956  0.0343  -5.6965  82  <.0001  -0.2639  -0.1273  ***
##
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Converting Fisher's z back to Pearson's r

```
round(convert_z2r(-0.1956), 2) # point estimate
```

```
## [1] -0.19
```

```
round(convert_z2r(-0.2639), 2) # lower CI
```

```
## [1] -0.26
```

```
round(convert_z2r(-0.1273), 2) # Upper CI
```

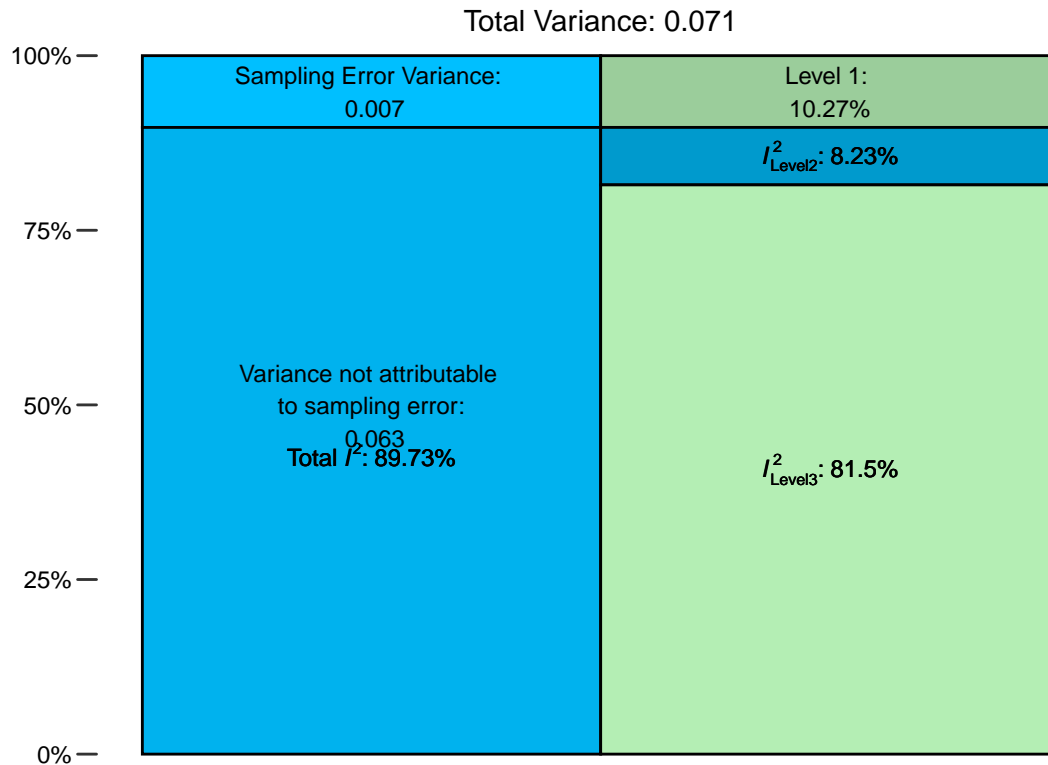
```
## [1] -0.13
```

Explore heterogeneity

```
i2 <- var.comp(m.cor)
summary(i2)
```

```
##           % of total variance    I2
## Level 1           10.274387  ---
## Level 2           8.226472  8.23
## Level 3           81.499141  81.5
## Total I2: 89.73%
```

```
plot(i2)
```



The sampling error variance on level 1 and the value of I^2 on level 2, i.e., the amount of heterogeneity variance within studies, were small (10.3% and 8.2%, respectively). The largest share of heterogeneity variance was from level 3, with between-study heterogeneity making up 81.5% of the total variation in our data. Overall, this indicates that there is considerable between-study heterogeneity, and less than one tenth of the variance can be explained by differences within studies.

Comparing models

Reduced model in which the level 3 variance (between-study heterogeneity) is set to 0, which assumes all effect sizes are independent.

```
l3.removed <- rma.mv(yi = z,
  V = var.z,
  slab = author,
  data = Kiphob_r,
  random = ~ 1 | author/cor_id,
  test = "t",
  method = "REML",
  sigma2 = c(0, NA))

summary(l3.removed)

##
## Multivariate Meta-Analysis Model (k = 83; method: REML)
##
##   logLik  Deviance      AIC      BIC     AICc
## -17.5166  35.0333  39.0333  43.8467  39.1852
##
## Variance Components:
##
##           estim      sqrt  nlvls  fixed      factor
```

```
## sigma^2.1  0.0000  0.0000    63   yes      author
## sigma^2.2  0.0661  0.2571    83   no   author/cor_id
##
## Test for Heterogeneity:
## Q(df = 82) = 564.7688, p-val < .0001
##
## Model Results:
##
## estimate      se      tval  df      pval      ci.lb      ci.ub
## -0.2159  0.0316  -6.8373  82  <.0001  -0.2787  -0.1531  ***
##
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Comparing full and reduced models.

```
anova(m.cor, 13.removed)
```

```
##
##          df      AIC      BIC      AICc  logLik      LRT  pval      QE
## Full      3 28.3664 35.5865 28.6741 -11.1832                564.7688
## Reduced   2 39.0333 43.8467 39.1852 -17.5166 12.6669 0.0004 564.7688
```

The 3-level model showed a better fit than the 2-level model with lower Akaike's information criterion (AIC) (28.4 vs. 39.0) and Bayesian information criterion (BIC) (35.6 vs. 43.8), indicating better performance. These lower AIC and BIC are consistent with the significant likelihood ratio test (LRT) comparing the two models ($2 = 12.67$, $p = 0.0004$). Therefore, although the 3-level model introduces an additional parameter, this added complexity has improved our estimate of the pooled effect.

Primary analysis using metacor

```
m <- metacor(cor = cor,
             n = n,
             studlab = author,
             data = Kiphob_r,
             cluster = cluster,
             fixed = FALSE,
             random = TRUE,
             method.tau = "REML",
             method.random.ci = "HK",
             prediction = TRUE,
             title = "Primary meta-analysis using metacor")
m

## Review:      Primary meta-analysis using metacor
##
## Number of studies: n = 63
## Number of estimates: k = 83
## Number of observations: o = 12278
##
##              COR              95%-CI      t  p-value
## Random effects model -0.1931 [-0.2579; -0.1266] -5.70 < 0.0001
## Prediction interval          [-0.6051;  0.3004]
##
## Quantifying heterogeneity:
```

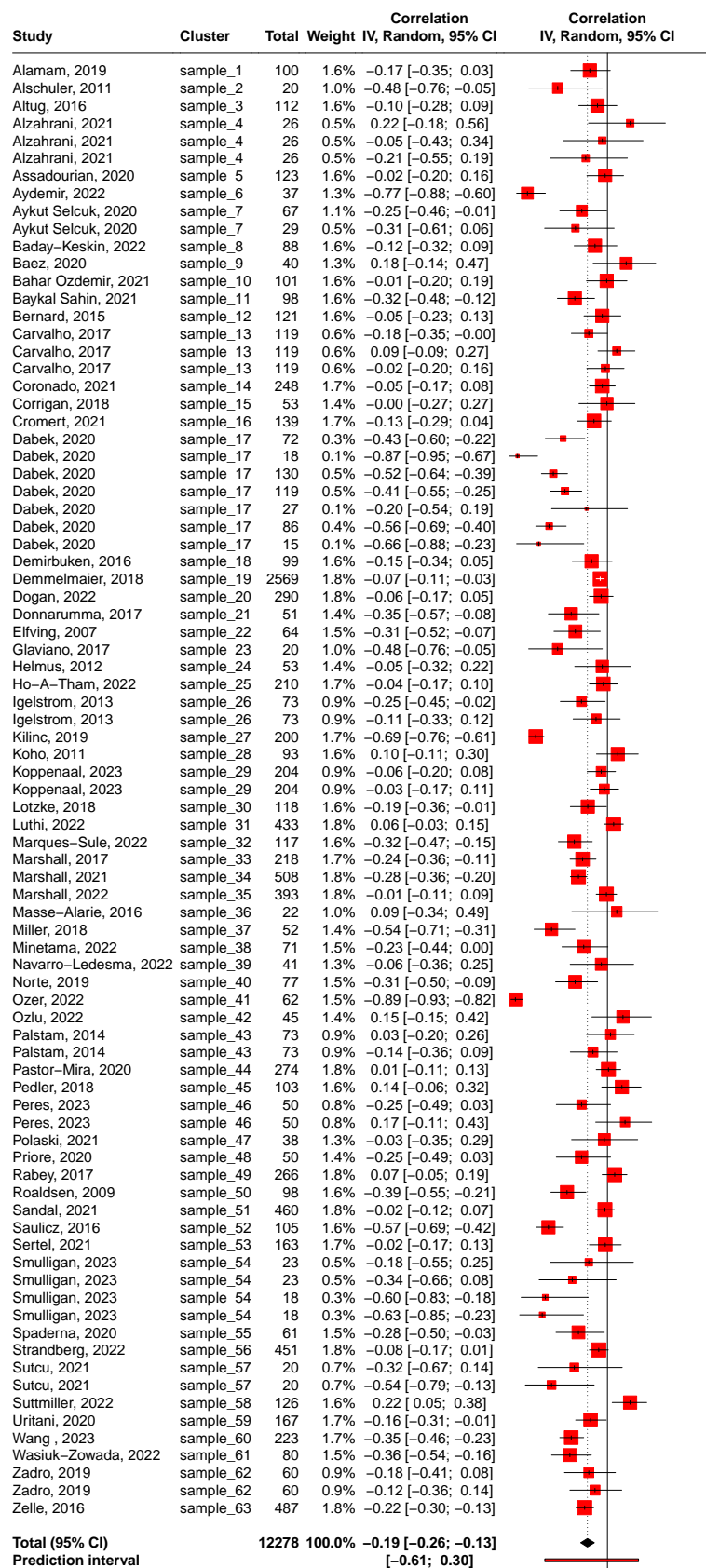
```
## tau^2.1 = 0.0576 [0.0227; 0.0950]; tau.1 = 0.2399 [0.1508; 0.3082] (between cluster)
## tau^2.2 = 0.0058 [0.0000; 0.0398]; tau.2 = 0.0762 [0.0000; 0.1995] (within cluster)
## I^2 = 85.5% [82.6%; 87.9%]; H = 2.62 [2.40; 2.87]
##
## Test of heterogeneity:
##      Q d.f.  p-value
## 564.77   82 < 0.0001
##
## Details on meta-analytical method:
## - Inverse variance method (three-level model)
## - Restricted maximum-likelihood estimator for tau^2
## - Profile-Likelihood method for confidence interval of tau^2 and tau
## - Random effects confidence interval based on t-distribution (df = 82)
## - Prediction interval based on t-distribution (df = 81)
## - Fisher's z transformation of correlations
```

Our main meta-analysis of 63 studies, 83 Pearson's r correlation estimates, and 12278 participants revealed a statistically significant small-to-moderate negative correlation between kinesiphobia and physical activity ($r = -0.19$; 95% confidence interval [95CI]: -0.26 to -0.13; $p < 0.0001$). However, we observed substantial-to-considerable between-study statistical heterogeneity ($\text{Tau}^2 = 0.06$, 95CI: 0.02 to 0.09; $I^2 = 85.5\%$, 95CI: 82.6 to 87.9%), and the prediction interval ranged from $r = -0.61$ to 0.30, indicating that a moderate positive correlation cannot be ruled out for future studies.

Primary meta-analysis forest plots

Forest plot with Pearson r correlation coefficients

```
# Generate the forest plot without shading
forest(m,
  layout = "Revman5",
  prediction = TRUE,
  print.tau2 = FALSE,
  leftlabs = c("Author", "n"),
  xlim=c(-1.0,1.0),
  fs.hetstat = 10,
  addrows.below.overall = 2)
```



Heterogeneity: $\tau^2 = 0.0634$; $\chi^2 = 564.77$, $df = 82$ ($P < 0.01$); $I^2 = 85\%$

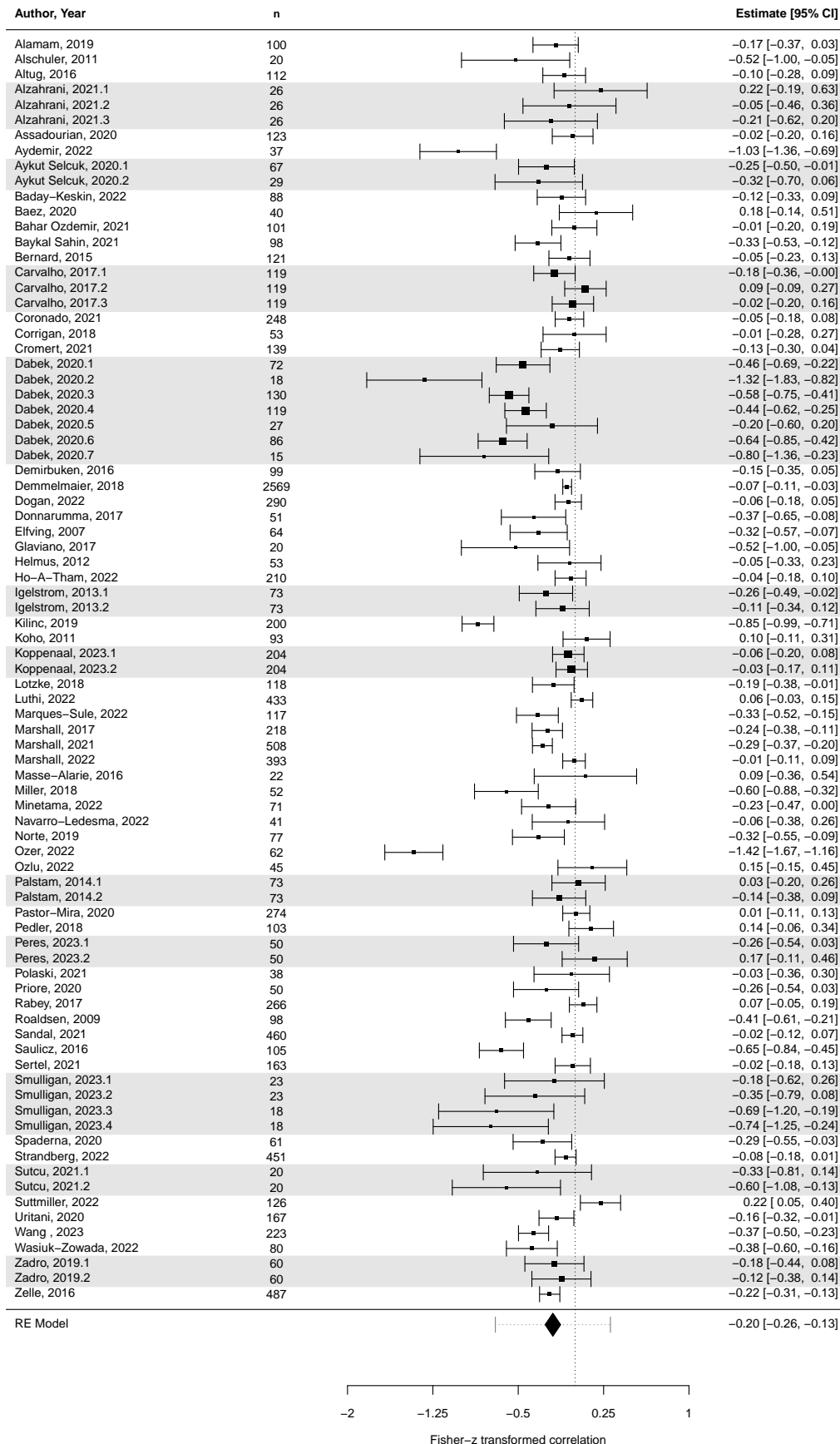
Save Forest plot:

```
png(file = "Main analysis forestplot.png",  
    width = 2800, height = 4000, res = 300)  
  
forest(m,  
       xlim = c(-1.0, 0.6),  
       layout = "Revman")  
  
dev.off()
```

```
## pdf  
## 2
```

Forest plot with Fisher z transformed correlation coefficients

```
forest(m.cor,  
       cex=0.8,  
       addpred=TRUE,  
       alim=c(-2,1),  
       xlab = "Fisher-z transformed correlation",  
       ilab = n,  
       header="Author, Year",  
       shade=c(2,3,8,9,12,13,14,15,23,24,27,28,42,43,46,47,56,57,58,59,60,61,62,66,67,68,74,75,78,79,80),  
       text(c(-2.62), m.cor$k+2, c("n"), cex=0.75, font=2)
```



Save plot:

```
png(file = "Main analysis Fisher z forest plot.png",
     width = 2800, height = 4000, res = 300)

forest(m.cor,
       cex=0.8,
       addpred=TRUE,
       alim=c(-2,2),
       xlab = "Fisher-z transformed correlation",
       ilab = n,
       header="Author(s), Year, estimate n",
       shade=c(1,2,4,5,12,13,14,15,23,24,34,35,36,40,41,44,45,50,51,52,53,54,55,56,65,66,69,70,71)) # f
text(c(-2.62), m.cor$k+2, c("n"), cex=0.75, font=2)

dev.off()

## pdf
## 2
```

Forest plot with aggregated Fisher z values

```
dat_r <- escalc(measure="ZCOR",
               yi=z,
               vi=var.z,
               slab = author,
               data=Kinpob_r)

agg <- aggregate(dat_r,
                 cluster=author,
                 V = vcov(m.cor,
                         type="obs"),
                 struct="ID",
                 addk=TRUE)

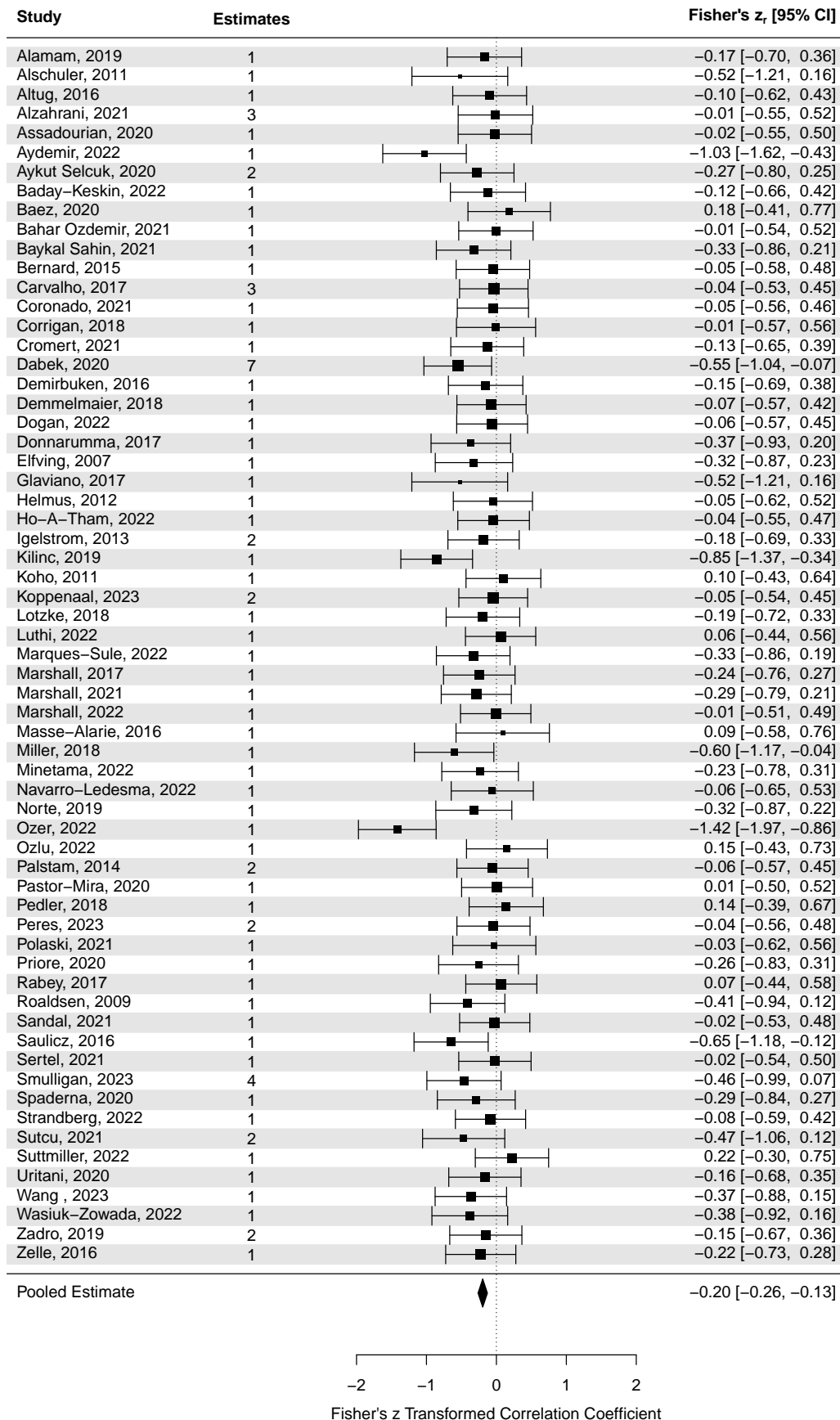
res <- rma(yi,
           vi,
           method="EE",
           data=agg,
           digits=2)

res

##
## Equal-Effects Model (k = 63)
##
## I2 (total heterogeneity / total variability): 4.39%
## H2 (total variability / sampling variability): 1.05
##
## Test for Heterogeneity:
## Q(df = 62) = 64.85, p-val = 0.38
##
## Model Results:
##
## estimate    se    zval  pval  ci.lb  ci.ub
##    -0.20  0.03  -5.70  <.01  -0.26  -0.13  ***
```

```
##
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

forest(res,
      xlim=c(-7,5),
      mlab="Pooled Estimate",
      header=TRUE,
      alim=c(-2,2),
      slab = author,
      ilab=ki,
      ilab.xpos=-3.5,
      shade = "zebra")
text(-3.5, res$k+2, "Estimates", font=2)
```



Save plot:

```
png(file = "Main analysis aggregated forestplot.png",
     width = 2800, height = 4000, res = 300)

forest(res,
       xlim=c(-7,5),
       mlab="Pooled Estimate",
       header=TRUE,
       alim=c(-2.5,2.5),
       slab = author,
       ilab=ki,
       ilab.xpos=-3.5,
       shade = "zebra")
text(-3.5, res$k+2, "Estimates", font=2)

dev.off()
```

```
## pdf
## 2
```

Secondary analysis based on Spearman's rho values

Read in rho data

```
Kinhob_rho <- read.csv("2024-05-02_meta_analysis_kinesiophobia_rho.csv",
                      header = TRUE, sep = ",")
```

Secondary analysis using metafor to compare models

```
m.cor.rho <- rma.mv(yi = z,
                   V = var.z,
                   slab = author,
                   data = Kinhob_rho,
                   random = ~ 1 | author/cor_id,
                   test = "t",
                   method = "REML")

summary(m.cor.rho)

##
## Multivariate Meta-Analysis Model (k = 21; method: REML)
##
##   logLik  Deviance      AIC      BIC      AICc
##   0.6454   -1.2908    4.7092    7.6964    6.2092
##
## Variance Components:
##
##           estim    sqrt  nlvls  fixed      factor
## sigma^2.1  0.0982  0.3134    12    no      author
## sigma^2.2  0.0000  0.0000    21    no  author/cor_id
##
## Test for Heterogeneity:
## Q(df = 20) = 146.0167, p-val < .0001
##
```

```
## Model Results:
##
## estimate      se      tval  df    pval    ci.lb    ci.ub
## -0.2018  0.0961  -2.1003  20  0.0486  -0.4023  -0.0014  *
##
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

round(convert_z2r(-0.0959), 3) # point estimate

## [1] -0.096

round(convert_z2r(-0.4017), 3) # lower CI

## [1] -0.381

round(convert_z2r(-0.0017), 3) # Upper CI

## [1] -0.002
```

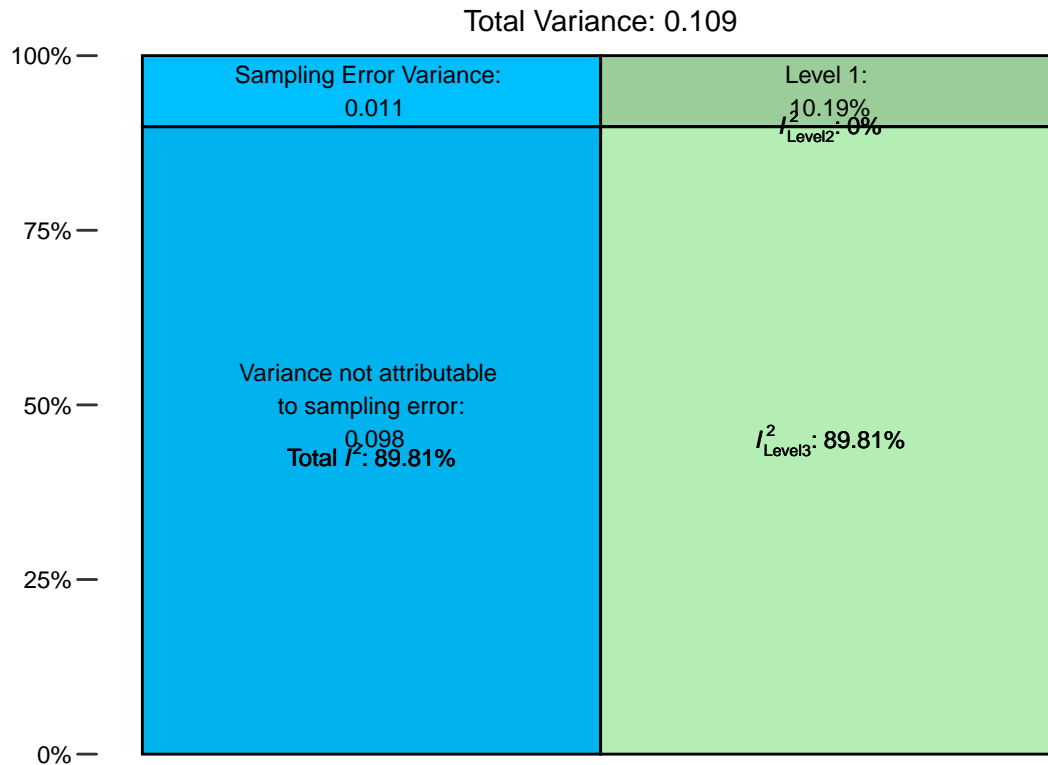
Results of the secondary meta-analysis of 12 studies, 21 Spearman's rho correlation estimates, and 2084 participants was consistent with the main meta-analysis as it showed a statistically significant small-to-moderate negative correlation between kinesiphobia and physical activity ($r = -0.20$; 95CI: -0.38 to -0.01; $p = 0.049$). However, we observed substantial-to-considerable between-study statistical heterogeneity ($\text{Tau}^2 = 0.10$, 95CI: 0.04 to 0.28; $I^2 = 86.3\%$) and the prediction interval ranged from $r = -0.71$ to 0.45, indicating that a moderate positive correlation cannot be ruled out for future studies.

Explore heterogeneity, rho

```
i2_rho <- var.comp(m.cor.rho)
summary(i2_rho)

##           % of total variance      I2
## Level 1           1.018861e+01    ---
## Level 2           2.089253e-10      0
## Level 3           8.981139e+01  89.81
## Total I2: 89.81%

plot(i2_rho)
```



The value of I^2 Level 2, the amount of heterogeneity variance within clusters (i.e. within studies), is very low, totaling roughly 0%. The largest share, however, falls to level 3. Between-cluster (here: between-study) heterogeneity makes up I^2 Level 3 = 96.8% of the total variation in our data. Overall, this indicates that there is considerable between-study heterogeneity on the third level, and very little of the variance can be explained by differences within studies.

Compare models, rho

```
l3.removed_rho <- rma.mv(yi = z,
  V = var.z,
  slab = author,
  data = Kiphob_rho,
  random = ~ 1 | author/cor_id,
  test = "t",
  method = "REML",
  sigma2 = c(0, NA))

summary(l3.removed_rho)

##
## Multivariate Meta-Analysis Model (k = 21; method: REML)
##
##   logLik  Deviance      AIC      BIC     AICc
##   -3.1233   6.2465   10.2465   12.2380   10.9524
##
## Variance Components:
##
##      estim    sqrt  nlvls  fixed      factor
## sigma^2.1  0.0000  0.0000   12    yes      author
## sigma^2.2  0.0649  0.2548   21    no   author/cor_id
```



```
##
## Test for Heterogeneity:
## Q(df = 20) = 146.0167, p-val < .0001
##
## Model Results:
##
## estimate      se      tval  df      pval      ci.lb      ci.ub
## -0.1782  0.0644  -2.7655  20   0.0119  -0.3126  -0.0438  *
##
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
anova(m.cor.rho, 13.removed_rho)
```

```
##
##          df      AIC      BIC      AICc  logLik      LRT      pval      QE
## Full      3  4.7092  7.6964  6.2092  0.6454
## Reduced   2 10.2465 12.2380 10.9524 -3.1233 7.5374 0.0060 146.0167
```

Modeling of the nested data structure was probably a good idea, and has improved our estimate of the pooled effect.

Secondary meta-analysis using metacor

```
m.rho <- metacor(cor = rho,
  n = n,
  studlab = author,
  data = Kinphob_rho,
  cluster = cluster,
  fixed = FALSE,
  random = TRUE,
  method.tau = "REML",
  method.random.ci = "HK",
  prediction = TRUE,
  title = "Secondary meta-analysis using metacor")
```

```
m.rho
```

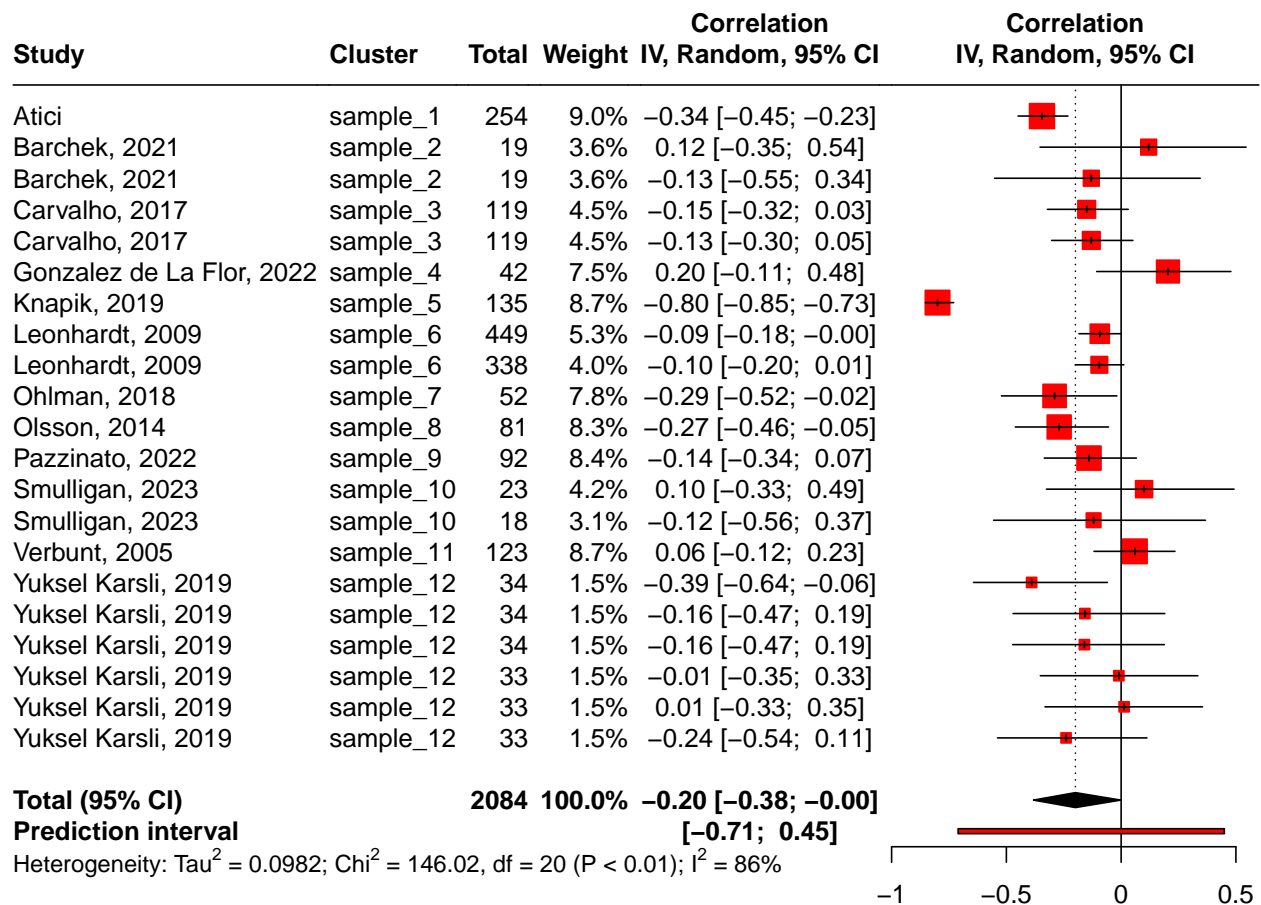
```
## Review:      Secondary meta-analysis using metacor
##
## Number of studies: n = 12
## Number of estimates: k = 21
## Number of observations: o = 2084
##
##              COR              95%-CI      t p-value
## Random effects model -0.1991 [-0.3819; -0.0014] -2.10  0.0486
## Prediction interval      [-0.7104;  0.4497]
##
## Quantifying heterogeneity:
## tau^2.1 = 0.0982 [0.0418; 0.2771]; tau.1 = 0.3134 [0.2044; 0.5264] (between cluster)
## tau^2.2 < 0.0001 [0.0000; 0.0172]; tau.2 < 0.0001 [0.0000; 0.1312] (within cluster)
## I^2 = 86.3% [80.4%; 90.4%]; H = 2.70 [2.26; 3.23]
##
## Test of heterogeneity:
##      Q d.f.  p-value
```

```
## 146.02 20 < 0.0001
##
## Details on meta-analytical method:
## - Inverse variance method (three-level model)
## - Restricted maximum-likelihood estimator for tau^2
## - Profile-Likelihood method for confidence interval of tau^2 and tau
## - Random effects confidence interval based on t-distribution (df = 20)
## - Prediction interval based on t-distribution (df = 19)
## - Fisher's z transformation of correlations
```

Secondary meta-analysis forest plots

Forest plot for secondary analysis with Pearson r correlation coefficients

```
forest(m.rho,
       xlim = c(-1.0, 0.6),
       prediction = TRUE,
       layout = "Revman")
```



Save plot:

```
png(file = "Secondary analysis forest plot with meta.png",
    width = 2800, height = 2500, res = 300)

forest(m.rho,
       xlim = c(-1.0, 0.6),
```

```
layout = "Revman")
```

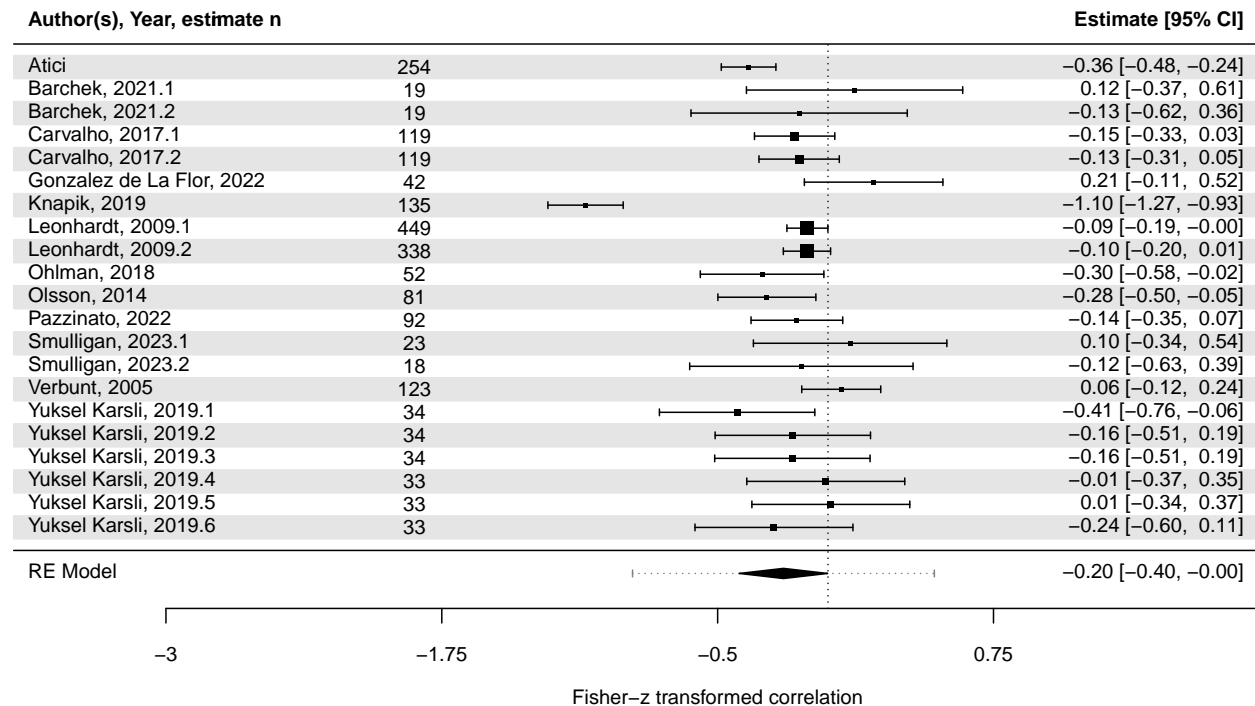
```
dev.off()
```

```
## pdf
```

```
## 2
```

Forest plot for secondary analysis with Fisher z using metafor

```
forest(m.cor.rho,
       cex=0.8,
       addpred=TRUE,
       alim=c(-3,2),
       xlab = "Fisher-z transformed correlation",
       ilab = n,
       header="Author(s), Year, estimate n",
       shade= "zebra")
text(c(-2.77), m.cor.rho$k+2, c("n"), cex=0.75, font=2)
```



Save plot:

```
png(file = "Rho analysis forestplot.png",
    width = 3000, height = 2600, res = 300)
```

```
forest(m.cor.rho,
       cex=0.8,
       addpred=TRUE,
       alim=c(-3,2),
       xlab = "Fisher-z transformed correlation",
       ilab = n,
       header="Author(s), Year, estimate n",
       shade= "zebra")
```

```
text(c(-2.77), m.cor.rho$k+2, c("n"), cex=0.75, font=2)

dev.off()
```

```
## pdf
## 2
```

Meta-analysis: subgroup analyses

Subgroup analysis by health status

```
Kinhob_r$Health_status <- as.factor(Kinhob_r$Health_status)
```

```
Health_subg <- rma.mv(yi = z,
                      V = var.z,
                      slab = author,
                      data = Kinhob_r,
                      random = ~ 1 | author/cor_id,
                      test = "t",
                      method = "REML",
                      mods = ~ Health_status)
```

```
summary(Health_subg)
```

```
##
## Multivariate Meta-Analysis Model (k = 83; method: REML)
##
##   logLik  Deviance      AIC      BIC      AICc
##   1.1209   -2.2418   27.7582   61.4857   36.6471
##
## Variance Components:
##
##              estim      sqrt  nlvls  fixed      factor
## sigma^2.1  0.0403  0.2008     63     no      author
## sigma^2.2  0.0044  0.0662     83     no  author/cor_id
##
## Test for Residual Heterogeneity:
## QE(df = 70) = 360.5491, p-val < .0001
##
## Test of Moderators (coefficients 2:13):
## F(df1 = 12, df2 = 70) = 3.1070, p-val = 0.0014
##
## Model Results:
##
##              estimate      se      tval  df      pval
## intrcpt          -0.1290  0.1807  -0.7137  70  0.4778
## Health_statusArthritis  -0.1295  0.1967  -0.6582  70  0.5126
## Health_statusCancer      0.0448  0.2821   0.1587  70  0.8743
## Health_statusCardiovascular condition -0.1813  0.2058  -0.8810  70  0.3813
## Health_statusChronic pain    0.0549  0.1860   0.2953  70  0.7687
## Health_statusFibromyalgia    0.0735  0.2869   0.2563  70  0.7985
## Health_statusNeurological condition -0.4570  0.2214  -2.0643  70  0.0427
```

```
## Health_statusObstructive Sleep Apnea      -0.0545  0.2869  -0.1899  70  0.8499
## Health_statusOlder adults                  -0.2919  0.2274  -1.2836  70  0.2035
## Health_statusPost-Partum Women             -0.0018  0.2911  -0.0060  70  0.9952
## Health_statusPulmonary condition           -0.7035  0.2450  -2.8710  70  0.0054
## Health_statusSurgery                       -0.0339  0.2097  -0.1617  70  0.8720
## Health_statusYoung adults                  0.1220  0.2959  0.4122  70  0.6814
##                                           ci.lb  ci.ub
## intrcpt                                  -0.4894  0.2315
## Health_statusArthritis                   -0.5218  0.2628
## Health_statusCancer                     -0.5179  0.6075
## Health_statusCardiovascular condition    -0.5919  0.2292
## Health_statusChronic pain                -0.3160  0.4258
## Health_statusFibromyalgia                -0.4987  0.6457
## Health_statusNeurological condition      -0.8985 -0.0155  *
## Health_statusObstructive Sleep Apnea     -0.6267  0.5177
## Health_statusOlder adults                -0.7454  0.1616
## Health_statusPost-Partum Women           -0.5823  0.5787
## Health_statusPulmonary condition         -1.1922 -0.2148  **
## Health_statusSurgery                     -0.4522  0.3844
## Health_statusYoung adults                -0.4682  0.7122
##
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

The Test of Moderators revealed a significant difference between subgroups, $F_{12,70} = 3.107$, with $p = 0.0014$.

Note that the model results are printed within a meta-regression framework. This means that we cannot directly extract the estimates in order to obtain the pooled effect sizes within subgroups. The first value, the intercept (intrcpt), shows the z value when the health status was acute pain ($z = -0.129$). The effect in the other groups can be obtained by adding their estimate to the one of the intercept. Thus, the effect in the arthritis group is $z = -0.1290 - 0.1295 = -0.258$, and the one in the Cancer group is $z = -0.1290 + 0.0448 = -0.0838$. The same is true for the upper and lower confidence intervals. These are also Fisher z scores.

```
Health_stat_meta <- update(m, subgroup = Health_status, tau.common = TRUE)
Health_stat_meta
```

```
## Review:      Primary meta-analysis using metacor
##
## Number of studies: n = 63
## Number of estimates: k = 83
## Number of observations: o = 12278
##
##              COR              95%-CI      t  p-value
## Random effects model -0.1931 [-0.2579; -0.1266] -5.70 < 0.0001
## Prediction interval      [-0.6051; 0.3004]
##
## Quantifying heterogeneity:
## tau^2.1 = 0.0576 [0.0227; 0.0950]; tau.1 = 0.2399 [0.1508; 0.3082] (between cluster)
## tau^2.2 = 0.0058 [0.0000; 0.0398]; tau.2 = 0.0762 [0.0000; 0.1995] (within cluster)
## I^2 = 85.5% [82.6%; 87.9%]; H = 2.62 [2.40; 2.87]
##
## Quantifying residual heterogeneity:
## tau^2 = 0.0392; tau = 0.1980; I^2 = 80.6%; H = 2.27
##
## Test of heterogeneity:
##      Q d.f.  p-value
```

```

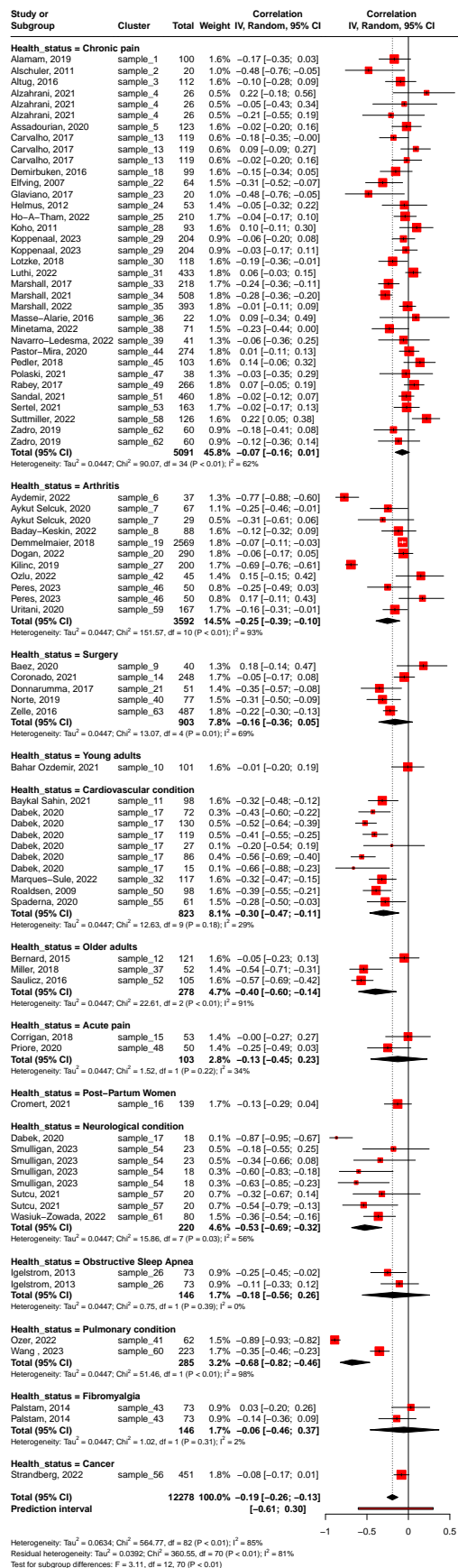
## 564.77 82 < 0.0001
##
## Results for subgroups (random effects model):
##
##          k      COR      95%-CI  tau^2
## Health_status = Chronic pain      35 -0.0739 [-0.1600; 0.0132] 0.0447
## Health_status = Arthritis         11 -0.2529 [-0.3913; -0.1032] 0.0447
## Health_status = Surgery            5 -0.1615 [-0.3585; 0.0494] 0.0447
## Health_status = Young adults       1 -0.0070 [-0.4417; 0.4303] 0.0447
## Health_status = Cardiovascular condition 10 -0.3007 [-0.4674; -0.1134] 0.0447
## Health_status = Older adults       3 -0.3977 [-0.6019; -0.1446] 0.0447
## Health_status = Acute pain         2 -0.1283 [-0.4538; 0.2274] 0.0447
## Health_status = Post-Partum Women  1 -0.1300 [-0.5268; 0.3134] 0.0447
## Health_status = Neurological condition 8 -0.5270 [-0.6863; -0.3194] 0.0447
## Health_status = Obstructive Sleep Apnea 2 -0.1814 [-0.5566; 0.2552] 0.0447
## Health_status = Pulmonary condition 2 -0.6818 [-0.8218; -0.4641] 0.0447
## Health_status = Fibromyalgia       2 -0.0554 [-0.4620; 0.3704] 0.0447
## Health_status = Cancer             1 -0.0840 [-0.4748; 0.3345] 0.0447
##
##          tau      Q      I^2
## Health_status = Chronic pain      0.2114  90.07 62.3%
## Health_status = Arthritis         0.2114 151.57 93.4%
## Health_status = Surgery            0.2114  13.07 69.4%
## Health_status = Young adults       0.2114   0.00  --
## Health_status = Cardiovascular condition 0.2114  12.63 28.7%
## Health_status = Older adults       0.2114  22.61 91.2%
## Health_status = Acute pain         0.2114   1.52 34.2%
## Health_status = Post-Partum Women  0.2114   0.00  --
## Health_status = Neurological condition 0.2114  15.86 55.9%
## Health_status = Obstructive Sleep Apnea 0.2114   0.75  0.0%
## Health_status = Pulmonary condition 0.2114  51.46 98.1%
## Health_status = Fibromyalgia       0.2114   1.02  2.2%
## Health_status = Cancer             0.2114   0.00  --
##
## Test for subgroup differences (random effects model):
##          F      d.f. p-value
## Between groups 3.11 12, 70 0.0014
##
## Details on meta-analytical method:
## - Inverse variance method (three-level model)
## - Restricted maximum-likelihood estimator for tau^2
##   (assuming common tau^2 in subgroups)
## - Profile-Likelihood method for confidence interval of tau^2 and tau
## - Random effects confidence interval based on t-distribution (df = 82)
## - Prediction interval based on t-distribution (df = 81)
## - Fisher's z transformation of correlations

```

The test of subgroup differences between health status was conducted on studies comprising people with chronic ($k = 35$) or acute pain ($k = 2$), arthritis ($k = 11$), a cardiovascular condition ($k = 10$), a neurological condition ($k = 8$), surgery ($k = 5$), older age ($k = 3$), obstructive sleep apnea ($k = 2$), a pulmonary condition ($k = 2$), fibromyalgia ($k = 2$), cancer ($k = 1$), as well as in post-partum women ($k = 1$) and healthy young adults ($k = 1$). We found a statistical moderating effect of health status ($p = 0.0014$). The relationship between kinesophobia and physical activity was statistically significant only in studies that included participants with cardiac condition ($r = -0.30$; 95CI: -0.47 to -0.11), arthritis ($r = -0.25$; 95CI: -0.39 to -0.10), a neurologic condition ($r = -0.53$; 95CI: -0.69 to -0.32), a pulmonary condition ($r = -0.68$; 95CI: -0.82 to -0.46), or older adults ($r = -0.40$; 95CI: -0.60 to -0.14). We found no evidence of an association

between kinesiophobia and physical activity in studies that included participants with chronic pain ($r = -0.07$; 95CI: -0.16 to 0.01) or acute pain ($r = -0.13$; 95CI: -0.45 to 0.23). Statistical heterogeneity was higher in the studies comprising people with a pulmonary condition ($I^2 = 98.1\%$), arthritis ($I^2 = 93.4\%$), or older adults ($I^2 = 91.2\%$) than in the studies comprising people with a cardiac ($I^2 = 28.7\%$) or neurologic condition ($I^2 = 55.9\%$).

```
forest(Health_stat_meta,
       layout = "Revman5",
       common = FALSE,
       xlim = c(-1.0, 0.6),
       prediction = TRUE,
       fs.hetstat = 10,
       col.subgroup = "black",
       addrows.below.overall = 2)
```




```
png(file = "Health condition forest plot.png",
     width = 2800, height = 7800, res = 300)
```

```
forest(Health_stat_meta,
       layout = "Revman5",
       sortvar = -TE,
       common = FALSE,
       xlim = c(-1.0, 0.6),
       prediction = TRUE,
       fs.hetstat = 10,
       col.subgroup = "black",
       addrows.below.overall = 2)
```

```
dev.off()
```

```
## pdf
```

```
## 2
```

Subgroup analysis by physical activity measure: device vs self-report

```
Kinhob_r$PA_objectivity <- as.factor(Kinhob_r$PA_objectivity)
```

```
PAobj_subg <- rma.mv(yi = z,
                    V = var.z,
                    slab = author,
                    data = Kinhob_r,
                    random = ~ 1 | author/cor_id,
                    test = "t",
                    method = "REML",
                    mods = ~ PA_objectivity)
```

```
summary(PAobj_subg)
```

```
##
```

```
## Multivariate Meta-Analysis Model (k = 83; method: REML)
```

```
##
```

```
##   logLik  Deviance      AIC      BIC      AICc
## -10.7304  21.4608   29.4608   39.0386   29.9871
```

```
##
```

```
## Variance Components:
```

```
##
```

```
##      estim    sqrt  nlvls  fixed      factor
## sigma^2.1  0.0623  0.2495   63    no      author
## sigma^2.2  0.0015  0.0392   83    no  author/cor_id
```

```
##
```

```
## Test for Residual Heterogeneity:
```

```
## QE(df = 81) = 558.5318, p-val < .0001
```

```
##
```

```
## Test of Moderators (coefficient 2):
```

```
## F(df1 = 1, df2 = 81) = 1.9042, p-val = 0.1714
```

```
##
```

```
## Model Results:
```

```
##
```

```
##              estimate      se    tval  df    pval
```

```
## intrcpt -0.1323 0.0573 -2.3106 81 0.0234
## PA_objectivitySelf-reported measure -0.0896 0.0649 -1.3799 81 0.1714
## ci.lb ci.ub
## intrcpt -0.2462 -0.0184 *
## PA_objectivitySelf-reported measure -0.2188 0.0396
##
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

The effect the objective measures group is $z = -0.132$, and in the self-report group it is $z = -0.132 - .090 = -0.222$.

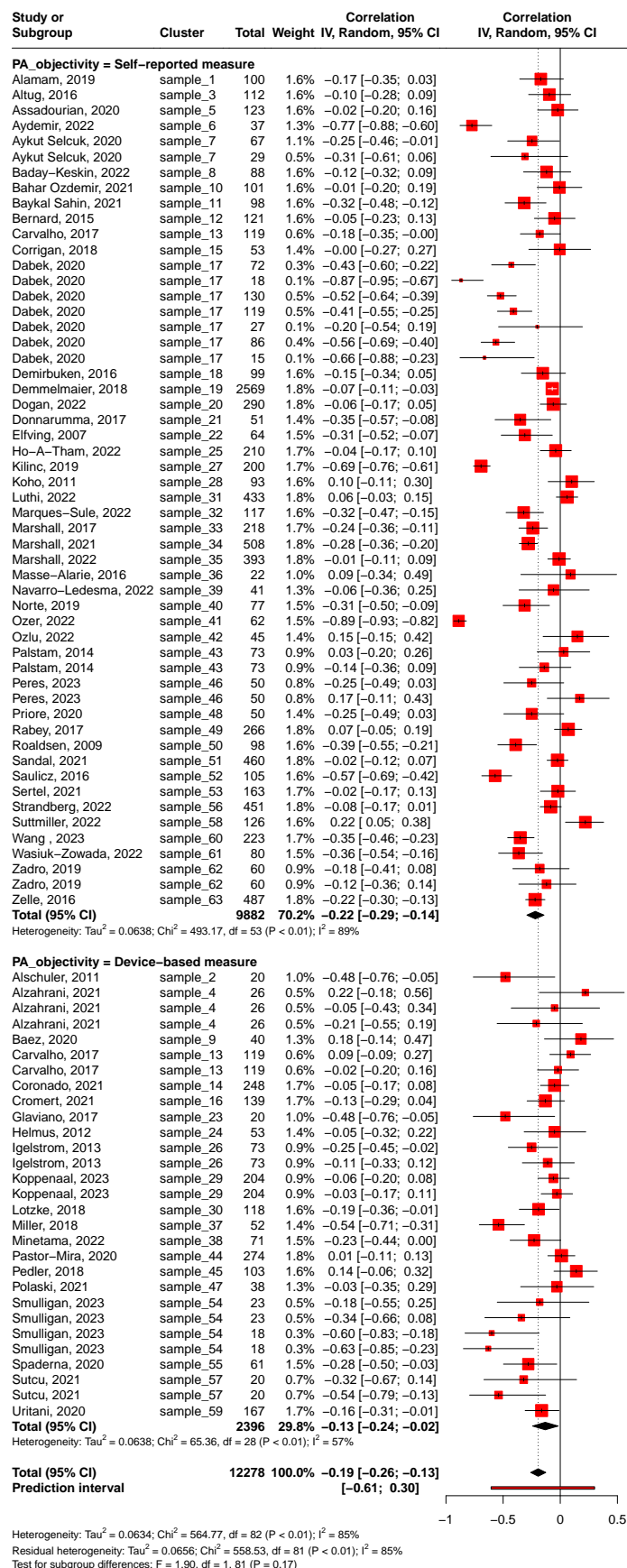
```
PAobj_meta <- update(m, subgroup = PA_objectivity, tau.common = TRUE)
PAobj_meta
```

```
## Review: Primary meta-analysis using metacor
##
## Number of studies: n = 63
## Number of estimates: k = 83
## Number of observations: o = 12278
##
## COR 95%-CI t p-value
## Random effects model -0.1931 [-0.2579; -0.1266] -5.70 < 0.0001
## Prediction interval [-0.6051; 0.3004]
##
## Quantifying heterogeneity:
## tau^2.1 = 0.0576 [0.0227; 0.0950]; tau.1 = 0.2399 [0.1508; 0.3082] (between cluster)
## tau^2.2 = 0.0058 [0.0000; 0.0398]; tau.2 = 0.0762 [0.0000; 0.1995] (within cluster)
## I^2 = 85.5% [82.6%; 87.9%]; H = 2.62 [2.40; 2.87]
##
## Quantifying residual heterogeneity:
## tau^2 = 0.0656; tau = 0.2560; I^2 = 85.5%; H = 2.63
##
## Test of heterogeneity:
## Q d.f. p-value
## 564.77 82 < 0.0001
##
## Results for subgroups (random effects model):
## k COR 95%-CI tau^2
## PA_objectivity = Self-reported measure 54 -0.2183 [-0.2919; -0.1422] 0.0638
## PA_objectivity = Device-based measure 29 -0.1315 [-0.2414; -0.0184] 0.0638
## tau Q I^2
## PA_objectivity = Self-reported measure 0.2526 493.17 89.3%
## PA_objectivity = Device-based measure 0.2526 65.36 57.2%
##
## Test for subgroup differences (random effects model):
## F d.f. p-value
## Between groups 1.90 1, 81 0.1714
##
## Details on meta-analytical method:
## - Inverse variance method (three-level model)
## - Restricted maximum-likelihood estimator for tau^2
## (assuming common tau^2 in subgroups)
## - Profile-Likelihood method for confidence interval of tau^2 and tau
## - Random effects confidence interval based on t-distribution (df = 82)
```

```
## - Prediction interval based on t-distribution (df = 81)
## - Fisher's z transformation of correlations
```

The test of subgroup differences between self-reported ($k = 54$) and device-based ($k = 29$) measures of physical activity showed no evidence of a moderating effect of the type of physical activity measure ($p = 0.171$). Both self-reported measures ($r = -0.22$; 95CI: -0.29 to -0.14; $I^2 = 89.3\%$) and device-based measures ($r = -0.13$; 95CI: -0.24 to -0.02; $I^2 = 57.2\%$) showed a negative association between kinesiophobia and physical activity.

```
forest(PAobj_meta,
       layout = "RevMan5",
       common = FALSE,
       xlim = c(-1.0, 0.6),
       prediction = TRUE,
       fs.hetstat = 10,
       col.subgroup = "black",
       addrows.below.overall = 2)
```



```
png(file = "PA objectivity forestplot.png",
     width = 2800, height = 5800, res = 300)
```

```
forest(PAobj_meta,
       layout = "RevMan5",
       common = FALSE,
       xlim = c(-1.0, 0.6),
       prediction = TRUE,
       fs.hetstat = 10,
       col.subgroup = "black",
       addrows.below.overall = 2)
```

```
dev.off()
```

```
## pdf
## 2
```

Subgroup analysis by physical activity measurement instruments

```
Kinhob_r$PA_measure <- as.factor(Kinhob_r$PA_measure)
```

```
PAmeas_subg <- rma.mv(yi = z,
                     V = var.z,
                     slab = author,
                     data = Kinhob_r,
                     random = ~ 1 | author/cor_id,
                     test = "t",
                     method = "REML",
                     mods = ~ PA_measure)
```

```
summary(PAmeas_subg)
```

```
##
## Multivariate Meta-Analysis Model (k = 83; method: REML)
##
##   logLik  Deviance      AIC      BIC      AICc
## -6.4184   12.8369   48.8369   88.5213   63.0869
##
## Variance Components:
##
##           estim      sqrt  nlvls  fixed      factor
## sigma^2.1  0.0626  0.2503    63    no      author
## sigma^2.2  0.0000  0.0000    83    no  author/cor_id
##
## Test for Residual Heterogeneity:
## QE(df = 67) = 490.8621, p-val < .0001
##
## Test of Moderators (coefficients 2:16):
## F(df1 = 15, df2 = 67) = 1.3310, p-val = 0.2092
##
## Model Results:
##
##           estimate      se      tval  df      pval      ci.lb
## intrcpt          -0.1711  0.0624  -2.7416  67  0.0078  -0.2957
```

```

## PA_measureBHPAQ      -0.1803  0.0940  -1.9184  67  0.0593  -0.3678
## PA_measureDiary       0.1491  0.2736   0.5449  67  0.5876  -0.3970
## PA_measureGLTEQ      -0.0300  0.2120  -0.1417  67  0.8877  -0.4531
## PA_measureGPAQ       0.1792  0.2155   0.8313  67  0.4087  -0.2510
## PA_measureIPAQ      -0.1151  0.0890  -1.2934  67  0.2003  -0.2926
## PA_measureJPAS       0.3937  0.2732   1.4409  67  0.1543  -0.1517
## PA_measureLTPAI      0.1156  0.2714   0.4261  67  0.6714  -0.4261
## PA_measureMLTPAQ     0.0989  0.1959   0.5048  67  0.6154  -0.2921
## PA_measurePASE       0.1501  0.2698   0.5564  67  0.5798  -0.3884
## PA_measurePedometer  0.1501  0.1716   0.8749  67  0.3847  -0.1923
## PA_measureQuestionnaire 0.0213  0.1437   0.1485  67  0.8824  -0.2654
## PA_measureRAPAQ      0.0198  0.2744   0.0722  67  0.9426  -0.5279
## PA_measureSGPALS     -0.0167  0.1499  -0.1117  67  0.9114  -0.3159
## PA_measureSQUASH     0.1292  0.2778   0.4652  67  0.6433  -0.4253
## PA_measureUCLA      -0.8566  0.3097  -2.7656  67  0.0073  -1.4749
##          ci.lb          ci.ub
## intrcpt          -0.0465    **
## PA_measureBHPAQ      0.0073    .
## PA_measureDiary      0.6952
## PA_measureGLTEQ      0.3930
## PA_measureGPAQ       0.6094
## PA_measureIPAQ       0.0625
## PA_measureJPAS       0.9391
## PA_measureLTPAI      0.6574
## PA_measureMLTPAQ     0.4898
## PA_measurePASE       0.6886
## PA_measurePedometer  0.4925
## PA_measureQuestionnaire 0.3081
## PA_measureRAPAQ      0.5676
## PA_measureSGPALS     0.2824
## PA_measureSQUASH     0.6837
## PA_measureUCLA      -0.2384    **
##
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

```

```

PAm <- update(m,
  subgroup = PA_measure,
  tau.common = TRUE)
PAm

```

```

## Review:      Primary meta-analysis using metacor
##
## Number of studies: n = 63
## Number of estimates: k = 83
## Number of observations: o = 12278
##
##              COR              95%-CI      t  p-value
## Random effects model -0.1931 [-0.2579; -0.1266] -5.70 < 0.0001
## Prediction interval      [-0.6051;  0.3004]
##
## Quantifying heterogeneity:
## tau^2.1 = 0.0576 [0.0227; 0.0950]; tau.1 = 0.2399 [0.1508; 0.3082] (between cluster)
## tau^2.2 = 0.0058 [0.0000; 0.0398]; tau.2 = 0.0762 [0.0000; 0.1995] (within cluster)
## I^2 = 85.5% [82.6%; 87.9%]; H = 2.62 [2.40; 2.87]

```

```

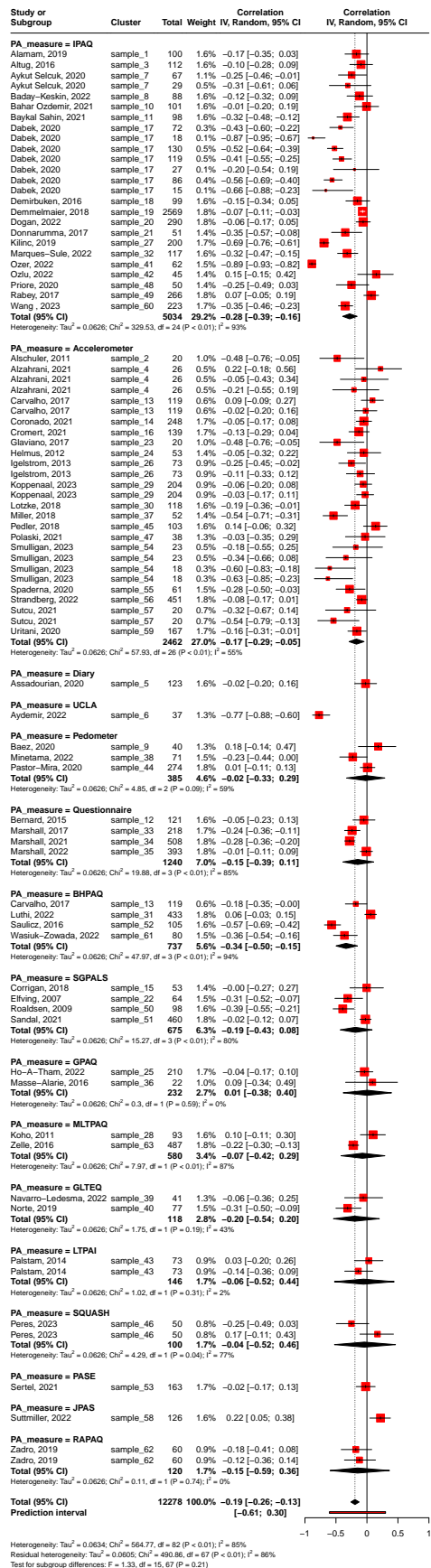
##
## Quantifying residual heterogeneity:
## tau^2 = 0.0605; tau = 0.2460; I^2 = 86.4%; H = 2.71
##
## Test of heterogeneity:
##      Q d.f.  p-value
## 564.77   82 < 0.0001
##
## Results for subgroups (random effects model):
##
##      k      COR      95%-CI  tau^2    tau      Q
## PA_measure = IPAQ      25 -0.2786 [-0.3908; -0.1583] 0.0626 0.2503 329.53
## PA_measure = Accelerometer 27 -0.1695 [-0.2874; -0.0465] 0.0626 0.2503 57.93
## PA_measure = Diary      1 -0.0220 [-0.5033; 0.4697] 0.0626 0.2503 0.00
## PA_measure = UCLA      1 -0.7730 [-0.9265; -0.3987] 0.0626 0.2503 0.00
## PA_measure = Pedometer  3 -0.0210 [-0.3275; 0.2895] 0.0626 0.2503 4.85
## PA_measure = Questionnaire 4 -0.1487 [-0.3868; 0.1081] 0.0626 0.2503 19.88
## PA_measure = BHPAQ      4 -0.3376 [-0.4988; -0.1538] 0.0626 0.2503 47.97
## PA_measure = SGPALS     4 -0.1857 [-0.4300; 0.0840] 0.0626 0.2503 15.27
## PA_measure = GPAQ       2 0.0081 [-0.3831; 0.3968] 0.0626 0.2503 0.30
## PA_measure = MLTPAQ     2 -0.0721 [-0.4160; 0.2898] 0.0626 0.2503 7.97
## PA_measure = GLTEQ      2 -0.1985 [-0.5409; 0.2004] 0.0626 0.2503 1.75
## PA_measure = LTPAI      2 -0.0554 [-0.5246; 0.4397] 0.0626 0.2503 1.02
## PA_measure = SQUASH     2 -0.0418 [-0.5243; 0.4609] 0.0626 0.2503 4.29
## PA_measure = PASE       1 -0.0210 [-0.4967; 0.4644] 0.0626 0.2503 0.00
## PA_measure = JPAS       1 0.2190 [-0.2990; 0.6373] 0.0626 0.2503 0.00
## PA_measure = RAPAQ      2 -0.1501 [-0.5945; 0.3645] 0.0626 0.2503 0.11
##
##      I^2
## PA_measure = IPAQ      92.7%
## PA_measure = Accelerometer 55.1%
## PA_measure = Diary      --
## PA_measure = UCLA      --
## PA_measure = Pedometer  58.8%
## PA_measure = Questionnaire 84.9%
## PA_measure = BHPAQ      93.7%
## PA_measure = SGPALS     80.4%
## PA_measure = GPAQ       0.0%
## PA_measure = MLTPAQ     87.4%
## PA_measure = GLTEQ      42.7%
## PA_measure = LTPAI      2.2%
## PA_measure = SQUASH     76.7%
## PA_measure = PASE       --
## PA_measure = JPAS       --
## PA_measure = RAPAQ      0.0%
##
## Test for subgroup differences (random effects model):
##      F    d.f.  p-value
## Between groups 1.33 15, 67 0.2092
##
## Details on meta-analytical method:
## - Inverse variance method (three-level model)
## - Restricted maximum-likelihood estimator for tau^2
##   (assuming common tau^2 in subgroups)
## - Profile-Likelihood method for confidence interval of tau^2 and tau
## - Random effects confidence interval based on t-distribution (df = 82)

```

```
## - Prediction interval based on t-distribution (df = 81)
## - Fisher's z transformation of correlations
```

We also found no evidence of a moderating effect of physical activity instruments ($p = 0.209$), physical activity outcome ($p = 0.685$), or kinesiophobia instrument ($p = 0.452$).

```
forest(PAm,
       layout = "Revman5",
       common = FALSE,
       xlim = c(-1.0, 1.0),
       prediction = TRUE,
       fs.hetstat = 10,
       col.subgroup = 'black',
       addrows.below.overall = 2)
```

```
png(file = "Physical activity measure forestplot.png",
     width = 2800, height = 8400, res = 300)
```

```
forest(PAm,
       layout = "Revman5",
       sortvar = -TE,
       common = FALSE,
       xlim = c(-1.0, 1.0),
       prediction = TRUE,
       fs.hetstat = 10,
       col.subgroup = 'black',
       addrows.below.overall = 2)
```

```
dev.off()
```

```
## pdf
```

```
## 2
```

Subgroup analysis by physical activity outcome

```
Kinhob_r$PA_outcome <- as.factor(Kinhob_r$PA_outcome)
```

```
PAout_subg <- rma.mv(yi = z,
                    V = var.z,
                    slab = author,
                    data = Kinhob_r,
                    random = ~ 1 | author/cor_id,
                    test = "t",
                    method = "REML",
                    mods = ~ PA_outcome)
```

```
summary(PAout_subg)
```

```
##
```

```
## Multivariate Meta-Analysis Model (k = 83; method: REML)
```

```
##
```

```
##   logLik  Deviance      AIC      BIC      AICc
## -11.7574  23.5148  39.5148  58.2652  41.6324
```

```
##
```

```
## Variance Components:
```

```
##
```

```
##           estim  sqrt  nlvls  fixed      factor
## sigma^2.1  0.0641  0.2531    63    no      author
## sigma^2.2  0.0017  0.0416    83    no author/cor_id
```

```
##
```

```
## Test for Residual Heterogeneity:
```

```
## QE(df = 77) = 553.8532, p-val < .0001
```

```
##
```

```
## Test of Moderators (coefficients 2:6):
```

```
## F(df1 = 5, df2 = 77) = 0.7209, p-val = 0.6098
```

```
##
```

```
## Model Results:
```

```
##
```

```
##           estimate      se    tval  df    pval    ci.lb    ci.ub
```

```

## intrcpt          -0.1085  0.0718 -1.5110  77  0.1349 -0.2514  0.0345
## PA_outcomeCounts/min -0.0372  0.1074 -0.3465  77  0.7299 -0.2510  0.1766
## PA_outcomeKcal/day   -0.2936  0.2202 -1.3331  77  0.1864 -0.7321  0.1449
## PA_outcomeMET-min/week -0.0917  0.0896 -1.0232  77  0.3094 -0.2702  0.0868
## PA_outcomeScore      -0.1353  0.0884 -1.5313  77  0.1298 -0.3113  0.0406
## PA_outcomeSteps/day  -0.0756  0.0933 -0.8104  77  0.4202 -0.2614  0.1102
##
## intrcpt
## PA_outcomeCounts/min
## PA_outcomeKcal/day
## PA_outcomeMET-min/week
## PA_outcomeScore
## PA_outcomeSteps/day
##
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

PAout <- update(m,
  subgroup = PA_outcome,
  tau.common = TRUE)
PAout

## Review:      Primary meta-analysis using metacor
##
## Number of studies: n = 63
## Number of estimates: k = 83
## Number of observations: o = 12278
##
##              COR              95%-CI      t  p-value
## Random effects model -0.1931 [-0.2579; -0.1266] -5.70 < 0.0001
## Prediction interval      [-0.6051;  0.3004]
##
## Quantifying heterogeneity:
## tau^2.1 = 0.0576 [0.0227; 0.0950]; tau.1 = 0.2399 [0.1508; 0.3082] (between cluster)
## tau^2.2 = 0.0058 [0.0000; 0.0398]; tau.2 = 0.0762 [0.0000; 0.1995] (within cluster)
## I^2 = 85.5% [82.6%; 87.9%]; H = 2.62 [2.40; 2.87]
##
## Quantifying residual heterogeneity:
## tau^2 = 0.0684; tau = 0.2616; I^2 = 86.1%; H = 2.68
##
## Test of heterogeneity:
##      Q d.f.  p-value
## 564.77  82 < 0.0001
##
## Results for subgroups (random effects model):
##              k      COR              95%-CI  tau^2    tau      Q
## PA_outcome = MET-min/week  35 -0.1975 [-0.2992; -0.0915] 0.0658 0.2565 371.71
## PA_outcome = Counts/min    5 -0.1447 [-0.3282;  0.0494] 0.0658 0.2565  4.37
## PA_outcome = Steps/day     13 -0.1820 [-0.3256; -0.0302] 0.0658 0.2565 27.04
## PA_outcome = Active time   12 -0.1080 [-0.2462;  0.0345] 0.0658 0.2565 46.81
## PA_outcome = Score         16 -0.2391 [-0.3583; -0.1122] 0.0658 0.2565 102.61
## PA_outcome = Kcal/day       2 -0.3817 [-0.6747;  0.0153] 0.0658 0.2565  1.32
##
##              I^2
## PA_outcome = MET-min/week 90.9%
## PA_outcome = Counts/min   8.4%

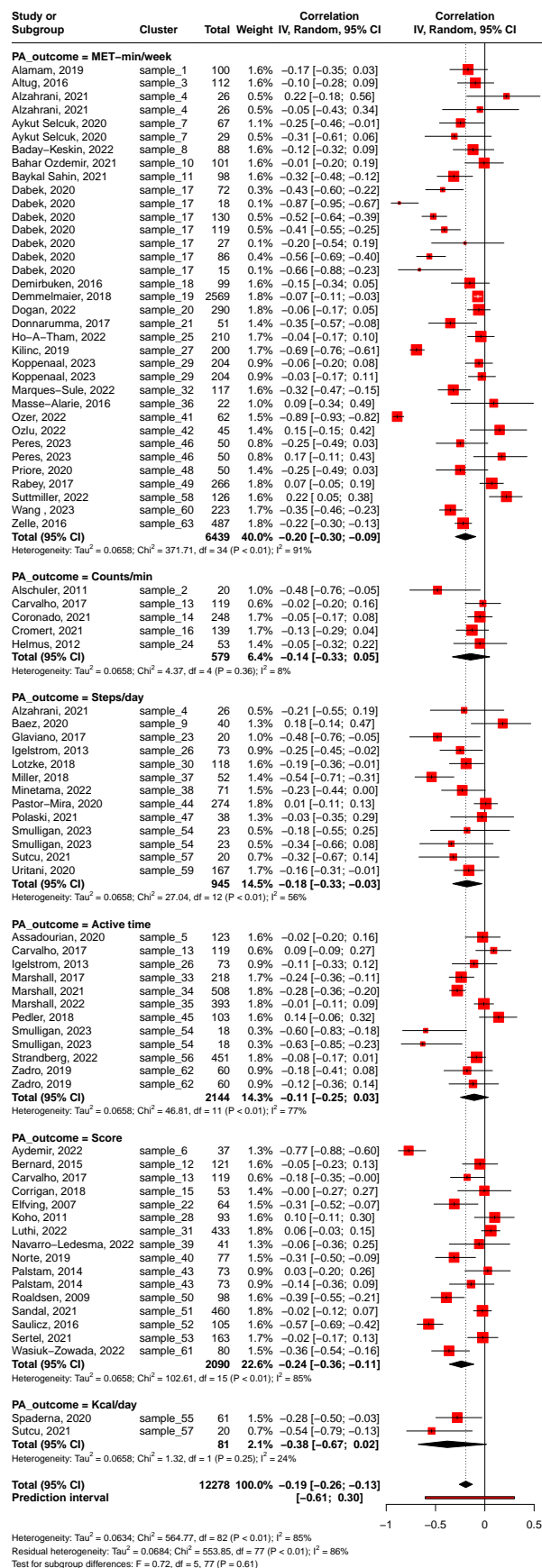
```

```

## PA_outcome = Steps/day      55.6%
## PA_outcome = Active time   76.5%
## PA_outcome = Score         85.4%
## PA_outcome = Kcal/day      24.1%
##
## Test for subgroup differences (random effects model):
##           F    d.f. p-value
## Between groups 0.72 5, 77 0.6098
##
## Details on meta-analytical method:
## - Inverse variance method (three-level model)
## - Restricted maximum-likelihood estimator for tau^2
##   (assuming common tau^2 in subgroups)
## - Profile-Likelihood method for confidence interval of tau^2 and tau
## - Random effects confidence interval based on t-distribution (df = 82)
## - Prediction interval based on t-distribution (df = 81)
## - Fisher's z transformation of correlations

forest(PAout,
       layout = "RevMan5",
       common = FALSE,
       xlim = c(-1.0, 0.6),
       prediction = TRUE,
       fs.hetstat = 10,
       col.subgroup = 'black',
       addrows.below.overall = 2)

```



```
png(file = "PA outcome forestplot.png",
     width = 2800, height = 6400, res = 300)
```

```
forest(PAout,
       layout = "RevMan5",
       sortvar = -TE,
       common = FALSE,
       xlim = c(-1.0, 0.6),
       prediction = TRUE,
       fs.hetstat = 10,
       col.subgroup = 'black',
       addrows.below.overall = 2)
```

```
dev.off()
```

```
## pdf
```

```
## 2
```

Subgroup analysis by kinesiophobia measure

```
Kinhob_r$Kinesiophobia_measure <- as.factor(Kinhob_r$Kinesiophobia_measure)
```

```
Kines_subg <- rma.mv(yi = z,
                    V = var.z,
                    slab = author,
                    data = Kinhob_r,
                    random = ~ 1 | author/cor_id,
                    test = "t",
                    method = "REML",
                    mods = ~ Kinesiophobia_measure)
```

```
summary(Kines_subg)
```

```
##
```

```
## Multivariate Meta-Analysis Model (k = 83; method: REML)
```

```
##
```

```
##   logLik  Deviance      AIC      BIC      AICc
## -9.5324  19.0648  45.0648  74.6615  51.3407
```

```
##
```

```
## Variance Components:
```

```
##
```

```
##           estim  sqrt  nlvls  fixed      factor
## sigma^2.1  0.0531  0.2305    63    no      author
## sigma^2.2  0.0095  0.0976    83    no  author/cor_id
```

```
##
```

```
## Test for Residual Heterogeneity:
```

```
## QE(df = 72) = 429.2840, p-val < .0001
```

```
##
```

```
## Test of Moderators (coefficients 2:11):
```

```
## F(df1 = 10, df2 = 72) = 0.9998, p-val = 0.4520
```

```
##
```

```
## Model Results:
```

```
##
```

```
##           estimate      se      tval  df      pval      ci.lb
```

```
## intrcpt -0.3654 0.2592 -1.4098 72 0.1629 -0.8822
## Kinesiophobia_measureBFOMS0 0.2010 0.3687 0.5451 72 0.5874 -0.5340
## Kinesiophobia_measureFABQ 0.2387 0.2694 0.8860 72 0.3785 -0.2984
## Kinesiophobia_measureFActS 0.0778 0.3835 0.2028 72 0.8399 -0.6867
## Kinesiophobia_measureKCS -0.2821 0.3737 -0.7549 72 0.4528 -1.0270
## Kinesiophobia_measureTSK_11 0.3237 0.2765 1.1704 72 0.2457 -0.2276
## Kinesiophobia_measureTSK_12 0.1591 0.3509 0.4535 72 0.6515 -0.5404
## Kinesiophobia_measureTSK_13 0.2081 0.3035 0.6858 72 0.4951 -0.3969
## Kinesiophobia_measureTSK_14 0.2812 0.3634 0.7739 72 0.4415 -0.4432
## Kinesiophobia_measureTSK_17 0.1317 0.2636 0.4995 72 0.6190 -0.3938
## Kinesiophobia_measureTSK_Heart -0.1916 0.3527 -0.5432 72 0.5887 -0.8947
## ci.ub
## intrcpt 0.1513
## Kinesiophobia_measureBFOMS0 0.9359
## Kinesiophobia_measureFABQ 0.7758
## Kinesiophobia_measureFActS 0.8423
## Kinesiophobia_measureKCS 0.4628
## Kinesiophobia_measureTSK_11 0.8749
## Kinesiophobia_measureTSK_12 0.8587
## Kinesiophobia_measureTSK_13 0.8132
## Kinesiophobia_measureTSK_14 1.0057
## Kinesiophobia_measureTSK_17 0.6571
## Kinesiophobia_measureTSK_Heart 0.5115
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
Kin_sub <- update(m,
  subgroup = Kinesiophobia_measure,
  tau.common = TRUE)
```

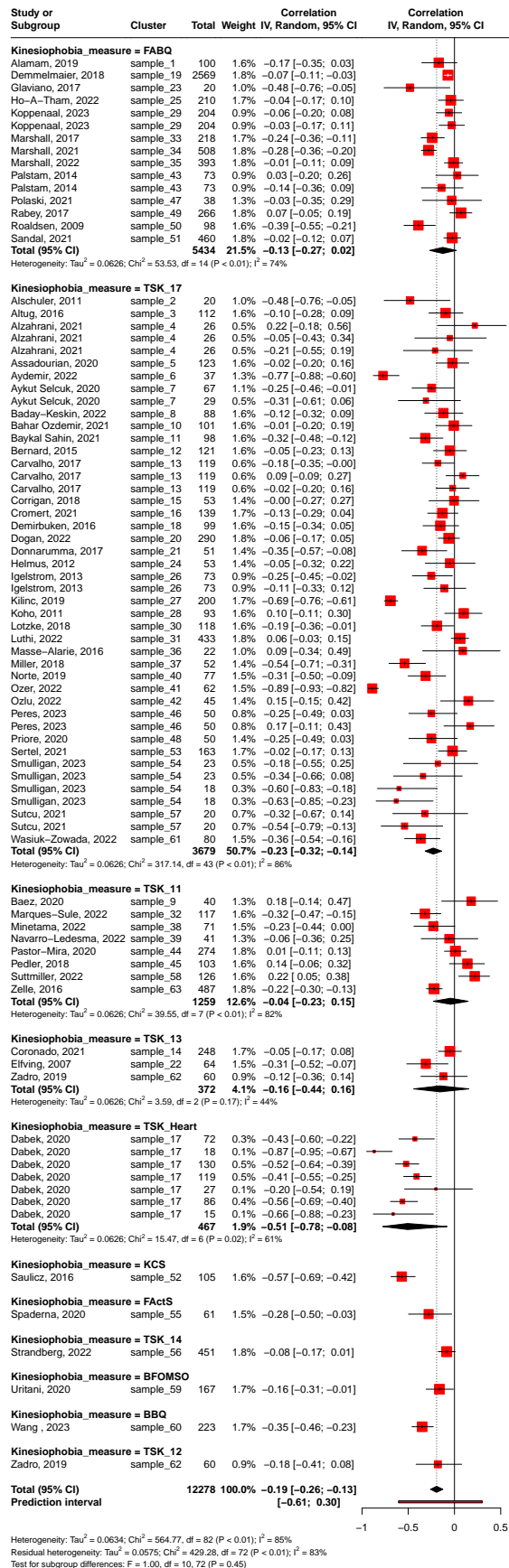
```
## Warning: Single-level factor(s) found in 'random' argument. Corresponding
## 'sigma2' value(s) fixed to 0.
```

```
Kin_sub
```

```
## Review: Primary meta-analysis using metacor
##
## Number of studies: n = 63
## Number of estimates: k = 83
## Number of observations: o = 12278
##
## COR 95%-CI t p-value
## Random effects model -0.1931 [-0.2579; -0.1266] -5.70 < 0.0001
## Prediction interval [-0.6051; 0.3004]
##
## Quantifying heterogeneity:
## tau^2.1 = 0.0576 [0.0227; 0.0950]; tau.1 = 0.2399 [0.1508; 0.3082] (between cluster)
## tau^2.2 = 0.0058 [0.0000; 0.0398]; tau.2 = 0.0762 [0.0000; 0.1995] (within cluster)
## I^2 = 85.5% [82.6%; 87.9%]; H = 2.62 [2.40; 2.87]
##
## Quantifying residual heterogeneity:
## tau^2 = 0.0575; tau = 0.2397; I^2 = 83.2%; H = 2.44
##
## Test of heterogeneity:
## Q d.f. p-value
```

```
## 564.77 82 < 0.0001
##
## Results for subgroups (random effects model):
##           k      COR           95%-CI  tau^2  tau
## Kinesiophobia_measure = FABQ      15 -0.1261 [-0.2666; 0.0197] 0.0626 0.2503
## Kinesiophobia_measure = TSK_17     44 -0.2296 [-0.3177; -0.1376] 0.0626 0.2503
## Kinesiophobia_measure = TSK_11      8 -0.0418 [-0.2297; 0.1492] 0.0626 0.2503
## Kinesiophobia_measure = TSK_13      3 -0.1560 [-0.4399; 0.1562] 0.0626 0.2503
## Kinesiophobia_measure = TSK_Heart    7 -0.5058 [-0.7755; -0.0800] 0.0626 0.2503
## Kinesiophobia_measure = KCS          1 -0.5700 [-0.8287; -0.1105] 0.0626 0.2503
## Kinesiophobia_measure = FActS        1 -0.2800 [-0.6917; 0.2690] 0.0626 0.2503
## Kinesiophobia_measure = TSK_14        1 -0.0840 [-0.5313; 0.3999] 0.0626 0.2503
## Kinesiophobia_measure = BFOMSO        1 -0.1630 [-0.5961; 0.3436] 0.0626 0.2503
## Kinesiophobia_measure = BBQ           1 -0.3500 [-0.7075; 0.1501] 0.0626 0.2503
## Kinesiophobia_measure = TSK_12        1 -0.2034 [-0.5901; 0.2592] 0.0626 0.2503
##           Q      I^2
## Kinesiophobia_measure = FABQ      53.53 73.8%
## Kinesiophobia_measure = TSK_17    317.14 86.4%
## Kinesiophobia_measure = TSK_11     39.55 82.3%
## Kinesiophobia_measure = TSK_13      3.59 44.2%
## Kinesiophobia_measure = TSK_Heart  15.47 61.2%
## Kinesiophobia_measure = KCS         0.00  --
## Kinesiophobia_measure = FActS       0.00  --
## Kinesiophobia_measure = TSK_14       0.00  --
## Kinesiophobia_measure = BFOMSO       0.00  --
## Kinesiophobia_measure = BBQ          0.00  --
## Kinesiophobia_measure = TSK_12       0.00  --
##
## Test for subgroup differences (random effects model):
##           F      d.f. p-value
## Between groups 1.00 10, 72 0.4520
##
## Details on meta-analytical method:
## - Inverse variance method (three-level model)
## - Restricted maximum-likelihood estimator for tau^2
##   (assuming common tau^2 in subgroups)
## - Profile-Likelihood method for confidence interval of tau^2 and tau
## - Random effects confidence interval based on t-distribution (df = 82)
## - Prediction interval based on t-distribution (df = 81)
## - Fisher's z transformation of correlations

forest(Kin_sub,
  layout = "Revman5",
  common = FALSE,
  xlim = c(-1.0, 0.6),
  prediction = TRUE,
  fs.hetstat = 10,
  col.subgroup = 'black',
  addrows.below.overall = 2)
```

```
png(file = "Kinesiophobia outcome forestplot.png",
     width = 2800, height = 7000, res = 300)
```

```
forest(Kin_sub,
       layout = "RevMan5",
       sortvar = -TE,
       common = FALSE,
       xlim = c(-1.0, 0.6),
       prediction = TRUE,
       fs.hetstat = 10,
       col.subgroup = 'black',
       addrows.below.overall = 2)
```

```
dev.off()
```

```
## pdf
```

```
## 2
```

Meta-analysis: meta-regression

Meta-regression by age

```
m.cor.reg.age <- metareg(m, ~Age)
```

```
## Warning: 11 rows with NAs omitted from model fitting.
```

```
m.cor.reg.age
```

```
##
```

```
## Multivariate Meta-Analysis Model (k = 72; method: REML)
```

```
##
```

```
## Variance Components:
```

```
##
```

```
##          estim      sqrt  nlvls  fixed    factor
```

```
## sigma^2.1 0.0580 0.2407    58    no      .id
```

```
## sigma^2.2 0.0033 0.0577    72    no  .id/.idx
```

```
##
```

```
## Test for Residual Heterogeneity:
```

```
## QE(df = 70) = 429.6753, p-val < .0001
```

```
##
```

```
## Test of Moderators (coefficient 2):
```

```
## F(df1 = 1, df2 = 70) = 0.8879, p-val = 0.3493
```

```
##
```

```
## Model Results:
```

```
##
```

```
##          estimate      se      tval  df    pval    ci.lb    ci.ub
```

```
## intrcpt  -0.0702 0.1323 -0.5309  70  0.5972 -0.3340 0.1936
```

```
## Age      -0.0024 0.0026 -0.9423  70  0.3493 -0.0075 0.0027
```

```
##
```

```
## ---
```

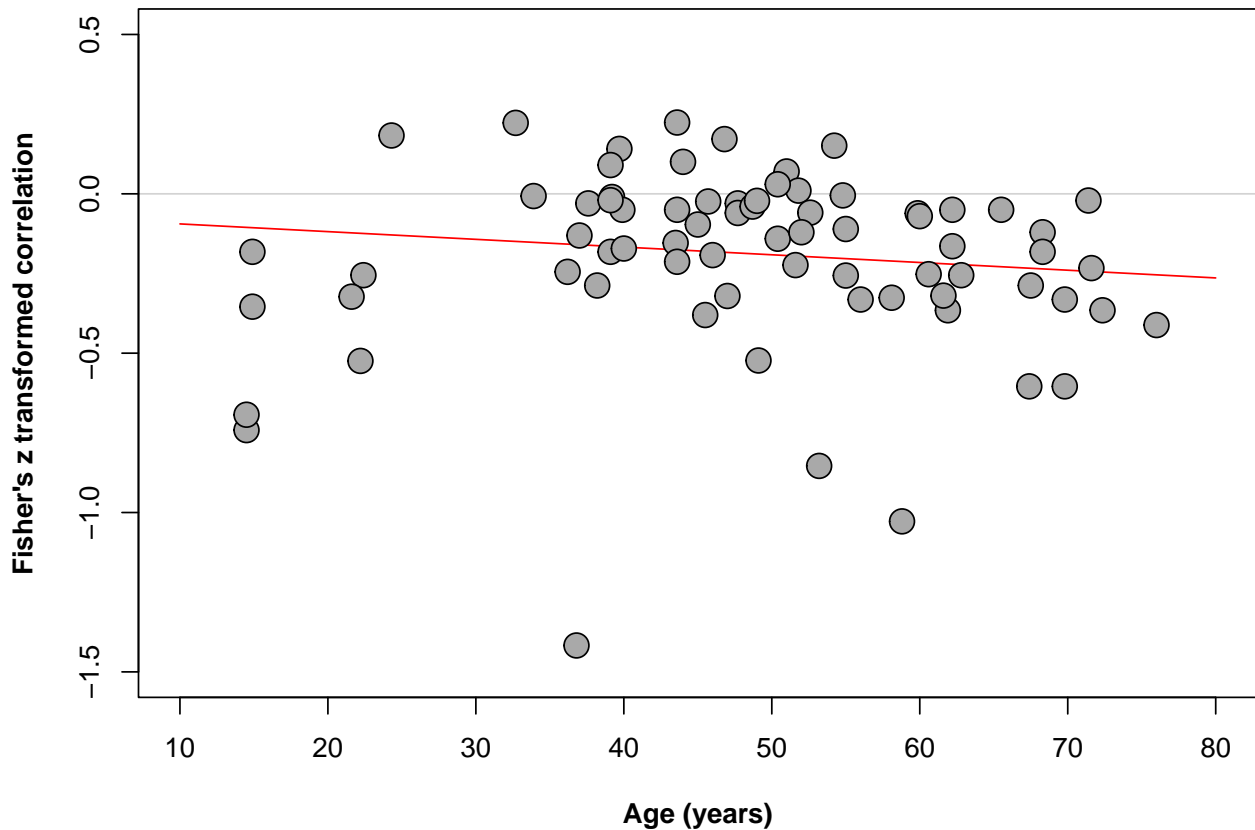
```
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Age did not statistically influence the correlation estimates of the meta-analysis studies (k = 72; p = 0.349).

```

bubble(m.cor.reg.age,
       xlim = c(10, 80),
       ylim = c(-1.5, 0.5),
       xlab = 'Age (years)',
       font.lab = 2,
       studlab = FALSE, # change to TRUE for study labels
       cex = 2,
       cex.studlab = 0.6,
       pos.studlab = 1,
       offset = 0.5,
       col.line = 'red')

```



Meta-regression by proportion of women

```
m.cor.reg.women <- metareg(m, ~Prop_women)
```

```
## Warning: 11 rows with NAs omitted from model fitting.
```

```
m.cor.reg.women
```

```
##
```

```
## Multivariate Meta-Analysis Model (k = 72; method: REML)
```

```
##
```

```
## Variance Components:
```

```
##
```

```
##      estim      sqrt  nlvls  fixed   factor
```

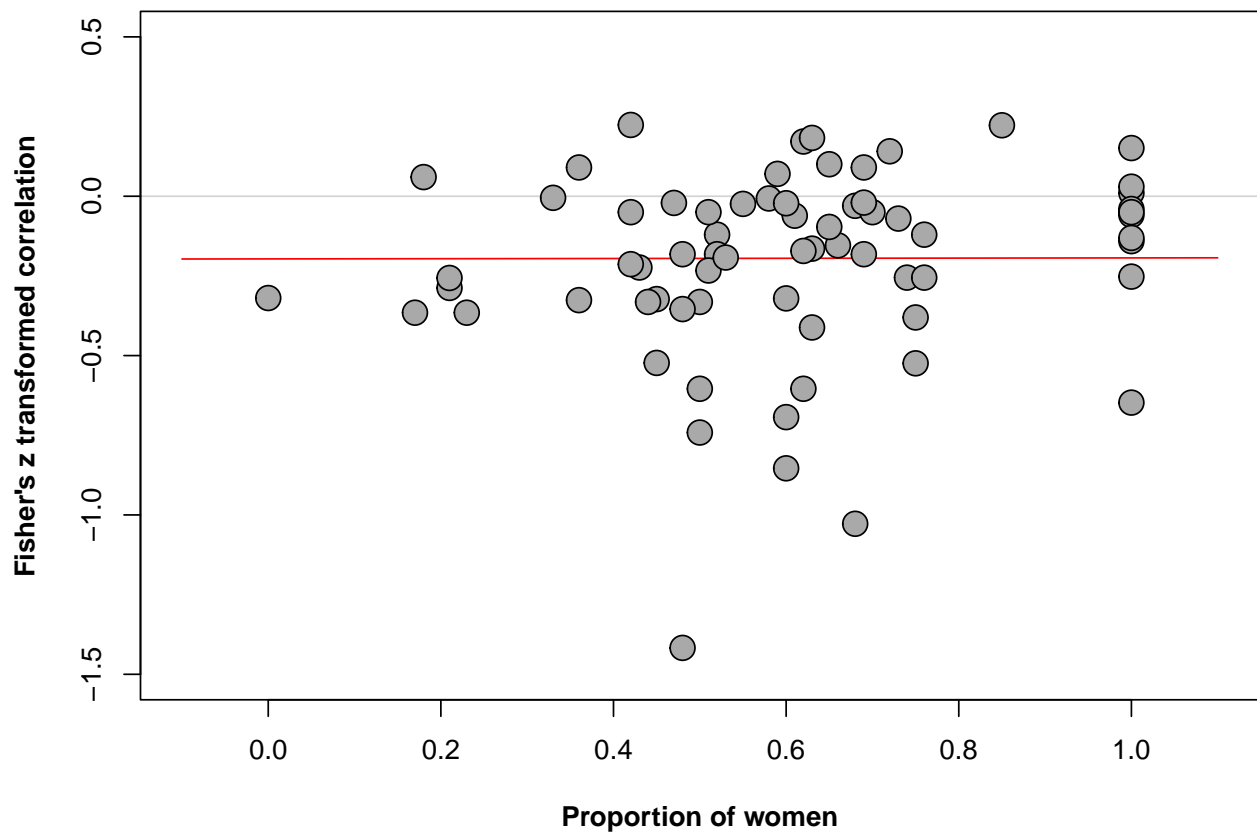
```
## sigma^2.1 0.0633 0.2515   58    no     .id
```

```
## sigma^2.2 0.0043 0.0659   72    no  .id/.idx
```

```
##
## Test for Residual Heterogeneity:
## QE(df = 70) = 449.6260, p-val < .0001
##
## Test of Moderators (coefficient 2):
## F(df1 = 1, df2 = 70) = 0.3528, p-val = 0.5545
##
## Model Results:
##
##           estimate      se      tval  df    pval    ci.lb    ci.ub
## intrcpt      -0.1966  0.0380  -5.1724  70  <.0001   -0.2725  -0.1208 ***
## Prop_women    0.0031  0.0052   0.5939  70  0.5545   -0.0073   0.0135
##
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Similarly, the proportion of women ($k = 72$; $p = 0.555$) and the mean level of pain in the studies ($k = 49$; $p = 0.481$) did not influence correlation estimates.

```
bubble(m.cor.reg.women,
       xlim = c(-0.1, 1.1),
       ylim = c(-1.5, 0.5),
       xlab = 'Proportion of women',
       font.lab = 2,
       studlab = F,
       cex = 2,
       cex.studlab = 0.6,
       pos.studlab = 1,
       offset = 0.5,
       col.line = 'red')
```



Meta-regression by pain

```
m.cor.reg.pain <- metareg(m, ~Pain)
```

```
## Warning: 34 rows with NAs omitted from model fitting.
```

```
m.cor.reg.pain
```

```
##
```

```
## Multivariate Meta-Analysis Model (k = 49; method: REML)
```

```
##
```

```
## Variance Components:
```

```
##
```

```
##      estim      sqrt  nlvls  fixed    factor
```

```
## sigma^2.1 0.0435 0.2085   41    no      .id
```

```
## sigma^2.2 0.0012 0.0349   49    no  .id/.idx
```

```
##
```

```
## Test for Residual Heterogeneity:
```

```
## QE(df = 47) = 274.6816, p-val < .0001
```

```
##
```

```
## Test of Moderators (coefficient 2):
```

```
## F(df1 = 1, df2 = 47) = 0.5039, p-val = 0.4813
```

```
##
```

```
## Model Results:
```

```
##
```

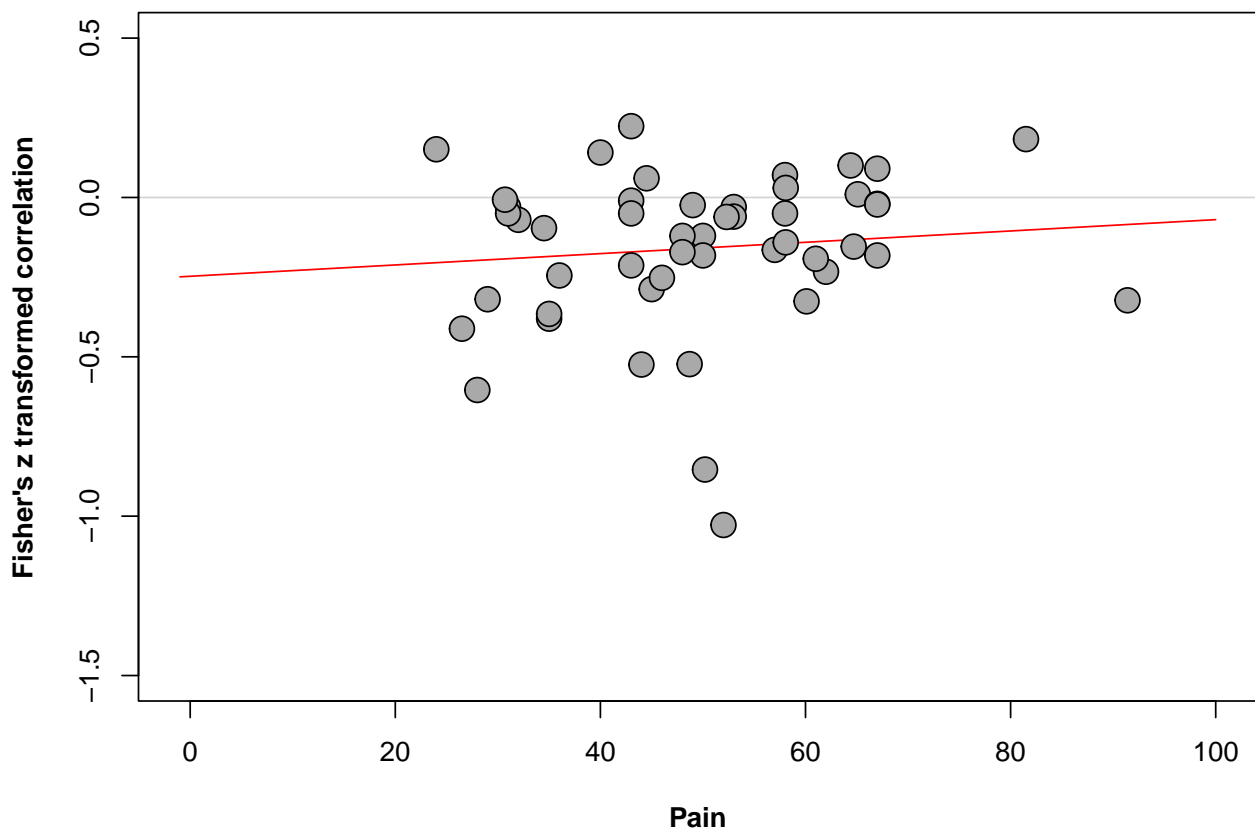
```
##      estimate      se      tval  df      pval      ci.lb      ci.ub
```

```
## intrcpt  -0.2473 0.1285  -1.9252 47      0.0603  -0.5058  0.0111 .
```

```
## Pain      0.0018 0.0025   0.7098 47      0.4813  -0.0033  0.0068
```

```
##
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

bubble(m.cor.reg.pain,
       xlim = c(-1, 100),
       ylim = c(-1.5, 0.5),
       xlab = 'Pain',
       studlab = FALSE,
       font.lab = 2,
       cex = 2,
       cex.studlab = 0.6,
       pos.studlab = 1,
       offset = 0.5,
       col.line = 'red')
```



Sensitivity analysis: meta-regression by axis score

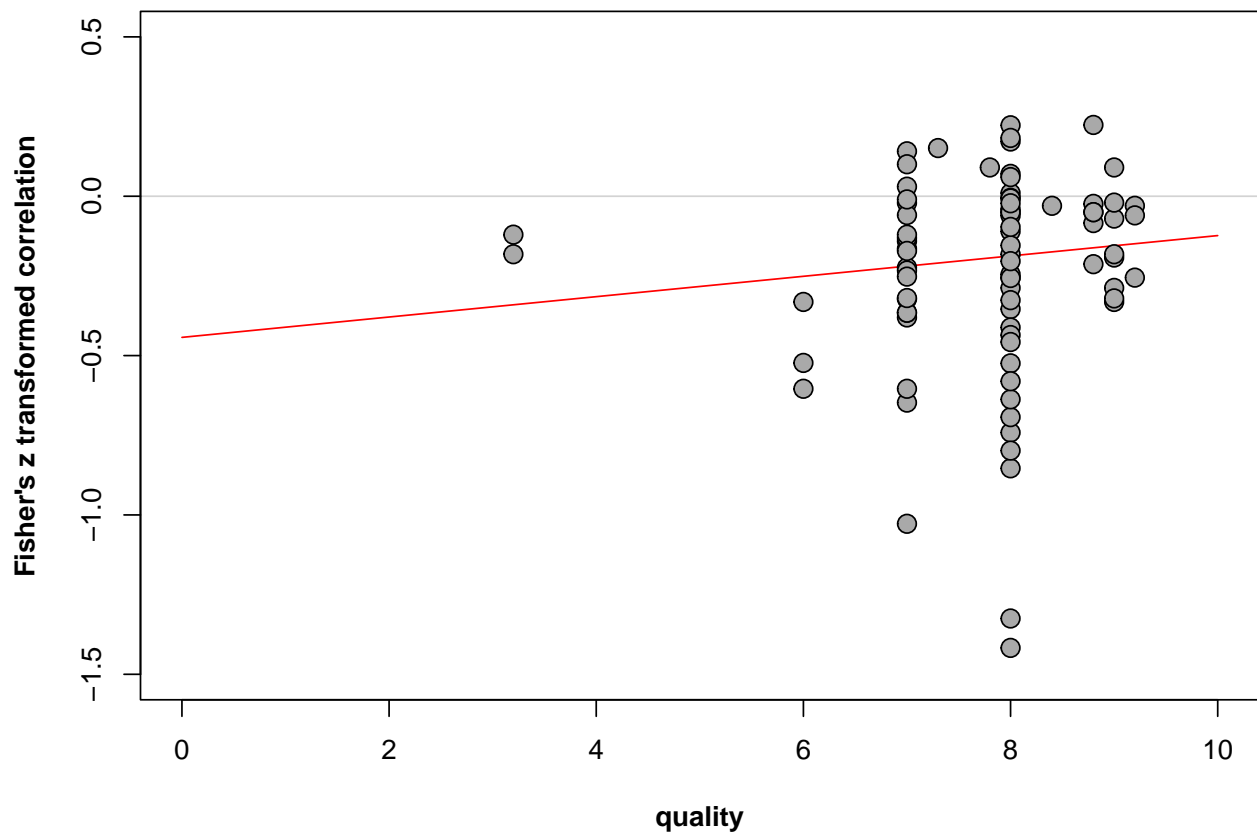
```
m.cor.reg.axis <- metareg(m, ~quality)
m.cor.reg.axis
```

```
##
## Multivariate Meta-Analysis Model (k = 83; method: REML)
##
## Variance Components:
##
##          estim      sqrt  nlvls  fixed    factor
```

```
## sigma^2.1  0.0582  0.2412    63    no      .id
## sigma^2.2  0.0057  0.0754    83    no  .id/.idx
##
## Test for Residual Heterogeneity:
## QE(df = 81) = 544.5239, p-val < .0001
##
## Test of Moderators (coefficient 2):
## F(df1 = 1, df2 = 81) = 0.8018, p-val = 0.3732
##
## Model Results:
##
##           estimate      se      tval  df    pval    ci.lb    ci.ub
## intrcpt   -0.4429  0.2783  -1.5914  81  0.1154  -0.9966  0.1108
## quality    0.0319  0.0357   0.8954  81  0.3732  -0.0390  0.1029
##
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

A study's quality score ($k = 83$) did not influence correlation values ($p = 0.3732$).

```
bubble(m.cor.reg.axis,
       xlim = c(0, 10),
       ylim = c(-1.5, 0.5),
       xlab = 'quality',
       studlab = FALSE,
       font.lab = 2,
       cex = 1.5,
       cex.studlab = 0.6,
       pos.studlab = 1,
       offset = 0.5,
       col.line = 'red')
```



Publication bias analysis

Small-study effects

Funnel plot

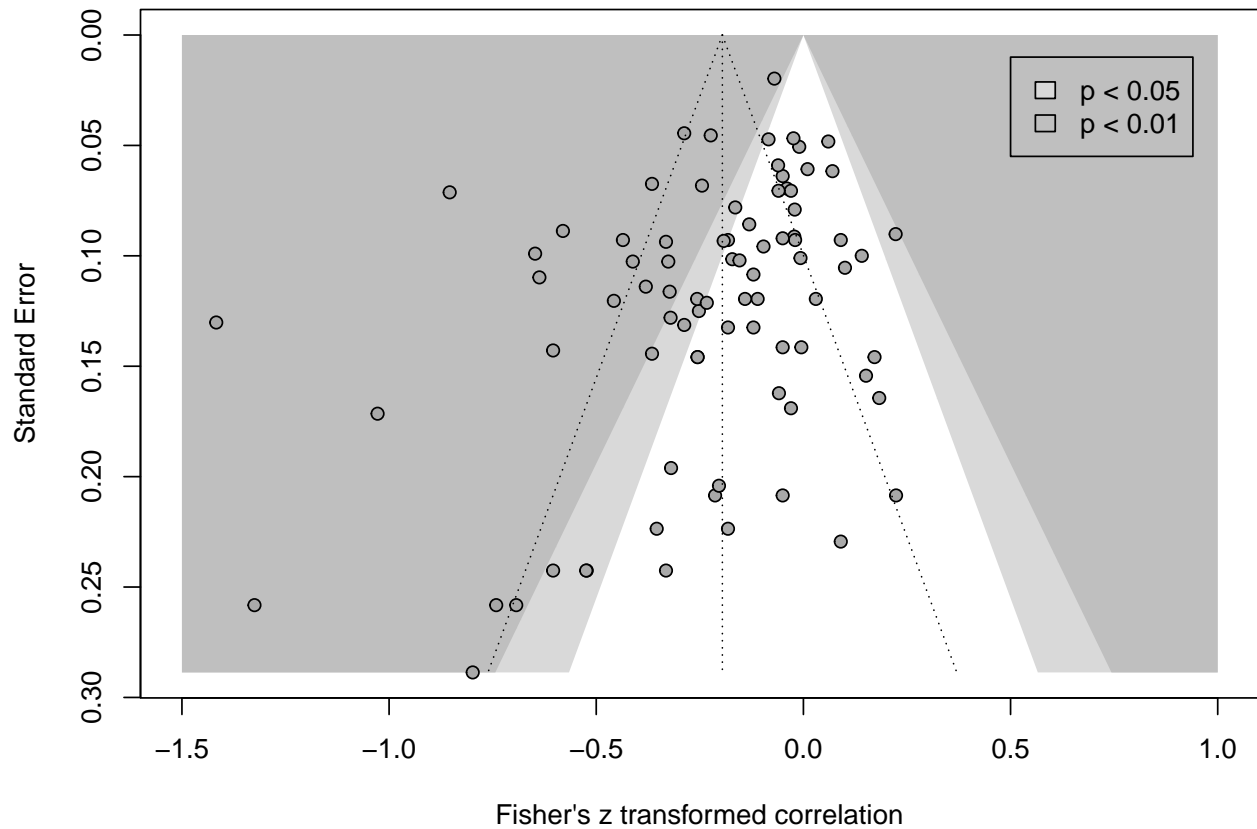
```
# Define fill colors for contour
col.contour = c("gray85", "gray75")

# Funnel plot
meta::funnel(m,
  xlim = c(-1.5, 1),
  contour = c(0.95, 0.99),
  col.contour = col.contour)

# legend
legend(x = 0.5, y = 0.01,
  legend = c("p < 0.05", "p < 0.01"),
  fill = col.contour)

# title
title("Contour-Enhanced Funnel Plot (Kinesiophobia and Physical Activity)")
```


Contour-Enhanced Funnel Plot (Kinesiophobia and Physical Activity)



```
png(file = "Funnel Plot.png", width = 2500, height = 2000, res = 300)

# Define fill colors for contour
col.contour = c("gray85", "gray75")

# Funnel plot
meta::funnel(m,
  xlim = c(-1.5, 1),
  contour = c(0.95, 0.99),
  col.contour = col.contour)

# legend
legend(x = 0.5, y = 0.01,
  legend = c("p < 0.05", "p < 0.01"),
  fill = col.contour)

# title
title("Contour-Enhanced Funnel Plot (Kinesiophobia and Physical Activity)")

dev.off()

## pdf
## 2
```

Egger's test

There is no direct function to conduct Egger's test for multi-level model. Alternatively, we calculate it by using the standard errors of the effect size estimates as a predictor in the meta-regression. (see: https://rstudio-pubs-static.s3.amazonaws.com/814435_6401e0cceb0b410c8208642dc5ee07f0.html)

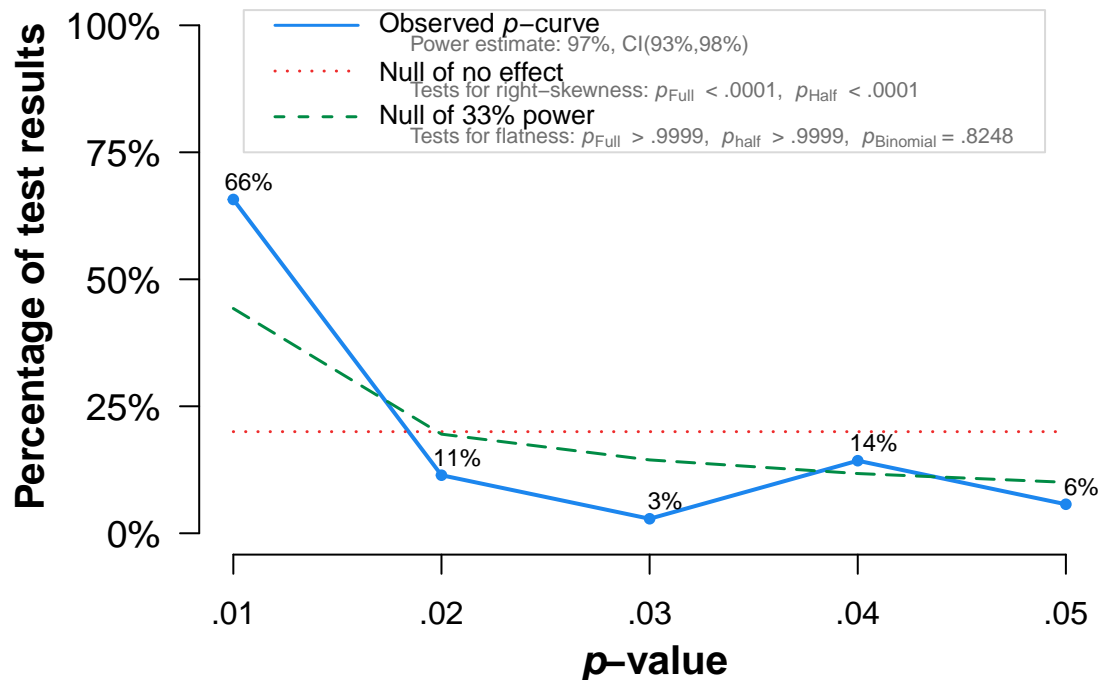
```
test.egger = rma.mv(z,var.z, mod = ~ sqrt(var.z), random = ~ 1 | author/cor_id, data = Kinphob_r, test.egger

##
## Multivariate Meta-Analysis Model (k = 83; method: REML)
##
## Variance Components:
##
##          estim      sqrt  nlvls  fixed      factor
## sigma^2.1  0.0534  0.2312    63    no      author
## sigma^2.2  0.0060  0.0772    83    no  author/cor_id
##
## Test for Residual Heterogeneity:
## QE(df = 81) = 506.5963, p-val < .0001
##
## Test of Moderators (coefficient 2):
## F(df1 = 1, df2 = 81) = 7.0537, p-val = 0.0095
##
## Model Results:
##
##          estimate      se      tval  df      pval      ci.lb      ci.ub
## intrcpt      -0.0333  0.0695  -0.4799  81  0.6326  -0.1716   0.1049
## sqrt(var.z)  -1.4968  0.5636  -2.6559  81  0.0095  -2.6181  -0.3754  **
##
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Egger's regression test using the standard errors of the effect size estimates as a predictor in the meta-regression showed that the coefficient of the standard error was significant ($b = -1.497$, 95% CI: -2.618 to -0.3754, $p = 0.0095$), suggesting that the data in the funnel plot was asymmetrical. This asymmetry may be explained by publication bias, but also by other potential causes, such as different study procedures and between-study heterogeneity, which was substantial-to-considerable here.

Pcurve analysis

```
pcurve(m, effect. estimation = FALSE, N, dmin = 0, dmax = 1)
```



Note: The observed p -curve includes 35 statistically significant ($p < .05$) results, of which 27 are $p < .025$. There were 48 additional results entered but excluded from p -curve because they were $p > .05$.

```
## P-curve analysis
## -----
## - Total number of provided studies: k = 83
## - Total number of p<0.05 studies included into the analysis: k = 35 (42.17%)
## - Total number of studies with p<0.025: k = 27 (32.53%)
##
## Results
## -----
##               pBinomial   zFull pFull   zHalf pHalf
## Right-skewness test      0.001 -13.811    0 -15.583    0
## Flatness test            0.825   9.421    1  14.677    1
## Note: p-values of 0 or 1 correspond to p<0.001 and p>0.999, respectively.
## Power Estimate: 97% (93.4%-98.5%)
##
## Evidential value
## -----
## - Evidential value present: yes
## - Evidential value absent/inadequate: no
```

The 83 Pearson's r correlation values were provided to the p -curve analysis. The observed p -curve included 35 statistically significant results ($p < 0.05$), 27 of which were highly significant ($p < 0.025$), and was visually right-skewed. The other results were excluded because they had a $p > 0.05$. The p -value of the right-skewness test was < 0.001 for both the half curve (curve of p values > 0.025) and the full curve (curve of p values < 0.05), confirming that the p -curve was right-skewed and suggesting that the effect of our meta-analysis is true, i.e., that the effect we estimated is not an artifact caused by selective reporting (e.g., p -hacking) in the literature 120. In addition, the statistical power of the studies that were included in the p -curve analysis was 97% (90% CI: 93 to 98%), suggesting that approximately 90% of the significant results are expected to be replicable.

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