Meta-analysis Criterion validity

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Introduction

This is part of a meta-analysis of a larger systematic review about the measurement properties of the Borg scales. In this analysis, we meta-analysed the correlation coefficients (treated as effect size) of the studies that about *Criterion Validity* of the *Borg 6-20 scale*.

Code & output

Install packages

library(knitr)

kable(borg, caption = "Borg dataset")

```
install.packages(c("readxl", "dplyr", "metafor"), dependencies = TRUE)
## Installing packages into '/cloud/lib/x86_64-pc-linux-gnu-library/4.5'
## (as 'lib' is unspecified)
library(readxl)
library(dplyr)
##
## 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
library(metafor)
## Loading required package: Matrix
## Loading required package: metadat
## Loading required package: numDeriv
## Loading the 'metafor' package (version 4.8-0). For an
## introduction to the package please type: help(metafor)
Import data
borg <- read_excel("Data_Borg.xlsx")</pre>
```

Table 1: Borg dataset

author_year	n	prom	$validity_type$	comparator	r
Liu,2000	36	Borg6-20	criterion	VO2peak	0.83
Kerrigan, 2016	24	Borg6-20	criterion	VO2R	0.91
Levinger,2004	14	Borg6-20	criterion	HRmax	0.43
Connolly,1996	9	Borg6-20	construct	VO2	0.8
Connolly,1996	9	Borg6-20	construct	$_{ m HR}$	0.49
Connolly,1996	9	Borg6-20	construct	VE	0.75
Tang,2016	97	Borg6-20	construct	$_{ m HR}$	0.44
Tang,2016	47	Borg6-20	construct	HR_beta_blockers	0.52
Shephard,1996	36	Borg6-20	responsiveness	VO2	NA
Buckley,2009	11	Borg6-20	reliability	ICC	0.85
Iellamo,2014	16	BorgCR10	criterion	TRIMPi	0.72

Cleaning (ensure numeric r and n)

Analysis. Filter: criterion validity + Borg 6-20

```
criterion <- borg %>%
  filter(validity_type == "criterion") %>%
  filter(prom != "BorgCR10")
```

Compute Fisher's Z effect sizes for r

```
criterion_es <- escalc(
  measure = "ZCOR",
  ri = r,
  ni = n,
  data = criterion
)</pre>
```

Fit random-effects meta-analysis

```
res <- rma(yi, vi, data = criterion_es, method = "REML")</pre>
```

```
print(summary(res))
## Random-Effects Model (k = 3; tau^2 estimator: REML)
## logLik deviance
                       AIC
                                 BIC
                                            AICc
## -1.5635
           3.1270
                       7.1270
                                 4.5133
                                         19.1270
##
## tau^2 (estimated amount of total heterogeneity): 0.2055 (SE = 0.2599)
## tau (square root of estimated tau^2 value): 0.4533
## I^2 (total heterogeneity / total variability): 80.27%
## H^2 (total variability / sampling variability): 5.07
## Test for Heterogeneity:
## Q(df = 2) = 8.2405, p-val = 0.0162
## Model Results:
## estimate
                             pval ci.lb ci.ub
             se zval
   1.0916 0.2940 3.7127 0.0002 0.5154 1.6679 ***
##
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
print(predict(res, transf = transf.ztor))
##
##
     pred ci.lb ci.ub pi.lb pi.ub
## 0.7975 0.4741 0.9313 0.0327 0.9733
Forest plot
slabs <- criterion$author_year</pre>
forest(
 res,
 slab = slabs,
 xlab = "Correlation (r)",
 transf = transf.ztor
```

