Supplemental Results

Risk of Bias Moderator Analyses

Allocation concealment: Q-M = .89, df = 2, p = .64

Incomplete outcome reporting: Q-M = 3.0, df =1, p = .08

Selective outcome reporting: Q-M = 1.72, df = 2, p = .42

Sequence generation: Q-M = 3.5, df = 1, p = .06

Blinding of outcome assessors: Q- M= 1.3, df = 2, p = .51

Blinding of participants and personnel: Q-M = 2.18, df = 2, p = .34

Other sources of bias: Q-M = 0.45, df = 1, p = .50

Sensitivity Analyses

Outlier Removed

Multilevel Estimates

Acquisition: g = .23, p = .10, 95% CI [-.04, .51]

Immediate retention: g = .12, p = .43, 95% CI [-.18, .41]

Delayed retention: g = .19, p = .10, 95% CI [-.04, .42]

Categorical Moderator Analyses

Age: Q-M = 5.42, df = 6, p = .49

Skill: Q-M = 1.1, df = 3, p = .78

Faded: Q-M = 2.17, df = 4, p =.70

Yoked: Q-M = 6.27, df = 4, p = .18

Feedback: Q-M = 7.64, df = 8, p = .47

Measure: Q-M = 30.65, df = 22, p = .10

Measure without test interaction: Q-M = 9.07, df = 8, pi = .34

Bandwidth: Q-M = 3.35, df = 5, p = .65

Mete-regression Analyses

Trials: Q-M = 5.80, df = 3, p = .12

Days: Q-M = 2.72, df = 3, p = .44

Frequency (overall analysis): Q-M = 1.99, df = 3, p = .57

Immediate retention interval: Q-M = .087, df = 2, p = .65

Test Time Moderators

Immediate retention vs. delayed retention: Q-M = .24, df = 1, p = .63.

Cluster Robust Inference Methods

Cluster Robust Multilevel Estimates

Acquisition: g = .19, p = .20, 95% CI [-.11, .50]

Immediate retention: g = .14, p = .93, 95% CI [-.29, .31]

Delayed retention q = .19, p = .15, 95% Ci [-.07, .46]

Correlated and Hierarchical Effects (CHE) Model Estimates with Approximate V Matrix (r = .7)

Acquisition: g = .19, p = .22, 95% CI [-.13, .51]

Immediate retention: q = .002, p = .99, 95% CI [-.34, .34]

Delayed retention: g = .20, p = .13, 95% CI [-.07, .48]

Cluster Robust Multilevel Estimates with Outlier Removed

Acquisition: g = .23, p = .11, 95% CI [-.06, .52]

Immediate retention: g = .12, p = .34, 95% CI [-.13, .37]

Delayed retention g = .19, p = .14, 95% Ci [-.07, .45]

CHE Model Estimates with Approximate V Matrix (r = .7) and Outlier Removed

Acquisition: g = .23, p = .13, 95% CI [-.08, .53]

Immediate retention: q = .12, p = .36, 95% CI [-.15, .38]

Delayed retention: q = .20, p = .13, 95% CI [-.06, .46]

Four Level Model: Measure Nested in Test Nested in Experiment

Multilevel Estimates

Acquisition: q = .15, p = .13, 95% CI [-.04, .34]

Immediate retention: g = .07, p = .56, 95% CI [-.15, .29]

Delayed retention: g = .18, p = .051, 95% CI [-.001, .36]

Four Level Model: Outlier Removed

Multilevel Estimates

Acquisition: g = .18, p = .07, 95% CI [-.01, .37]

Immediate retention: g = .12, p = .27, 95% CI [-.09, .34]

Delayed retention: g = .18, p = .051, 95% CI [-.001, .36]

Moderator Analyses

Age: Q-M = 7.81, df = 6, p = .25

Age (outlier removed): Q-M = 6.06, df = 6, p = .42

Skill: Q-M = 1.75, df = 3, p = .63

Skill (outlier removed): Q-M = .98, df = 3, p = .81

Task: Q-M = 19.47, df = 8, p = .01

Task (Drews et al. 2021 removed): Q-M = 9.4, df = 8, p = .31

Bandwidth: Q-M = 1.74, df = 5, p = .88

Bandwidth (outlier removed): Q-M = 1.19, df = 5, p = .94

Faded: Q-M = 3.03, df = 5, p = .70

Faded (outlier removed): Q-M = 2.36, df = 5, p = .80

Yoked: Q-M = 6.49, df = 4, p = .17

Yoked (outlier removed): Q-M = 6.22, df = 4, p = .18

Feedback: Q-M = 11.66, df = 16, p = .77

Feedback (outlier removed): Q-M = 8.53, df = 16, p = .93

Feedback (interaction removed): Q-M = 4.15, df = 6, p = .66

Feedback Collapsed: Spatial error, temporal error, variable error, movement time, form, and other.

Feedback: Q-M = 11.41, df = 14, p = .78

Feedback (outlier removed): Q-M = 8.71, df = 14, p = .92

Feedback (interaction removed): Q-M = 2.21, df = 5, p = .82

Feedback (delayed retention only): Q-M = 2.07, df = 5, p = .84

Measure: Q-M = 16.17, df = 28, p = .96

Measure (outlier removed): Q-M = 14.49, df = 28, p = .98.

Measure (interaction removed): Q-M = 5.24, df = 10, p = .87.

Do Measures Selected as Primary Differ from Secondary Measures

Full sample: Q-M = 1.85, df = 5, p = .87

Outlier removed: Q-M = 1.08, df = 5, p = .96

Meta-regression Analyses

Trials: Q-M = 9.83, df = 5, p = .08

Trials (outlier removed): Q-M = 9.92, df = 5, p = .08

Days: Q-M = 7.75, df = 5, p = .17

Days (outliers removed): Q-M = 7.70, df = 5, p = .17

Days: Q-M = 7.75, df = 5, p = .17

Univariate Analysis of Transfer Test Data

Estimate: *g* = .15, *p* = .45, 95% CI [-.24, .55]

Heterogeneity: Q = 68.47, df = 14, p < .0001. $\tau^2 = .57$

Estimate (outlier removed): g = .05, p = .83, 95% CI [-.38, .48]

Heterogeneity (outliers removed): Q = 55.19, df = 13, p < .0001. $\tau^2 = .42$

Moderator Analyses of Spatial Error Subset and Delayed Retention Time Point

Age: Q-M = 1.72, df = 1, p = .19

Skill: Q-M = .61, df = 1, p = .44

Task: Q-M = 1.77, df = 2, p = .41

Faded: Q-M = .72, df = 1, p = .40

Yoked: Q-M = .29, df = 1, p = .59

Feedback: Q-M = 5.08, df = 4, p = .28

```
Trials: Q-M < .001, df = 1, p = .98

Days: Q-M = .002, df = 1, p = .96

Frequency: Q-M = 1.14, df = 1, p = .29
```

Simulation Study

```
2. #Meta-analysis of dependent effect sizes simulation #
4. library(MBESS)
5. library(metafor)
library(faux)
7. library(tidyverse)
8. set.seed(1000)
9.
10.
11.
12. # individual meta-analysis simulation function
14. onemeta <- function(nSims, cxa,exa, cxb, exb, rho, sample_lb, sample_ub){</pre>
15.
     esd_1 <-numeric(nSims) #empty container for all simulated ES</pre>
     esd 2 <-numeric(nSims) #empty container for all simulated ES
     SSn1 <-numeric(nSims) #empty container for random sample sizes group 1
     SSn2 <-numeric(nSims) #empty container for random sample sizes group 2
18.
19
     for(i in 1:nSims){ #for each simulated experiment
20.
       SampleSize<-sample(sample lb:sample ub, 1) #randomly draw a sample between lb and ub
       ge <-rnorm multi(n = SampleSize, 2, 0, 1, r = rho, varnames = c("t1", "t2")) #sample
21.
   from a multivariate normal
       gc <-rnorm_multi(n = SampleSize, 2, 0, 1, r = rho, varnames = c("t1", "t2")) #sample</pre>
22.
   from a multivariate normal
       gc$t1 <- gc$t1 + cxa # add effect to time 1 for control</pre>
23.
24.
       ge$t1 <- ge$t1 + exa # add effect to time 1 for experimental</pre>
       gc$t2 <- gc$t2 + cxb # add effect to time 2 for experimental</pre>
25.
       ge$t2 <- ge$t2 + exb # add effect to time 2 for experimental</pre>
26.
27.
       SSn1[i]<-SampleSize #save sample size group 1</pre>
28.
       SSn2[i]<-SampleSize #save sample size group 2</pre>
29.
       esd 1[i] < -smd(Mean.1 = mean(ge$t1), Mean.2 = mean(gc$t1), s.1 = sd(ge$t1), s.2 = sd(gc$t1),
   n.1=SampleSize, n.2=SampleSize, Unbiased=TRUE) #Use MBESS to calc Hedges g
       esd_2[i] < -smd(Mean.1 = mean(ge$t2), Mean.2 = mean(gc$t2), s.1 = sd(ge$t2), s.2 = sd(gc$t2),
   n.1=SampleSize, n.2=SampleSize, Unbiased=TRUE)}
31.
    #Insert effect sizes and sample sizes
32. n1<-c(SSn1)
33. n2 < -c(SSn2)
     J<-1-3/(4*(SSn1+SSn2-2)-1) #correction for bias
     esdv_1 < -(((SSn1+SSn2)/(SSn1*SSn2))+(esd_1^2/(2*(SSn1+SSn2))))*J^2
     esdv_2 <-(((SSn1+SSn2)/(SSn1*SSn2)) + (esd_2^2/(2*(SSn1+SSn2))))*J^2
36.
37.
     dat <- as.data.frame(cbind(esd_1, esdv_1, esd_2, esdv_2, n1, n2))</pre>
38.
     dat$id <- seq.int(nrow(dat))</pre>
39.
     dat<-dat %>%
40.
       pivot_longer(
41.
         cols = !c(n1,n2,id),
         names_to = c(".value", "num"),
names_sep = "_")
42.
43.
44.
     return(as.data.frame(dat))
45.
46.
     }
```

```
47.
48.
49.
50. # simulation study ## multilevel models
52. iters <- 100 #number of simulated meta-analyses
53. mpval <- rep(NA, iters) #empty container for moderator p-values
54. difb <- rep(NA, iters) #empty container for moderator beta values
55. nSims <- 32 #number of experiments per meta-analysis
56. cxa <- 0 #intervention effect for control group at first time point
57. exa <- 0 #intervention effect for experimental group at first time point
58. cxb <- 0 #intervention effect for control group at second time point
59. exb <- 0 #intervention effect for experimental group at second time point
60. rho <- .5 #correlation between time points in the population
61. sample_lb <- 10 #lower bound for number of participants per experiment
62. sample ub <- 30 #upper bound for number of participants per experiment
63.
64.
65.
66.
67. for (i in 1:iters){
     dat <-onemeta(nSims = nSims, cxa = cxa, exa = exa, cxb = cxb, exb = exb, rho = rho,
   sample_lb = sample_lb, sample_ub = sample_ub)
69.
     tryCatch(
70.
         res <- rma.mv(esd, esdv, mods = ~factor(num), random = ~1|id/num, data = dat) #fit
71.
   multilevel model with time points nested in experiments
72.
         mpval[i] <- res$QMp</pre>
73.
          difb[i] <- res$b[2]},
74.
       error=function(error_message) {
75.
         message(error_message)
         return(NA) #when the model fails to converge, store as NA rather than stopping the
76.
   loop
77.
78.
     )
79. }
81. mean(mpval <= .05, na.rm = TRUE) # mean rejection rate with alpha = .05
82. mean(difb, na.rm = TRUE) # mean estimate of difference in intervention effect between time
   points
83.
84.
85.
86.
87.
89. # simulation study ## cluster robust methods
91. iters <- 100 #number of simulated meta-analyses
92. mpval <- rep(NA, iters) #empty container for moderator p-values
93. difb <- rep(NA, iters) #empty container for moderator beta values
94. nSims <- 32 #number of experiments per meta-analysis
95. cxa <- 0 #intervention effect for control group at first time point
96. exa <- 0 #intervention effect for experimental group at first time point
97. cxb <- 0 #intervention effect for control group at second time point
98. exb <- 0 #intervention effect for experimental group at second time point
99. rho <- .5 #correlation between time points in the population
100. sample_lb <- 10 #lower bound for number of participants per experiment
101. sample_ub <- 30 #upper bound for number of participants per experiment
102.
103.
104.
105. for (i in 1:iters){
       dat <-onemeta(nSims = nSims, cxa = cxa, exa = exa, cxb = cxb, exb = exb, rho = rho,
   sample_lb = sample_lb, sample_ub = sample_ub)
```

```
107.
        tryCatch(
108.
109.
        res <- rma.mv(esd, esdv, mods = ~factor(num), random = ~1 id/num, data = dat)
110.
        rob <-robust(res, cluster = dat$id) #use cluster robust inference methods</pre>
111.
       mpval[i] <- rob$QMp</pre>
112.
        difb[i] <- rob$b[2]},
113.
        error=function(error_message) {
114.
          message(error_message)
115.
         return(NA)
116.
117.
118. }
119.
120.
     mean(mpval <= .05, na.rm = TRUE) # mean rejection rate with alpha = .05</pre>
121. mean(difb, na.rm = TRUE) # mean estimate of difference in intervention effect between time
122.
123.
```

To conduct a simulation, adjust the following settings:

Iters: Number of meta-analyses to simulate (warning, the simulation can take a long time to run if this number is large)

nSims: Number of studies per meta-analysis.

cxa: The size of the effect to add to the first time point for the control group. Effects are scaled to Cohen's d.

exa: The size of the effect to add to the first time point for the experimental group.

cxb: The size of the effect to add to the second time point for the control group.

exb: The size of the effect to add to the second time point for the experimental group.

rho: The correlation between time points in the population, prior to adding intervention effects.

sample_lb: The lower bound for number of participants per group in an experiment included in a simulated meta-analysis. Groups will be equal in size for each experiment.

sample_ub: The upper bound for number of participants per group.

This simulation can be extended to return additional results from meta-analysis models by adding containers above the simulation and specifying the output to save within the simulation.