## Smith Replication: Primary Data Analysis

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#### Session Information

```
# References for R packages
# install.packages(devtools)
# require(devtools)
# devtools::install_qithub("crsh/papaja")
# require(papaja)
# papaja::r_refs(file = "results/r-package-refs.bib") #<-- writes bib file w/refs
#Save package versions
#renv::snapshot()
sessionInfo()
## R version 4.2.1 (2022-06-23 ucrt)
## Platform: x86_64-w64-mingw32/x64 (64-bit)
## Running under: Windows 10 x64 (build 19045)
## Matrix products: default
##
## locale:
## [1] LC COLLATE=English United States.utf8 LC CTYPE=English United States.utf8
## [3] LC_MONETARY=English_United States.utf8 LC_NUMERIC=C
## [5] LC_TIME=English_United States.utf8
## attached base packages:
## [1] stats
                graphics grDevices datasets utils
                                                         methods
                                                                    base
##
## other attached packages:
## [1] labeling_0.4.2
                               MBESS_4.9.2
                                                      highr_0.9
                                                                              farver_2.1.1
## [5] ggstance_0.3.5
                               scales_1.2.1
                                                      rlang_1.0.6
                                                                              cowplot_1.1.1
## [9] readxl_1.4.1
                               reshape2_1.4.4
                                                      superb_0.95.0
                                                                              psychReport_3.0.2
                                                      BayesFactor_0.9.12-4.4 coda_0.19-4
## [13] apaTables_2.0.8
                               psych_2.2.9
                               afex_1.2-0
## [17] ez_4.4-0
                                                      lme4_1.1-31
                                                                              Matrix 1.5-1
## [21] forcats_0.5.2
                               stringr_1.4.1
                                                      dplyr_1.0.10
                                                                              purrr_0.3.5
## [25] readr_2.1.3
                               tidyr_1.2.1
                                                      tibble_3.1.8
                                                                              ggplot2_3.4.0
                                                                              knitr_1.40
## [29] tidyverse_1.3.2
                               plyr_1.8.7
                                                      rmarkdown_2.17
## [33] pacman_0.5.1
##
```

```
## loaded via a namespace (and not attached):
##
  [1] googledrive_2.0.0
                            minqa_1.2.5
                                                 colorspace_2.0-3
                                                                      ellipsis_0.3.2
   [5] fs 1.5.2
                            rstudioapi 0.14
                                                                     fansi 1.0.3
                                                 MatrixModels 0.5-1
## [9] mvtnorm_1.1-3
                            lubridate_1.8.0
                                                 xml2_1.3.3
                                                                      splines_4.2.1
## [13] mnormt_2.1.1
                            jsonlite_1.8.3
                                                 nloptr_2.0.3
                                                                      lsr_0.5.2
## [17] broom 1.0.1
                                                 shiny_1.7.3
                                                                      compiler 4.2.1
                            dbplyr_2.2.1
## [21] httr 1.4.4
                                                                      fastmap 1.1.0
                            backports 1.4.1
                                                 assertthat_0.2.1
## [25] gargle_1.2.1
                            cli 3.4.1
                                                 later_1.3.0
                                                                     htmltools_0.5.3
## [29] tools 4.2.1
                            lmerTest_3.1-3
                                                 gtable_0.3.1
                                                                      glue_1.6.2
## [33] Rcpp_1.0.9
                            carData_3.0-5
                                                 cellranger_1.1.0
                                                                      vctrs_0.5.0
## [37] nlme_3.1-157
                            xfun_0.34
                                                 rbibutils_2.2.9
                                                                     rvest_1.0.3
## [41] mime_0.12
                            lifecycle_1.0.3
                                                 renv_0.16.0
                                                                      googlesheets4_1.0.1
                                                                      shinyBS_0.61.1
## [45] MASS_7.3-57
                            hms_1.1.2
                                                 promises_1.2.0.1
                                                                      stringi_1.7.8
## [49] parallel_4.2.1
                            yaml_2.3.6
                                                 pbapply_1.5-0
## [53] boot_1.3-28
                            Rdpack_2.4
                                                 pkgconfig_2.0.3
                                                                      evaluate_0.17
## [57] lattice_0.20-45
                            tidyselect_1.2.0
                                                 magrittr_2.0.3
                                                                      R6_2.5.1
## [61] generics_0.1.3
                            DBI_1.1.3
                                                 pillar_1.8.1
                                                                     haven_2.5.1
## [65] foreign 0.8-82
                            withr 2.5.0
                                                 mgcv 1.8-40
                                                                      abind 1.4-5
## [69] modelr 0.1.9
                                                 car_3.1-1
                            crayon_1.5.2
                                                                     utf8_1.2.2
## [73] tzdb 0.3.0
                            grid 4.2.1
                                                 reprex 2.0.2
                                                                     digest 0.6.30
## [77] xtable_1.8-4
                            httpuv_1.6.6
                                                 numDeriv_2016.8-1.1 munsell_0.5.0
```

#### Unconventional Analysis:

Time-Outs (>=1500 ms to respond) are treated as errors. The primary results are reported using this atypical approach to be to be consistent with the original paper by Smith et al. Near the end we also present a *Conventional Analysis* which drops the time-outs for Percent Error (PE) results.

## Experiment 1 - Stroop

#### Import and clean data

##		posture	SITTING			STANDING		
##		congruency	congruent	incongruent	neutral	congruent	incongruent	neutral
##	sj							
##	1		60	60	60	60	60	60
##	2		60	60	60	60	60	60
##	3		60	60	60	60	60	60
##	4		60	60	60	60	60	60
##	5		60	60	60	60	60	60
	6		60	60	60	60	60	60
	7		60	60	60	60	60	60
##	8		60	60	60	60	60	60
##	9		60	60	60	60	60	60
	10		60	60	60	60	60	60 60
##	11 12		60 60	60 60	60 60	60 60	60 60	60 60
##	13		60	60	60	60	60	60
##	14		60	60	60	60	60	60
##	15		60	60	60	60	60	60
##	16		60	60	60	60	60	60
##	17		60	60	60	60	60	60
##	18		60	60	60	60	60	60
##	19		60	60	60	60	60	60
##	20		60	60	60	60	60	60
##	21		60	60	60	60	60	60
##	22		120	120	120	120	120	120
##	24		60	60	60	60	60	60
##	25		60	60	60	60	60	60
##	26		60	60	60	60	60	60
##	27		60	60	60	60	60	60
##	28		60	60	60	60	60	60
##	29		60	60	60	60	60	60
##	30		60	60	60	60	60	60
##	31		60	60	60	60	60	60
##	32		60	60	60	60	60	60
	33		60	60	60	60	60	60
##	34		60	60	60	60	60	60
##	35		60	60	60	60	60	60 60
##	36		60	60	60	60	60	60 60
##	37		60	60	60	60	60	60 60
##	38 39		60 60	60 60	60 60	60 60	60 60	60 60
##			60	60	60	60		60 60
##	40		60	60	60	60	60	60

## 41	60	60	60	60	60	60
## 42	60	60	60	60	60	60
## 43	60	60	60	60	60	60
## 44	60	60	60	60	60	60
## 45	60	60	60	60	60	60
## 46	60	60	60	60	60	60
## 47	60	60	60	60	60	60
## 48	60	60	60	60	60	60
## 49	60	60	60	60	60	60
## 50	60	60	60	60	60	60

### ftable(blockType~sj, mergedStroopData)

##		blockType	experimental	practice
##	sj			
##	1		288	72
##	2		288	72
##	3		288	72
##	4		288	72
##	5		288	72
##	6		288	72
##	7		288	72
##	8		288	72
##	9		288	72
##	10		288	72
##	11		288	72
##	12		288	72
##	13		288	72
##	14		288	72
##	15		288	72
##	16		288	72
##	17		288	72
##	18		288	72
##	19		288	72
##	20		288	72
##	21		288	72
##	22		576	144
##	24		288	72
##	25		288	72
##	26		288	72
##	27		288	72
##	28		288	72
##	29		288	72
##	30		288	72
##	31		288	72
##	32		288	72
##	33		288	72
##	34		288	72
##	35		288	72
##	36		288	72
##	37		288	72
##	38		288	72
##	39		288	72
##	40		288	72

```
## 42
                         288
                                   72
## 43
                         288
                                   72
## 44
                         288
                                   72
                                   72
## 45
                         288
## 46
                         288
                                   72
## 47
                                   72
                         288
                                   72
## 48
                         288
                                   72
## 49
                         288
## 50
                         288
                                   72
\#...need to fix SJ - same one was used with two counterbalances
mergedStroopData$sj = paste(mergedStroopData$sj,"_",mergedStroopData$cb,sep="")
```

#...check for missing data
mergedStroopData[!complete.cases(mergedStroopData),]

## 41

##		sj	cb	blockNumber	blockType	trialNum	congruency	posture	wordStim	inkColour	rt	cResp
##	646	10_2	2	8	experimental	34	incongruent	STANDING	RED	green	0	2
##	1081	12_2	2	1	practice	1	neutral	SITTING	XXXXX	green	0	2
##	1117	12_2	2	2	experimental	1	incongruent	SITTING	GREEN	red	0	1
##	1445	13_1	1	1	practice	5	incongruent	STANDING	GREEN	red	0	1
##	1801	14_2	2	1	practice	1	neutral	SITTING	XXX	green	0	2
##	2162	15_1	1	1	practice	2	congruent	STANDING	GREEN	green	0	2
##	2163	15_1	1	1	practice	3	neutral	STANDING	XXXXX	green	0	2
##	2615	16_2	2	3	experimental	23	${\tt incongruent}$	SITTING	RED	green	0	2
##	2720	16_2	2	6	practice	20	neutral	STANDING	XXX	red	0	1
##	2737	16_2	2	7	experimental	1	${\tt incongruent}$	STANDING	GREEN	red	0	1
##	2885	17_1	1	1	practice	5	incongruent	STANDING	RED	green	0	2
##	3360	18_2	2	4	experimental	12	${\tt incongruent}$	SITTING	GREEN	red	0	1
##	3457	18_2	2	7	experimental	1	${\tt incongruent}$	STANDING	RED	green	0	2
##	3601	19_1	1	1	practice	1	congruent	STANDING	RED	red	0	1
##	4370	20_2	2	2	experimental	14	neutral	SITTING	XXXXX	red	0	1
##	4505	20_2	2	6	practice	5	incongruent	STANDING	GREEN	red	0	1
##	4681	21_1	1	1	practice	1	congruent	STANDING	RED	red	0	1
##	4682	21_1	1	1	practice	2	neutral	STANDING	XXX	red	0	1
##	4692	21_1	1	1	practice	12	incongruent	STANDING	RED	green	0	2
##	5041	22_1	1	1	practice	1	incongruent	STANDING	GREEN	red	0	1
##	5042	22_1	1	1	practice	2	incongruent	STANDING	RED	green	0	2
##	5043	22_1	1	1	practice	3	neutral	STANDING	XXX	red	0	1
##	5422	22_2	2	1	practice	22	incongruent	SITTING	GREEN	red	0	1
##	5428	22_2	2	1	practice	28	incongruent	SITTING	RED	green	0	2
##	5501	22_2	2	3	experimental	29	neutral	SITTING	XXXXX	red	0	1
##	5530	22_2	2	4	experimental	22	incongruent	SITTING	RED	green	0	2
##	5533	22_2	2	4	experimental	25	incongruent	SITTING	RED	green	0	2
##	5608	22_2	2	6	practice	28			GREEN	red	0	1
##	5621	22_2	2	7	experimental	5		STANDING	XXXXX	green	0	2
##	5644	22_2	2	7	experimental	28	neutral	STANDING	XXX	green	0	2
##	5668	22_2	2	8	experimental	16	incongruent		GREEN	red	0	1
##	5684	22_2	2	8	experimental	32		STANDING	XXXXX	red	0	1
##	5741	22_2	2	10	experimental	17	neutral	STANDING	XXX	green	0	2
##	5761	24_2	2	1	practice	1	congruent	SITTING	RED	red	0	1
	5763	24_2	2	1	practice	3	congruent	SITTING	GREEN	green	0	2
##	5768	24_2	2	1	practice	8	neutral	SITTING	XXX	green	0	2

	5770	24_2	2	1	practice		incongruent	SITTING	RED	green	0	2
##	5772	24_2	2	1	practice	12	congruent	SITTING	GREEN	green	0	2
##	5773	24_2	2	1	practice	13	neutral	SITTING	XXX	red	0	1
##	5775	24_2	2	1	practice		incongruent	SITTING	GREEN	red	0	1
##	5776	24_2	2	1	practice	16	neutral	SITTING	XXXXX	green	0	2
##	5797	24_2	2	2	experimental	1	neutral	SITTING	XXXXX	green	0	2
##	5798	24_2	2	2	${\tt experimental}$	2	incongruent	SITTING	RED	green	0	2
##	6018	24_2	2	8	experimental	6	congruent	STANDING	RED	red	0	1
##	6121	25_1	1	1	practice	1	neutral	STANDING	XXX	red	0	1
##	6482	26_2	2	1	practice	2	neutral	SITTING	XXXXX	red	0	1
##	6518	26_2	2	2	${\tt experimental}$	2	incongruent	SITTING	RED	green	0	2
##	6841	27_1	1	1	practice	1	congruent	STANDING	RED	red	0	1
##	6842	27_1	1	1	practice	2	neutral	STANDING	XXXXX	green	0	2
##	6843	27_1	1	1	practice	3	neutral	STANDING	XXXXX	red	0	1
##	7202	28_2	2	1	practice	2	incongruent	SITTING	GREEN	red	0	1
##	7921	3_1	1	1	practice	1	congruent	STANDING	GREEN	green	0	2
##	7957	3_1	1	2	experimental	1	neutral	STANDING	XXXXX	red	0	1
##	8858	31_1	1	7	experimental	2	congruent	SITTING	RED	red	0	1
##	9253	32_2	2	8	experimental	1	neutral	STANDING	XXX	green	0	2
##	9361	33_1	1	1	practice	1	congruent	STANDING	RED	red	0	1
##	9364	33_1	1	1	practice	4	incongruent	STANDING	GREEN	red	0	1
##	9386	33_1	1	1	practice	26	congruent	STANDING	GREEN	green	0	2
##	9390	33_1	1	1	practice	30	neutral	STANDING	XXX	red	0	1
##	9505	33_1	1	5	experimental	1	neutral	STANDING	XXXXX	red	0	1
##	10441	36_2	2	1	practice	1	congruent	SITTING	GREEN	green	0	2
##	10444	36_2	2	1	practice	4	incongruent	SITTING	GREEN	red	0	1
##	10447	36_2	2	1	practice	7	incongruent	SITTING	RED	green	0	2
##	10448	36_2	2	1	practice	8	neutral	SITTING	XXX	green	0	2
##	10535	36_2	2	3	experimental	23	incongruent	SITTING	GREEN	red	0	1
##	10639	36_2	2	6	practice	19	neutral	STANDING	XXXXX	green	0	2
##	10785	36_2	2	10	experimental	21	congruent	STANDING	GREEN	green	0	2
##	11294	38_2	2	4	experimental	26	incongruent	SITTING	RED	green	0	2
##	11387	38_2	2	7	experimental	11	neutral	STANDING	XXX	green	0	2
##	11418	38_2	2	8	experimental	6	incongruent	STANDING	RED	green	0	2
##	11421	38_2	2	8	experimental	9	incongruent	STANDING	GREEN	red	0	1
##	12202	4_2	2	9	experimental	34	incongruent	STANDING	GREEN	red	0	1
##	12241	40_2	2	1	practice	1	incongruent	SITTING	RED	green	0	2
##	13603	43_1	1	8	experimental	31	incongruent	SITTING	RED	green	0	2
##	13688	44_2	2	1	practice	8	incongruent	SITTING	RED	green	0	2
##	13689	44_2	2	1	practice	9	incongruent	SITTING	GREEN	red	0	1
##	13694	44_2	2	1	practice	14	incongruent	SITTING	GREEN	red	0	1
##	13695	44_2	2	1	practice	15	neutral	SITTING	XXXXX	red	0	1
##	13696	44_2	2	1	practice	16	incongruent	SITTING	RED	green	0	2
##	13702	44_2	2	1	practice	22	congruent	SITTING	GREEN	green	0	2
##	13709	44_2	2	1	practice	29	incongruent	SITTING	GREEN	red	0	1
##	13713	44_2	2	1	practice	33	incongruent	SITTING	RED	green	0	2
##	13715	44_2	2	1	practice	35	incongruent	SITTING	GREEN	red	0	1
##	13716	44_2	2	1	practice	36	congruent	SITTING	GREEN	green	0	2
##	13717	44_2	2	2	experimental	1	congruent	SITTING	RED	red	0	1
##	13721	44_2	2	2	experimental	5	incongruent	SITTING	GREEN	red	0	1
##	14041	45_1	1	1	practice	1	congruent	STANDING	GREEN	green	0	2
##	14379	45_1	1	10	experimental	15	neutral	SITTING	XXXXX	green	0	2
##	14864	47_1	1		experimental	32	congruent	STANDING	RED	red	0	1
##	14901	47_1	1		experimental	33	_	STANDING	XXX	green	0	2
										-		

	14958	_	1	6	practice		incongruent	SITTING	RED	green	0
	15121			1	practice	1	congruent	SITTING	GREEN	green	0
	15842	5_1	1	1	practice	2	${\tt incongruent}$		GREEN	red	0
##	15843	5_1	1	1	practice	3	congruent		RED	red	0
##	15845	5_1	1	1	practice	5	congruent		GREEN	green	0
##	15846	5_1	1	1	practice	6	incongruent	STANDING	RED	green	0
	15847	5_1	1	1	practice	7	neutral	STANDING	XXX	green	0
	16107	5_1	1	8	experimental	15	neutral	SITTING	XXX	red	0
##	16201	_	2	1	practice	1	incongruent	SITTING	GREEN	red	0
##	16565	6_2	2	1	practice	5	congruent	SITTING	RED	red	0
##	16957	7_1	1	2	experimental	1	neutral	STANDING	XXX	green	0
##	17282	8_2	2	1	practice	2	incongruent		GREEN	red	0
##	17643	9_1	1	1	practice	3	congruent	STANDING	GREEN	green	0
##	17644	9_1	1	1	practice	4	incongruent	STANDING	RED	green	0
##		resp									
	646		NA								
	1081		NA								
	1117		NA								
	1445		NA								
	1801		NA								
	2162	0	NA								
	2163	0	NA								
	2615		NA								
	2720		NA								
	2737		NA								
	2885		NA								
	3360		NA								
	3457		NA								
	3601		NA								
	4370		NA								
	4505		NA								
	4681		NA								
	4682		NA								
	4692		NA								
	5041		NA								
	5042		NA								
	5043		NA								
	5422		NA								
	5428		NA								
	5501		NA								
	5530		NA								
##	5533	0	NA								

2

1

1

2

2

2

1

1

1 2

1

2

2

## 5608

## 5621

## 5644

## 5668

## 5684

## 5741

## 5761

## 5763

## 5768

## 5770

## 5772

## 5773

O NA

```
## 5775
             O NA
## 5776
             O NA
## 5797
             O NA
## 5798
             O NA
## 6018
             O NA
## 6121
             O NA
## 6482
             O NA
             O NA
## 6518
## 6841
             O NA
## 6842
             O NA
## 6843
             O NA
## 7202
             O NA
## 7921
             O NA
## 7957
             O NA
## 8858
             O NA
## 9253
             O NA
## 9361
             O NA
## 9364
             O NA
## 9386
             O NA
## 9390
             O NA
## 9505
             O NA
## 10441
             O NA
## 10444
             O NA
## 10447
             O NA
## 10448
             O NA
## 10535
             O NA
## 10639
             O NA
## 10785
             O NA
## 11294
             O NA
## 11387
             O NA
## 11418
             O NA
## 11421
             O NA
## 12202
             O NA
## 12241
             O NA
## 13603
             O NA
## 13688
             O NA
## 13689
             O NA
## 13694
             O NA
## 13695
             O NA
## 13696
             O NA
## 13702
             O NA
## 13709
             O NA
## 13713
             O NA
## 13715
             O NA
## 13716
             O NA
## 13717
             O NA
## 13721
             O NA
## 14041
             O NA
## 14379
             O NA
## 14864
             O NA
## 14901
             O NA
## 14958
             O NA
## 15121
             O NA
## 15842
             O NA
```

```
## 15843
           O NA
## 15845 O NA
## 15846 O NA
## 15847 O NA
## 16107
         O NA
## 16201 O NA
## 16565 O NA
## 16957 O NA
## 17282
          O NA
## 17643 O NA
## 17644
           O NA
#...THERE IS MISSING DATA, BUT IT IS EXPECTED:
\#...There are trials where the the experiment times out
#...can be identified as RT == 0
#...trials where a response was not made have an RT =0, resp=0 and ac = NA
#...get the number of time outs
dim(mergedStroopData[!complete.cases(mergedStroopData),])[1]
## [1] 104
#...how are the missing trials distributed???
timeOutStroopData = mergedStroopData[!complete.cases(mergedStroopData),]
ftable(posture~congruency, timeOutStroopData)
              posture SITTING STANDING
## congruency
## congruent
                           10
                                     14
                            29
## incongruent
                                     18
## neutral
                            13
                                     20
ftable(blockType~sj, timeOutStroopData)
##
       blockType experimental practice
## sj
## 10_2
                                     0
                            1
## 12_2
                            1
                                      1
## 13_1
                            0
                                     1
## 14_2
                            0
                                     1
## 15_1
                            0
                                      2
## 16_2
                            2
                                     1
                            0
## 17_1
                                     1
                            2
## 18_2
                                     0
## 19_1
                            0
                                     1
## 20 2
                            1
                                     1
## 21_1
                            0
                                     3
## 22_1
                            0
                                     3
```

8

1

8

3

0

## 22\_2

## 24\_2

## 25\_1

```
## 26 2
                             1
                                      1
## 27_1
                             0
                                      3
## 28 2
                             0
                                      1
## 3_1
                             1
                                      1
## 31 1
                             1
                                      0
## 32 2
                             1
                                      0
## 33 1
                             1
## 36 2
                             2
                                      5
## 38 2
                             4
                                      0
## 4_2
                             1
                                      0
## 40_2
                             0
                                      1
## 43_1
                             1
                                      0
## 44_2
                             2
                                     10
## 45_1
                             1
                                      1
## 47_1
                             2
                                      1
## 48_2
                             0
                                      1
## 5_1
                             1
                                      5
## 50 2
                             0
## 6_2
                             0
                                      1
## 7 1
                             1
                                      0
## 8_2
                             0
                                      1
## 9_1
                             0
                                      2
#...this code changes the "time-out" trials as errors
#...see Davoli et al.
mergedStroopData$ac[mergedStroopData$rt==0] = 0
#...remove practice trials
mergedStroopData <- mergedStroopData[!mergedStroopData$blockType=="practice",]</pre>
#...check that only experimental trials are left
unique(mergedStroopData$blockType)
## [1] "experimental"
totalStroopTrials = dim(mergedStroopData)[1]
observationDataStroop = data.frame(ftable(blockType~sj, mergedStroopData))[,c(1,3)]
#...remove trials faster than 100ms
# mergedStroopData= mergedStroopData[!mergedStroopData$rt==0,] #...greater that 1500ms
mergedStroopData= mergedStroopData[!(mergedStroopData$rt<=100 & mergedStroopData$rt > 0),]
validStroopRTTrials = dim(mergedStroopData)[1]
observationDataStroop$validTrials = data.frame(ftable(blockType~sj, mergedStroopData))[,c(3)]
print(paste("percent invalid trials = ",
            ((totalStroopTrials-validStroopRTTrials)/totalStroopTrials)*100))
## [1] "percent invalid trials = 0.00694444444444444"
#Drops "time-out" trials
mergedStroopData.conventional <- mergedStroopData %>%
                                  filter(rt != 0 & resp != 0)
```

```
write.csv(mergedStroopData, file = "results/exp1_merged_stroop_data.csv",
           row.names = F)
#Conventional analysis drops the time out trials, rather than treating as errors
write.table(mergedStroopData.conventional, file = "results/exp1_merged_stroop_data_conv.txt",
           row.names = F)
stroopCorrect = mergedStroopData[mergedStroopData$ac == 1, ]
# mergedDataSet = mergedDataSet[mergedDataSet$ac ==1,]
errorsRemoved = dim(stroopCorrect)[1] #...total remaining trials
observationDataStroop$correctTrials = data.frame(ftable(blockType~sj, stroopCorrect))[,c(3)]
trimInfo = data.frame(totalStroopTrials, validStroopRTTrials,errorsRemoved)
head(trimInfo)
    total Stroop Trials\ valid Stroop RTT rials\ errors Removed
## 1
                14400
                                    14399
                                                  13852
#...percent of error trials lost
print(paste("percent errors removed = ",
           (((validStroopRTTrials-errorsRemoved)/totalStroopTrials)*100)))
## [1] "percent errors removed = 3.79861111111111"
######################################
#...CHECK 20% CRITERION
observationDataStroop$percentLoss =
  ((observation Data Stroop\$Freq-observation Data Stroop\$correct Trials)/\\
    observationDataStroop$Freq)*100
observationDataStroop$percentLoss
## [1] 3.4722222 1.7361111 0.3472222 6.9444444 1.7361111 4.1666667 4.5138889 1.7361111
## [9] 1.0416667 1.7361111 1.7361111 3.1250000 2.0833333 4.5138889 1.7361111 10.0694444
## [17] 8.6805556 0.3472222 2.0833333 4.5138889 4.8611111 1.3888889 1.3888889 3.4722222
## [25] 16.6666667 1.7361111 15.9722222 1.0416667 0.0000000 5.5555556 2.0833333 8.6805556
## [33] 0.3472222 3.1250000 1.7361111 2.0833333 4.1666667 2.7777778 2.4305556 7.6388889
## [41] 1.0416667 4.1666667 2.7777778 1.0416667 5.5555556 3.1250000 10.0694444 1.7361111
## [49] 3.4722222 3.8194444
sum(observationDataStroop$percentLoss>20)
## [1] 0
#...None!
#...RUN TRIMMING PROCEDURE
tempList = pjRecursiveTrim2(stroopCorrect, #...dataset
                           "rt", #...dependent variables
```

```
c("sj",
                             "cb",
                             "congruency",
                             "posture")) #.independent variables
trimmedStroopData=tempList[[1]]
totalStroopN = tempList[[2]]
rejectedStroop = tempList[[3]]
percentTrimmedStroop = tempList[[4]]
NcellsStroop = tempList[[5]]
#...get the trimming info
trimOutputStroop= data.frame(totalStroopN, rejectedStroop,percentTrimmedStroop,NcellsStroop)
head(trimOutputStroop)
## totalStroopN rejectedStroop percentTrimmedStroop NcellsStroop
## 1
          13852
                            292
                                           2.107999
                                                             300
stroopRT = plyr::ddply(trimmedStroopData,
                .(sj, cb,congruency,posture),
                summarise,
                meanRT = mean(rt))
head(stroopRT)
     sj cb congruency posture meanRT
## 1 1_1 1 congruent SITTING 471.6458
## 2 1_1 1 congruent STANDING 400.0638
## 3 1_1 1 incongruent SITTING 430.0455
## 4 1_1 1 incongruent STANDING 439.8444
## 5 1_1 1 neutral SITTING 454.5455
## 6 1_1 1
              neutral STANDING 408.4565
#...get error data
stroopPE = plyr::ddply(mergedStroopData,
                    .(sj, cb,congruency,posture),
                    summarise,
                    meanPE = 100 - (mean(ac)*100))
head(stroopPE)
     sj cb congruency posture
## 1 1_1 1 congruent SITTING 0.000000
## 2 1_1 1 congruent STANDING 0.000000
## 3 1_1 1 incongruent SITTING 4.166667
## 4 1_1 1 incongruent STANDING 6.250000
## 5 1_1 1 neutral SITTING 6.250000
## 6 1_1 1 neutral STANDING 4.166667
```

```
#...combine the RT and error data
stroopCombined = cbind(stroopRT,meanPE =stroopPE$meanPE)
head(stroopCombined)
     sj cb congruency posture
                                  meanRT
## 1 1_1 1 congruent SITTING 471.6458 0.000000
## 2 1_1 1 congruent STANDING 400.0638 0.000000
## 3 1_1 1 incongruent SITTING 430.0455 4.166667
## 4 1 1 1 incongruent STANDING 439.8444 6.250000
## 5 1_1 1
              neutral SITTING 454.5455 6.250000
## 6 1 1 1
               neutral STANDING 408.4565 4.166667
#...set as factors
stroopCombined$sj = factor(stroopCombined$sj)
stroopCombined$cb = factor(stroopCombined$cb)
Reaction time results
rtModelStroop <- ezANOVA(stroopCombined,</pre>
                  dv = .(meanRT),
                  wid=.(sj),
                  within=.(posture,congruency),
                  detailed=TRUE,
                  type=3,
                  return_aov=TRUE)
## Warning: Converting "posture" to factor for ANOVA.
## Warning: Converting "congruency" to factor for ANOVA.
rtModelStroop$ANOVA
##
                Effect DFn DFd
                                        SSn
                                                   SSd
                                                                               p p<.05
## 1
           (Intercept) 1 49 6.530862e+07 1322013.63 2.420643e+03 2.233955e-43
                                                                                     * 0.9748634585
## 2
               posture 1 49 8.221421e+02 156217.37 2.578776e-01 6.138604e-01
                                                                                       0.0004879807
                         2 98 7.093105e+04 154676.49 2.247026e+01 9.278220e-09
                                                                                     * 0.0404190166
            congruency
                         2 98 8.430066e+01 51054.35 8.090852e-02 9.223396e-01
## 4 posture:congruency
                                                                                       0.0000500584
rtStroopMSE = rtModelStroop$ANOVA$SSd/rtModelStroop$ANOVA$DFd
#...print ANOVA in nice format
paste(rtModelStroop$ANOVA$Effect,": F(",
     rtModelStroop$ANOVA$DFn,
     rtModelStroop$ANOVA$DFd,
     ") = ",
     round(rtModelStroop$ANOVA$F,3),
     ", MSE = ",
```

```
round(rtStroopMSE,3),
      ", p = ",
     round(rtModelStroop$ANOVA$p,3),
      ", partialEtaSq = ",
      round(rtModelStroop$ANOVA$SSn/(rtModelStroop$ANOVA$SSn+rtModelStroop$ANOVA$SSd),4),
## [1] "(Intercept): F(1, 49) = 2420.643, MSE = 26979.87, p = 0, partialEtaSq = 0.9802"
## [2] "posture: F(1, 49) = 0.258, MSE = 3188.11, p = 0.614, partialEtaSq = 0.0052"
## [3] "congruency: F(2, 98) = 22.47, MSE = 1578.332, p = 0, partialEtaSq = 0.3144"
## [4] "posture:congruency: F(2, 98) = 0.081, MSE = 520.963, p = 0.922, partialEtaSq = 0.0016"
#...CALCULATE THE BAYES FACTORS FOR THE RT ANALYSIS
stroopBF = stroopCombined
stroopBF$posture = factor(stroopBF$posture)
stroopBF$congruency = factor(stroopBF$congruency)
bfValues1 = anovaBF(meanRT~congruency*posture+sj,
                  data = stroopBF,
                   whichRandom = "sj",
                  method="laplace")
bfValues1
## Bayes factor analysis
## [1] congruency + sj
                                                     : 35335703 ±NA%
                                                     : 0.1461731 ±NA%
## [2] posture + sj
## [3] congruency + posture + sj
                                                      : 5410998
                                                                  ±NA%
## [4] congruency + posture + congruency:posture + sj : 346149.8 ±NA%
## Against denominator:
   meanRT ~ sj
## ---
## Bayes factor type: BFlinearModel, JZS
#...get the Bayes factor for the Null Interaction
bfValues1[3]/bfValues1[4]
## Bayes factor analysis
## [1] congruency + posture + sj : 15.63195 \pm NA\%
##
## Against denominator:
## meanRT ~ congruency + posture + congruency:posture + sj
## Bayes factor type: BFlinearModel, JZS
1/(bfValues1[3]/bfValues1[4])
## Bayes factor analysis
## -----
## [1] congruency + posture + congruency:posture + sj : 0.06397154 ±NA%
```

```
##
## Against denominator:
## meanRT ~ congruency + posture + sj
## ---
## Bayes factor type: BFlinearModel, JZS
# Difference scores and paired t-tests
#... stroop effect (incongruent - congruent) FOR Standing
standingStroop = stroopCombined[stroopCombined$posture=="STANDING", ]
standingStroop = standingStroop[standingStroop$congruency!="neutral", ]
t.test(standingStroop$meanRT[standingStroop$congruency=="congruent"],
       standingStroop$meanRT[standingStroop$congruency=="incongruent"],
      paired=TRUE )
## Paired t-test
##
## data: standingStroop$meanRT[standingStroop$congruency == "congruent"] and standingStroop$meanRT[sta
## t = -4.3805, df = 49, p-value = 6.226e-05
## alternative hypothesis: true mean difference is not equal to 0
## 95 percent confidence interval:
## -53.81756 -19.96796
## sample estimates:
## mean difference
         -36.89276
##
#... stroop effect (incongruent - congruent) FOR SITTING
sittingStroop = stroopCombined[stroopCombined$posture=="SITTING", ]
sittingStroop = sittingStroop[sittingStroop$congruency!="neutral", ]
t.test(sittingStroop$meanRT[sittingStroop$congruency=="congruent"],
       sittingStroop$meanRT[sittingStroop$congruency=="incongruent"],
       paired=TRUE )
##
##
  Paired t-test
## data: sittingStroop$meanRT[sittingStroop$congruency == "congruent"] and sittingStroop$meanRT[sittin
## t = -5.1209, df = 49, p-value = 5.104e-06
## alternative hypothesis: true mean difference is not equal to 0
## 95 percent confidence interval:
## -52.27703 -22.81052
## sample estimates:
## mean difference
##
        -37.54377
Percent error results
errModelStroop <- ezANOVA(stroopCombined,</pre>
                   dv = .(meanPE),
```

wid=.(sj),

```
within=.(posture,congruency),
                   detailed=TRUE,
                   type=3,
                   return_aov = TRUE)
## Warning: Converting "posture" to factor for ANOVA.
## Warning: Converting "congruency" to factor for ANOVA.
errStroopMSE = errModelStroop$ANOVA$SSd/errModelStroop$ANOVA$DFd
exp1 ANOVA acc uncon <-
paste(errModelStroop$ANOVA$Effect,": F(",
      errModelStroop$ANOVA$DFn,
      ", ",
      errModelStroop$ANOVA$DFd,
      ") = ",
      round(errModelStroop$ANOVA$F,3),
      ", MSE = ",
     round(errStroopMSE,3),
      ", p = ",
     round(errModelStroop$ANOVA$p,3),
      ", partialEtaSq = ",
     round(errModelStroop$ANOVA$SSn/(errModelStroop$ANOVA$SSn+errModelStroop$ANOVA$SSd),4),
      sep="")
exp1_ANOVA_acc_uncon
## [1] "(Intercept): F(1, 49) = 57.526, MSE = 75.297, p = 0, partialEtaSq = 0.54"
## [2] "posture: F(1, 49) = 0.007, MSE = 16.562, p = 0.934, partialEtaSq = 1e-04"
## [3] "congruency: F(2, 98) = 11.598, MSE = 9.222, p = 0, partialEtaSq = 0.1914"
## [4] "posture:congruency: F(2, 98) = 1.59, MSE = 6.228, p = 0.209, partialEtaSq = 0.0314"
write.csv(exp1_ANOVA_acc_uncon, "results/exp1_ANOVA_PE_uncon.csv")
#BF for errors
bfValues1.error = anovaBF(meanPE~congruency*posture+sj,
                          data = stroopBF,
                          whichRandom = "sj",
                          method="laplace")
bfValues1.error
## Bayes factor analysis
## -----
## [1] congruency + sj
                                                      : 776.6647 ±NA%
## [2] posture + sj
                                                      : 0.1172445 ±NA%
## [3] congruency + posture + sj
                                                      : 91.51836 ±NA%
## [4] congruency + posture + congruency:posture + sj : 13.90884 ±NA%
## Against denominator:
## meanPE ~ sj
## Bayes factor type: BFlinearModel, JZS
```

```
#...get the Bayes factor for the Null Interaction
bfValues1.error[3]/bfValues1.error[4]
## Bayes factor analysis
## -----
## [1] congruency + posture + sj : 6.579869 ±NA%
## Against denominator:
    meanPE ~ congruency + posture + congruency:posture + sj
##
## ---
## Bayes factor type: BFlinearModel, JZS
1/(bfValues1.error[3]/bfValues1.error[4])
## Bayes factor analysis
## -----
## [1] congruency + posture + congruency:posture + sj : 0.1519787 ±NA%
##
## Against denominator:
## meanPE ~ congruency + posture + sj
## Bayes factor type: BFlinearModel, JZS
#...ERRORs
#... stroop effect (incongruent - congruent) FOR Standing
t.test(standingStroop$meanPE[standingStroop$congruency=="congruent"],
       standingStroop$meanPE[standingStroop$congruency=="incongruent"],
      paired=TRUE )
##
## Paired t-test
##
## data: standingStroop$meanPE[standingStroop$congruency == "congruent"] and standingStroop$meanPE[sta
## t = -2.0681, df = 49, p-value = 0.04393
## alternative hypothesis: true mean difference is not equal to 0
## 95 percent confidence interval:
## -2.79325655 -0.04007678
## sample estimates:
## mean difference
##
        -1.416667
#... stroop effect (incongruent - congruent) FOR SITTING
t.test(sittingStroop$meanPE[sittingStroop$congruency=="congruent"],
       sittingStroop$meanPE[sittingStroop$congruency=="incongruent"],
       paired=TRUE )
##
## Paired t-test
##
## data: sittingStroop$meanPE[sittingStroop$congruency == "congruent"] and sittingStroop$meanPE[sittin
## t = -4.6535, df = 49, p-value = 2.51e-05
```

```
## alternative hypothesis: true mean difference is not equal to 0
## 95 percent confidence interval:
## -3.758593 -1.491407
## sample estimates:
## mean difference
## -2.625
```

### Make plots for Stroop

inxn.rt.MSE = rtStroopMSE[4]

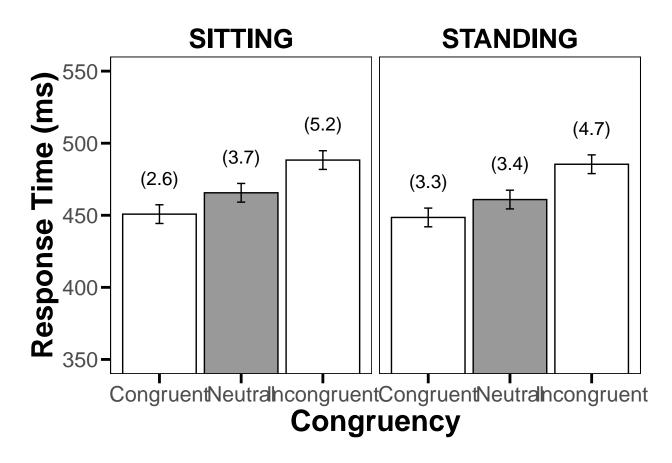
```
#...pull out summary statistics per condition averaged across subjects for graph
graphRT = describeBy(stroopCombined$meanRT,
                     list(stroopCombined$posture,stroopCombined$congruency),
                    mat=TRUE,
                    digits = 1)
graphPE = describeBy(stroopCombined$meanPE,
                     list(stroopCombined$posture,stroopCombined$congruency),
                     mat=TRUE,
                     digits = 1)
head(graphRT)
##
       item
             group1
                          group2 vars n mean
                                                 sd median trimmed mad
                                                                          min
                                                                                max range skew
## X11
         1 SITTING
                       congruent
                                   1 50 450.8 56.3 443.0 446.5 45.5 348.3 598.3 250.0 0.7
## X12
          2 STANDING
                                   1 50 448.5 60.6 440.3 442.2 48.6 341.3 624.4 283.0 1.0
                       congruent
## X13
         3 SITTING incongruent
                                   1 50 488.3 91.9 471.8
                                                            476.7 74.4 351.1 803.1 452.0
## X14
         4 STANDING incongruent
                                   1 50 485.4 100.6 458.7
                                                             470.5 79.0 358.3 861.5 503.2
                                                             460.1 49.7 357.8 702.6 344.7 1.0
## X15
          5 SITTING
                        neutral
                                   1 50 465.6 66.3 456.6
                        neutral
                                   1 50 460.9 67.6 452.7
                                                             453.6 51.5 346.2 695.4 349.2 1.4
## X16
         6 STANDING
      kurtosis
                 se
           0.2 8.0
## X11
## X12
           0.9 8.6
## X13
           1.7 13.0
## X14
           3.8 14.2
## X15
           1.9 9.4
## X16
           2.7 9.6
#...get rid of irrelevant columns
graphRT = graphRT[,c("group1", "group2", "mean", "se")]
graphPE = graphPE[,c("group1", "group2", "mean", "se")]
#...rename the variables
names(graphRT) = c("posture", "congruency", "mean", "se")
names(graphPE) = c("posture", "congruency", "mean", "se")
#...make sure posture is in UPPERCASE
graphRT$posture = str_to_upper(graphRT$posture)
#..calculate the within subjects confidence intervals based on loftus and masson
#...the confidence intervals are based on the interaction term.
```

```
inxn.err.MSE = errStroopMSE[4]
graphRT$se = sqrt((inxn.rt.MSE)/length(unique(stroopCombined$sj)))
graphPE$se= sqrt((inxn.err.MSE)/length(unique(stroopCombined$sj)))
critT = qt(p=.025, df=length(unique(stroopCombined$sj))-2,lower.tail =FALSE)
#---add the min and max for the confidence intervals
graphRT$min = graphRT$mean - (graphRT$se*critT)
graphRT$max = graphRT$mean + (graphRT$se*critT)
####GET AC DATA FROM twoAnimalWordsPRPac.R
graphRT$ac = paste("(",format(round(graphPE$mean,digits=1),nsmall = 1),")",sep="")
head(graphRT)
        posture congruency mean
                                        se
## X11 SITTING
                  congruent 450.8 3.227887 444.3099 457.2901 (2.6)
## X12 STANDING
                  congruent 448.5 3.227887 442.0099 454.9901 (3.3)
## X13 SITTING incongruent 488.3 3.227887 481.8099 494.7901 (5.2)
## X14 STANDING incongruent 485.4 3.227887 478.9099 491.8901 (4.7)
## X15 SITTING
                    neutral 465.6 3.227887 459.1099 472.0901 (3.7)
## X16 STANDING
                   neutral 460.9 3.227887 454.4099 467.3901 (3.4)
#...used for positioning the accuracy data on the graph
graphRT$vAdj = 25 #down
graphRT$vAdj[graphRT$congruency=="incongruent"]=25 #up
graphRT$hAdj = 0 #right
#qraphRT$hAdj[qraphRT$posture=="SITTING"]=-60 #left
graphRT$congruency = factor(graphRT$congruency, labels = c("Congruent", "Incongruent", "Neutral"))
graphRT$congruency = factor(graphRT$congruency, levels=c("Congruent", "Neutral", "Incongruent"))
interactionPlot <- ggplot(graphRT, aes(congruency, mean, group=posture)) +</pre>
  theme(legend.position = "none")+
  scale fill manual(values=c("#FFFFFF","#999999","#FFFFFF","#999999")) +
  coord_cartesian(ylim=c(350,550),expand=TRUE) +
  scale_y_continuous(breaks = round(seq(350, 550, by = 50),0)) +
  geom_text(aes(label=ac),nudge_x=graphRT$hAdj,nudge_y =graphRT$vAdj, size=5) +
  geom bar(stat="identity", aes(fill=interaction(congruency)),colour="black")+
  geom_errorbar(aes(ymin=min,ymax=max,group=interaction(posture,congruency)), width=.1)+
  labs(x = "Congruency", y = "Response Time (ms)") +
  theme(axis.ticks = element_line(size = 1, colour = "black", linetype = "solid"),
       axis.ticks.length = unit(.25, "cm"),
       #axis.line = element_line(size = 1, colour = "black", linetype = "solid"),
       panel.background = element_rect(fill = "white", colour = "white", size = 1),
       axis.text=element_text(size=16),
       axis.title=element_text(size=22,face="bold"),
        strip.text = element_text(size = 20, face = "bold",colour = "black", angle = 0),
        panel.border = element_rect(colour = "black", fill = NA, size = 0.50),
        strip.background = element_rect(fill=NA, colour="NA", size = 2))+
  facet_grid(~posture)
```

## Warning: The 'size' argument of 'element\_line()' is deprecated as of ggplot2 3.4.0.
## i Please use the 'linewidth' argument instead.

## Warning: The 'size' argument of 'element\_rect()' is deprecated as of ggplot2 3.4.0.
## i Please use the 'linewidth' argument instead.

 ${\tt interactionPlot}$ 



##

```
##
## Means and standard deviations for meanRT as a function of a 3(congruency) X 2(posture) design
##
##
                                    M_95%_CI
                          М
                                                  SD
##
     posture:SITTING
##
          congruency
##
           congruent 450.77 [434.78, 466.76]
                                               56.26
##
         incongruent 488.31 [462.20, 514.42]
                                               91.87
##
             neutral 465.62 [446.78, 484.47]
                                               66.31
##
##
   posture:STANDING
##
          congruency
##
           congruent 448.51 [431.30, 465.72]
##
         incongruent 485.40 [456.81, 513.99] 100.61
##
             neutral 460.86 [441.64, 480.08] 67.64
##
## Note. M and SD represent mean and standard deviation, respectively.
## LL and UL indicate the lower and upper limits of the
## 95% confidence interval for the mean, respectively.
## The confidence interval is a plausible range of population means
## that could have created a sample mean (Cumming, 2014).
```

### Experiment 2 - Task-switching

### Import and clean data

```
###read in data
ts_path <- "/data/Experiment 2 Data/task-switching-replication-recoded-2.csv"
task_switching_raw <- read.csv(paste0(workingdir, ts_path))</pre>
head(task switching raw)
     participant session condition trialType posture blockNum trialNum switchTrialType
## 1
               1
                        1
                                  1 experiment standing
                                                                1
                                                                          1
                                                                                     buffer
## 2
               1
                                                                          2
                                                                                   noswitch
                        1
                                  1 experiment standing
                                                                1
## 3
                                  1 experiment standing
                                                                          3
                        1
                                                                1
                                                                                     switch
## 4
               1
                                  1 experiment standing
                                                                          4
                                                                                   noswitch
                        1
                                                                1
## 5
               1
                        1
                                  1 experiment standing
                                                                1
                                                                          5
                                                                                   noswitch
## 6
               1
                        1
                                  1 experiment standing
                                                                1
                                                                          6
                                                                                   noswitch
     congruentTrialType cueType shapeType shapeColor response correctResponse correct reactionTime
## 1
                           solid
                                                  blue
            incongruent
                                    square
                                                          right
                                                                            left
                                                                                      no
                                                                                            0.9088130
                                                                                            0.5947349
## 2
            incongruent
                           solid
                                    square
                                                  blue
                                                           left
                                                                            left
                                                                                     yes
## 3
                                                  blue
            incongruent
                         dashed
                                    square
                                                          right
                                                                           right
                                                                                     yes
                                                                                            0.7084870
## 4
                                                  blue
                                                          right
            incongruent
                         dashed
                                    square
                                                                           right
                                                                                     yes
                                                                                            0.5995200
## 5
              congruent
                         dashed
                                    square
                                                yellow
                                                          right
                                                                           right
                                                                                             0.4399409
                                                                                     yes
## 6
              congruent
                         dashed
                                    square
                                               yellow
                                                          right
                                                                           right
                                                                                            0.3847258
                                                                                     yes
##
                    date
                             utcTime
## 1 2021-11-10 10:22:00 1636561737
## 2 2021-11-10 10:22:00 1636561744
## 3 2021-11-10 10:22:00 1636561746
## 4 2021-11-10 10:22:00 1636561748
## 5 2021-11-10 10:22:00 1636561750
```

#### ## 6 2021-11-10 10:22:00 1636561752

```
### check data
#does every person have 392 trials?
ntrials_sub <- task_switching_raw %>%
  group_by(participant) %>%
  summarize(ntrials = n()) %>%
  pull(ntrials)
all(ntrials_sub == 392)
## [1] TRUE
#does every block start with a buffer and have 49 trials?
task_switching_raw <- task_switching_raw %>%
  mutate(condblock = paste0(posture, blockNum))
blocktrials <- task_switching_raw %>%
  group_by(participant, condblock) %>%
  summarize(ntrials = n(), firsttrial = first(switchTrialType))
## 'summarise()' has grouped output by 'participant'. You can override using the '.groups' argument.
all(blocktrials$ntrials == 49)
## [1] TRUE
all(blocktrials$firsttrial == "buffer")
## [1] TRUE
### clean data
#Drop buffer trials
task_switching_raw2 <- task_switching_raw %>%
  filter(switchTrialType != "buffer")
#Recode Correct to 1 and Incorrect to 0
task_switching_raw2$correct_bin <- recode(task_switching_raw2$correct,
                                          "no" = 0,
                                          "ves" = 1)
#Change RTs from seconds to milliseconds
task_switching_raw2 <- task_switching_raw2 %>%
  mutate(reactionTime = reactionTime * 1000)
#Calc overall acc by participant
ts_overall_acc <- task_switching_raw2 %>%
  group_by(participant) %>%
  summarize(Accuracy = mean(correct_bin))
```

```
#find participants with less than 80% accuracy
#2, 8, 15, 44, 49, 51
#First exclusion criteria
low_acc_subs <- ts_overall_acc %>% filter(Accuracy < 0.80) %>%
  pull(participant)
task_switching_raw3 <- task_switching_raw2 %>%
  filter(!(participant %in% low_acc_subs))
#Summary of time-out trials
timeOutTaskSwitch = task_switching_raw3 %>%
                                     filter(is.na(reactionTime))
ftable(posture~congruentTrialType, timeOutTaskSwitch)
##
                      posture sitting standing
## congruentTrialType
                                    82
## congruent
                                             92
## incongruent
                                    98
                                            117
ftable(posture~congruentTrialType~switchTrialType, timeOutTaskSwitch)
##
                   congruentTrialType congruent incongruent
## switchTrialType
## noswitch
                                              76
                                                          88
## switch
                                              98
                                                          127
ftable(congruentTrialType ~ participant, timeOutTaskSwitch)
##
               congruentTrialType congruent incongruent
## participant
## 1
                                           8
                                                       7
## 3
                                           2
                                                       3
## 4
                                           3
                                                       8
## 7
                                           1
                                                       2
                                           2
## 9
                                                       0
## 10
                                                       1
                                           1
## 11
                                           0
                                                       1
## 12
                                           1
                                                       2
## 13
                                                       2
                                           4
## 14
                                           3
                                                       0
## 16
                                           3
                                                        4
## 17
                                           1
                                                       1
                                                       2
## 18
                                           3
## 19
                                          14
                                                      21
## 20
                                           1
                                                       2
## 21
                                           3
                                                      10
```

6

5

2

1

5

3

3

## 22

## 23

## 24

## 25

##	26	1	2
##	27	3	5
##	28	4	4
##	29	4	6
##	30	5	6
##	31	11	13
##	32	7	2
##	33	1	4
##	34	3	1
##	35	2	2
##	36	2	2
##	37	1	4
##	38	6	3
##	39	2	5
##	41	3	3
##	42	5	12
##	43	4	5
##	45	0	8
##	46	5	5
##	47	2	0
##	48	10	7
##	50	2	3
##	52	1	0
##	53	1	1
##	54	3	3
	55	13	27
##	56	6	2
##	57	3	2

## ftable(switchTrialType ~ participant, timeOutTaskSwitch)

##		switchTrialType	noswitch	switch
##	participant			
##	1		6	9
##	3		4	1
##	4		7	4
##	7		2	1
##	9		0	2
##	10		1	1
##	11		1	0
##	12		0	3
##	13		1	5
##	14		0	3
##	16		1	6
##	17		1	1
##	18		3	2
##	19		19	16
##	20		2	1
##	21		6	7
##	22		1	1
##	23		4	7
##	24		1	7
##	25		3	2
##	26		1	2

```
## 27
                                           5
                                                    3
## 28
                                            3
                                                    5
## 29
                                           6
                                                    4
## 30
                                           5
                                                    6
## 31
                                          13
                                                   11
## 32
                                           5
                                                    4
## 33
                                           1
                                                    4
                                                    2
## 34
                                            2
## 35
                                            1
                                                    3
## 36
                                            2
                                                    2
## 37
                                            1
                                                    4
                                                    5
## 38
                                            4
## 39
                                            3
                                                    4
                                                    3
## 41
                                            3
## 42
                                            6
                                                   11
## 43
                                            5
                                                    4
## 45
                                            3
                                                    5
                                            5
                                                    5
## 46
## 47
                                            0
                                                    2
                                                    9
## 48
                                           8
## 50
                                            0
                                                    5
## 52
                                            1
                                                    0
## 53
                                                    2
                                           0
## 54
                                            2
                                                    4
                                          11
                                                   29
## 55
## 56
                                            4
                                                    4
## 57
                                            1
                                                    4
```

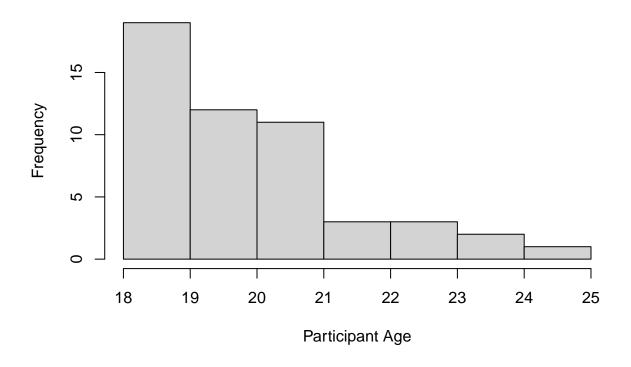
## 'summarise()' has grouped output by 'participant', 'posture', 'congruentTrialType'. You can
## override using the '.groups' argument.

```
values_from =
ts_acc_mean <- ts_acc_mean %>%
  ungroup() %>%
  mutate(across(posture:switchTrialType, as.factor))
str(ts acc mean)
## tibble [408 x 6] (S3: tbl_df/tbl/data.frame)
## $ participant
                       : int [1:408] 1 1 1 1 1 1 1 3 3 ...
## $ posture
                        : Factor w/ 2 levels "sitting", "standing": 1 1 1 1 2 2 2 2 1 1 ...
## $ congruentTrialType: Factor w/ 2 levels "congruent", "incongruent": 1 1 2 2 1 1 2 2 1 1 ...
## $ switchTrialType : Factor w/ 2 levels "noswitch", "switch": 1 2 1 2 1 2 1 2 1 2 ...
## $ Accuracy
                        : num [1:408] 0.96 0.978 0.957 0.88 0.981 ...
## $ PE
                        : num [1:408] 4 2.17 4.35 12 1.89 ...
#Total N = 51 (6 dropped for total acc < 80%)
length(unique(ts_acc_mean$participant))
## [1] 51
Summarize Demographics
demo_raw <- read.csv(paste0(workingdir, "/data",</pre>
                            "/Experiment 2 Data/Task Switching February 24, 2022 13.05.csv"),
                     skip = 1) \% > \%
  slice(-1) %>%
  select(-c(Response.Type,IP.Address, Recipient.Last.Name:Distribution.Channel))
colnames(demo_raw)[10:15] <- c("Gender.Pick", "Gender.Text", "Age", "Race.Pick", "Race.Text", "Eng.First")
dim(demo raw)
## [1] 59 15
#59 records
#first two are test data
# need to match up the 6 dropped participants from behavioral data
demo_df <- demo_raw %>%
 filter(!(X %in% c("test", low_acc_subs)))
dim(demo df)
## [1] 51 15
demo_df <- demo_df %>%
  mutate(Gender.New = ifelse(Gender.Pick %in% c("Man","Woman"), Gender.Pick, Gender.Text),
         Eng.First = toupper(Eng.First))
#gender breakdown
```

switchTrialType),

```
gender_table <- demo_df %>%
  group_by(Gender.New) %>%
  summarize(n = n())
gender_table
## # A tibble: 3 x 2
##
     Gender.New
##
     <chr>
                 <int>
## 1 Man
                    23
## 2 non binaary
                    1
## 3 Woman
                    27
#age breakdown
hist(as.numeric(demo_df$Age),
     main = "Histogram of Participant Ages",
    xlab = "Participant Age")
```

## **Histogram of Participant Ages**



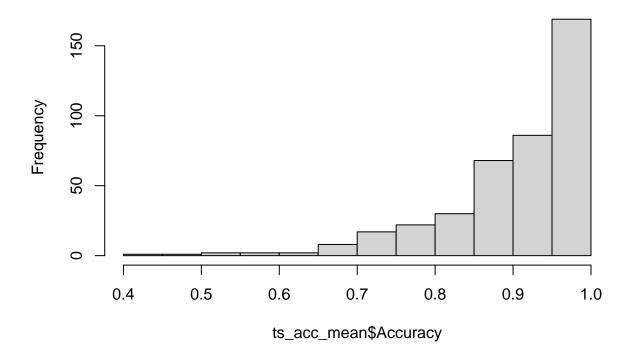
```
age_table <- demo_df %>%
  group_by(Age) %>%
  summarize(n = n())
age_table
```

```
## # A tibble: 8 x 2
## Age n
```

```
<chr> <int>
## 1 18
## 2 19
              10
## 3 20
              12
## 4 21
              11
## 5 22
               3
## 6 23
## 7 24
               2
## 8 25
#age mean and sd
mean_age <- mean(as.numeric(demo_df$Age))</pre>
sd_age <- sd(as.numeric(demo_df$Age))</pre>
kable(matrix(c(mean_age, sd_age), nrow = 1), col.names = c("Mean of Age", "SD of Age"))
                                   Mean of Age
                                                SD of Age
                                       20.21569
                                                   1.73567
#race breakdown
race_table <- demo_df %>%
  group_by(Race.Pick) %>%
  summarize(n = n()) \%>\%
 arrange(desc(n))
race_table
## # A tibble: 6 x 2
     Race.Pick
                                            n
##
     <chr>>
                                        <int>
## 1 White /European American
                                           22
## 2 Black / African American
                                           11
## 3 Hispanic/Latino/Latina/Latinx
                                           11
## 4 Asian /South Pacific Islander
                                            3
## 5 Central Asian /Indian /Pakistani
                                            3
## 6 Native American / American Indian
                                            1
#language breakdown
lang_table <- demo_df %>%
 group_by(Eng.First) %>%
 summarize(n=n())
lang_table
## # A tibble: 2 x 2
##
     Eng.First
                   n
##
     <chr>
              <int>
## 1 NO
                   8
## 2 YES
                  43
```

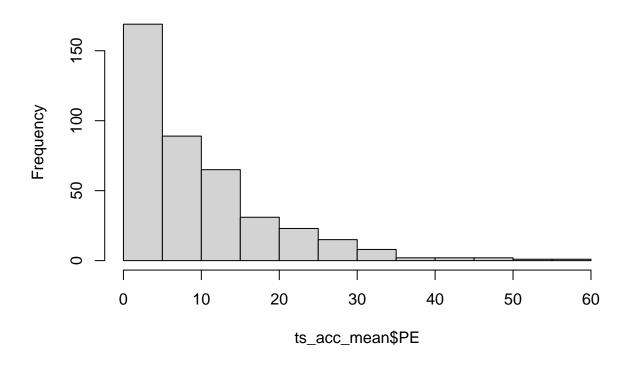
#### Accuracy results

# Histogram of ts\_acc\_mean\$Accuracy



hist(ts\_acc\_mean\$PE)

# Histogram of ts\_acc\_mean\$PE



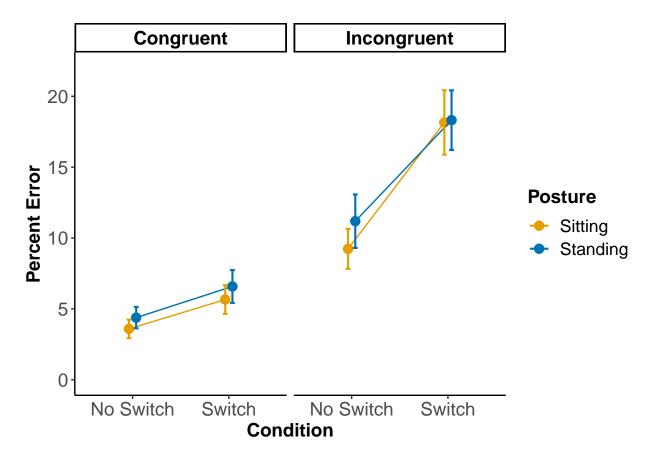
## Warning: Converting "participant" to factor for ANOVA.

## Anova Table (Type 3 tests)

```
##
## Response: PE
##
                                          Effect
                                                    df
                                                         MSE
                                                                         pes p.value
                                        posture 1, 50 88.41
## 1
                                                                        .021
                                                                                 .308
                                                                  1.06
## 2
                             congruentTrialType 1, 50 86.07 99.66 ***
                                                                         .666
                                                                                <.001
## 3
                                switchTrialType 1, 50 28.61 92.04 ***
                                                                        .648
                                                                               <.001
## 4
                     posture:congruentTrialType 1, 50 41.84
                                                                  0.02 < .001
                        posture:switchTrialType 1, 50 24.00
## 5
                                                                  0.74
                                                                        .015
                                                                                 .395
             congruentTrialType:switchTrialType 1, 50 15.15 58.43 ***
                                                                        .539
                                                                                <.001
                                                                                 .268
## 7 posture:congruentTrialType:switchTrialType 1, 50 18.80
                                                                  1.26 .024
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '+' 0.1 ' 1
#Calculate confidence interval: PES for posture x switch/condition interaction
#using ANOVA results
                                 (partial eta-squared)
interaction_effect_CI <- get.ci.partial.eta.squared(accModelTS$anova_table$F[5],</pre>
                                                      accModelTS$anova table$`num Df`[5],
                                                      accModelTS$anova_table$`den Df`[5],
                                                      conf.level = 0.90)
                                                     #90% CI is the convention for PES
interaction_effect_CI
## $LL
## [1] 0
## $UL
## [1] 0.1073579
congruent.labs <- c("Congruent", "Incongruent")</pre>
names(congruent.labs) <- c("1", "2")</pre>
#make plot like Smith et al's
acc_plot <-
  superbPlot(ts_acc_mean_wide,
             WSFactors = c("Condition(2)", "Congruent(2)", "Posture(2)"),
             variables = colnames(ts acc mean wide)[2:9],
             errorbar = "SE", #Tempted to change to CI, should stay SE to be consistent with SMith
             plotStyle = "line",
             factorOrder = c("Condition", "Posture", "Congruent"),
             adjustments = list(purpose = "difference")) +
 theme_classic() +
 theme(axis.text=element_text(size=14),
       panel.background = element_rect(fill = "white", colour = "white", size = 1),
       axis.title = element_text(size=14, face="bold"),
       strip.text = element_text(size = 14, face = "bold",colour = "black", angle = 0),
       legend.text = element_text(size = 13),
       legend.title = element_text(size= 14, face = "bold")) +
  ylim(0, 22) + #Trying to make ylim same as the Smith w/o cutting off error bars
  facet_wrap(vars(Congruent), labeller = labeller(Congruent = congruent.labs)) +
  scale_x_discrete(labels=c("1" = "No Switch", "2" = "Switch"))+
  scale_color_manual(values=c("#E69F00", "#0072B2"),
                     labels = c("Sitting", "Standing")) +
  labs(y = "Percent Error")
```

## superb::FYI: Here is how the within-subject variables are understood:

```
Condition Congruent Posture
##
                                                        variable
##
            1
                                     sitting_congruent_noswitch
            2
##
                       1
                               1
                                       sitting_congruent_switch
##
            1
                       2
                                  sitting_incongruent_noswitch
            2
                       2
                                     sitting_incongruent_switch
##
                               1
##
                               2
                                    standing_congruent_noswitch
##
            2
                       1
                                      standing_congruent_switch
            1
                       2
##
                               2 standing_incongruent_noswitch
##
            2
                       2
                                    standing_incongruent_switch
```



```
#...CALCULATE THE BAYES FACTORS FOR THE ACC ANALYSIS

taskswitchBF = data.frame(ts_acc_mean)
```

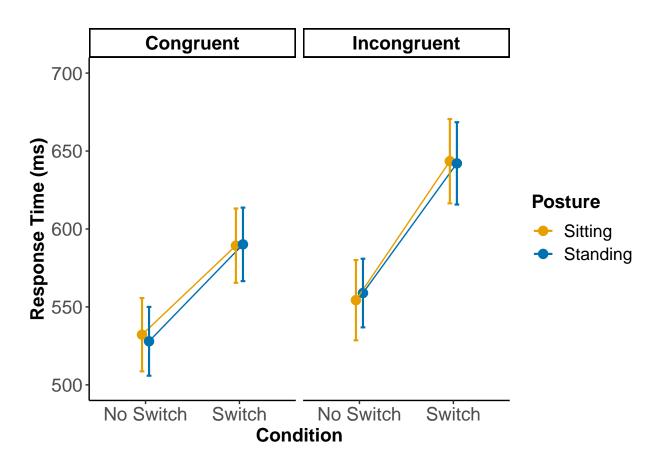
```
taskswitchBF$participant = factor(ts_acc_mean$participant)
taskswitchBF$posture = factor(ts_acc_mean$posture)
taskswitchBF$congruentTrialType = factor(taskswitchBF$congruentTrialType)
taskswitchBF$switchTrialType = factor(taskswitchBF$switchTrialType)
#str(taskswitchBF)
bfValues2 = anovaBF(Accuracy~ posture*switchTrialType+participant,
                  data = taskswitchBF,
                  whichRandom = "participant",
                  method="laplace")
bfValues2
## Bayes factor analysis
## -----
## [1] posture + participant
                                                                        : 0.1836549 ±NA%
## [2] switchTrialType + participant
                                                                        : 5429475 ±NA%
## [3] posture + switchTrialType + participant
                                                                        : 1064550 ±NA%
## [4] posture + switchTrialType + posture:switchTrialType + participant : 168897.1 ±NA%
## Against denominator:
   Accuracy ~ participant
## ---
## Bayes factor type: BFlinearModel, JZS
#...get the Bayes factor for the Null Interaction (vs. model w/no interaction)
bfValues2[3]/bfValues2[4]
## Bayes factor analysis
## -----
## [1] posture + switchTrialType + participant : 6.302952 ±NA%
##
## Against denominator:
## Accuracy ~ posture + switchTrialType + posture:switchTrialType + participant
## ---
## Bayes factor type: BFlinearModel, JZS
1/(bfValues2[3]/bfValues2[4])
## Bayes factor analysis
## [1] posture + switchTrialType + posture:switchTrialType + participant : 0.1586558 ±NA%
## Against denominator:
## Accuracy ~ posture + switchTrialType + participant
## Bayes factor type: BFlinearModel, JZS
```

#### Reaction time results

```
#look at reaction time for correct trials
ts_correct_only <- task_switching_raw3 %>%
  filter(correct_bin == 1)
#Second exclusion criteria
#How many trials faster than 100 ms? Only a single one
sum(ts_correct_only$reactionTime < 100)</pre>
## [1] 1
dim(ts_correct_only)
## [1] 17699
                20
ts_correct_only2 <- ts_correct_only %>% filter(reactionTime >= 100)
#Sanity check, one trial is dropped. Now have 17,698 trials
dim(ts_correct_only2)
## [1] 17698
                20
trimOutputTS = pjRecursiveTrim2(dataSet = ts_correct_only2,
                                  dv = "reactionTime",
                                  splitvars = c("participant",
                                                 "posture",
                                                 "switchTrialType",
                                                 "congruentTrialType"))
trimmedTSData=trimOutputTS[[1]]
totalN.TS = trimOutputTS[[2]]
rejectedTS = trimOutputTS[[3]]
percentTrimmedTS = trimOutputTS[[4]] #this is very close to the percentage trimmed for stroop
#2.14% of trials
percentTrimmedTS
## [1] 2.141485
NcellsTS = trimOutputTS[[5]] # 51 participants * 8 conditions
trimmed_rt_mean_TS <- trimmedTSData %>%
  group_by(participant,
           posture,
           congruentTrialType,
           switchTrialType) %>%
  summarize(mean rt = mean(reactionTime))
```

## 'summarise()' has grouped output by 'participant', 'posture', 'congruentTrialType'. You can
## override using the '.groups' argument.

```
#Convert data to wide format
trimmed_rt_mean_TS_wide <- trimmed_rt_mean_TS %>%
  pivot wider(names from = c(posture,
                            congruentTrialType,
                            switchTrialType),
             values from =
                            mean_rt)
trimmed_RT_plot <-</pre>
  superbPlot(trimmed_rt_mean_TS_wide,
            WSFactors = c("Condition(2)", "Congruent(2)", "Posture(2)"),
            variables = colnames(trimmed_rt_mean_TS_wide)[2:9],
            errorbar = "SE",
            plotStyle = "line",
            factorOrder = c("Condition", "Posture", "Congruent"),
            adjustments = list(purpose = "difference"))+
 theme classic() +
 theme(axis.text=element_text(size=14),
       panel.background = element_rect(fill = "white", colour = "white", size = 1),
       axis.title = element_text(size=14, face="bold"),
       strip.text = element_text(size = 14, face = "bold",colour = "black", angle = 0),
      legend.text = element text(size = 13),
       legend.title = element_text(size= 14, face = "bold")) +
  facet_wrap(vars(Congruent), labeller = labeller(Congruent = congruent.labs)) +
  scale x discrete(labels=c("1" = "No Switch", "2" = "Switch"))+
  scale_color_manual(values=c("#E69F00", "#0072B2"), labels = c("Sitting", "Standing")) +
  ylim(500, 700) +
  labs(y = "Response Time (ms)")
## superb::FYI: Here is how the within-subject variables are understood:
## Condition Congruent Posture
##
           1
                     1
                             1
                                  sitting_congruent_noswitch
           2
##
                     1
                                    sitting_congruent_switch
##
           1
                     2
                             1 sitting_incongruent_noswitch
           2
                     2
##
                             1
                                 sitting_incongruent_switch
##
           1
                    1
                            2 standing_congruent_noswitch
##
           2
                    1
                            2
                                   standing_congruent_switch
##
           1
                    2
                            2 standing_incongruent_noswitch
##
                             2 standing_incongruent_switch
ggsave(trimmed_RT_plot,
      file = "results/plots/fig2_exp2_ts_trimmed_RT_plot.pdf",
      units = "in",
      width = 6.62,
      height = 5.50,
      dpi = 600)
trimmed_RT_plot
```



- ## Warning: Converting "participant" to factor for ANOVA.
- ## Warning: Converting "posture" to factor for ANOVA.
- ## Warning: Converting "congruentTrialType" to factor for ANOVA.
- ## Warning: Converting "switchTrialType" to factor for ANOVA.

```
write.csv(rt.stats.TS[ ,-7],
          file = "results/exp2_Descriptives_trimmed_RT.csv",
         row.names = F)
write.csv(rtModelTS$anova_table, file = "results/exp2_ANOVA_trimmed_RT.csv")
rtModelTS
## Anova Table (Type 3 tests)
## Response: mean_rt
##
                                         Effect
                                                   df
                                                          MSE
                                                                       F
                                                                           pes p.value
                                        posture 1, 50 9922.05
## 1
                                                                    0.00 <.001
                                                                                 .995
                                                                         . 495
## 2
                             congruentTrialType 1, 50 3302.21 48.98 ***
                                                                                 <.001
## 3
                                switchTrialType 1, 50 4166.22 130.17 ***
                                                                                <.001
                                                                    0.17 .003
## 4
                     posture:congruentTrialType 1, 50 1592.27
                                                                                  .679
## 5
                        posture:switchTrialType 1, 50 1519.08
                                                                    0.00 < .001
                                                                                  .951
             congruentTrialType:switchTrialType 1, 50 1252.33 14.32 *** .223
                                                                                 <.001
## 7 posture:congruentTrialType:switchTrialType 1, 50 1552.36
                                                                    0.50 .010
                                                                                .483
## ---
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '+' 0.1 ' ' 1
trimmed_rt_mean_TS2 <- data.frame(trimmed_rt_mean_TS)</pre>
trimmed_rt_mean_TS2$participant = factor(trimmed_rt_mean_TS2$participant)
trimmed_rt_mean_TS2$posture = factor(trimmed_rt_mean_TS2$posture)
trimmed_rt_mean_TS2$congruentTrialType = factor(trimmed_rt_mean_TS2$congruentTrialType)
trimmed_rt_mean_TS2$switchTrialType = factor(trimmed_rt_mean_TS2$switchTrialType)
bfValues2.RT = anovaBF(mean rt ~ posture*switchTrialType+participant,
                   data = trimmed_rt_mean_TS2,
                   whichRandom = "participant",
                   method="laplace")
bfValues2.RT
## Bayes factor analysis
## -----
## [1] posture + participant
                                                                         : 0.1007388
                                                                                        ±NA%
## [2] switchTrialType + participant
                                                                         : 4.227658e+24 ±NA%
## [3] posture + switchTrialType + participant
                                                                         : 4.265145e+23 ±NA%
## [4] posture + switchTrialType + posture:switchTrialType + participant : 6.012989e+22 ±NA%
## Against denominator:
   mean_rt ~ participant
## Bayes factor type: BFlinearModel, JZS
#...get the Bayes factor for the Null Interaction (vs. model w/no interaction)
bfValues2.RT[3]/bfValues2.RT[4]
## Bayes factor analysis
## [1] posture + switchTrialType + participant : 7.093219 \pm NA\%
##
```

```
## Against denominator:
## mean_rt ~ posture + switchTrialType + posture:switchTrialType + participant
## ---
## Bayes factor type: BFlinearModel, JZS

1/(bfValues2.RT[3]/bfValues2.RT[4])

## Bayes factor analysis
## ------
## [1] posture + switchTrialType + posture:switchTrialType + participant : 0.1409797 ±NA%
##
## Against denominator:
## mean_rt ~ posture + switchTrialType + participant
## ---
## Bayes factor type: BFlinearModel, JZS
```

# Experiment 3 - Visual search

### Import and clean data

```
vs_files = list.files(path = "data/Experiment 3 Data/", full.names = T)
vs_files = vs_files[str_detect(vs_files,pattern="(?=.*SJ)(?=.*.txt)")]
merged.VS.data <- ldply(vs_files,</pre>
                         read.delim,
                         header=FALSE,
                         stringsAsFactors = FALSE,
                         sep = "") #for each item in the list apply the function read.delim
#..ADD HEADERS
names(merged.VS.data) = c("sj",
                          "cb",
                         "blockNumber",
                         "blockType",
                         "trialNum",
                         "target",
                          "targetImage",
                          "distractor",
                         "distractorImage",
                         "posture",
                         "setSize",
                         "rt",
                         "resp",
                         "cresp",
                          "ac")
#...look at unique values from both columns
unique(merged.VS.data[c('sj')])
```

## sj

```
## 1
          1
## 265
         10
## 529
         11
## 793
         12
## 1057
         13
## 1321
         14
## 1585
         15
## 1849
         16
## 2113
         17
## 2377
         18
## 2641
         19
## 2905
          2
## 3169
        20
## 3433
         21
## 3697
         22
## 3961
         23
## 4225
         24
## 4489
         25
## 4753
         26
## 5017
         27
## 5281
         28
## 5545
## 5809
          3
## 6073
         30
## 6337
         31
## 6601
         32
## 6865
         33
## 7129
         34
## 7393
         35
## 7657
         36
## 7921
         37
## 8185
         38
## 8449
         39
## 8713
## 8977
         40
## 9241
         41
## 9505
## 9769
        43
## 10033 44
## 10297 45
## 10561 46
## 10825 47
## 11089 48
## 11353 49
## 11617 5
## 11881 50
## 12145
## 12409
## 12673
          8
## 12937
unique(merged.VS.data[c('blockType')])
```

## blockType

```
## 1 practice
## 9 experimental
```

# $\textit{\#..DOES} \ \textit{EACH} \ \textit{SUBJECT} \ \textit{HAVE} \ \textit{THE} \ \textit{SAME} \ \textit{NUMBER} \ \textit{OF} \ \textit{TRIALS}$

ftable(blockType~sj, merged.VS.data)

##	blockType	experimental	practice
## sj	-	-	-
## 1		256	8
## 2		256	8
## 3		256	8
## 4		256	8
## 5		256	8
## 6		256	8
## 7		256	8
## 8		256	8
## 9		256	8
## 10		256	8
## 11		256	8
## 12		256	8
## 13		256	8
## 14		256	8
## 15		256	8
## 16		256	8
## 17		256	8
## 18		256	8
## 19		256	8
## 20		256	8
## 21		256	8
## 22		256	8
## 23		256	8
## 24		256	8
## 25		256	8
## 26		256	8
## 27		256	8
## 28		256	8
## 29		256	8
## 30		256	8
## 31		256	8
## 32		256	8
## 33		256	8
## 34		256	8
## 35		256	8
## 36		256	
## 30		256	8 8
## 38		256	8
			8
## 39 ## 40		256	8
		256	
## 41 ## 42		256	8
		256	8
## 43		256	8
## 44		256	8
## 45		256	8
## 46		256	8

```
## 47
                         256
## 48
                         256
                                    8
                                    8
## 49
                         256
## 50
                         256
                                    8
#...DO WE HAVE EQUAL OBSERVATIONS FOR EACH COUNTERBALANCE
ftable(blockType~cb, merged.VS.data)
      blockType experimental practice
## cb
## 1
                                  200
                        6400
## 2
                        6400
                                  200
#...LOOK FOR MISSING DATA
merged.VS.data[!complete.cases(merged.VS.data),]
## [1] sj
                        cb
                                         blockNumber
                                                         blockType
                                                                         trialNum
                                                                                          target
## [7] targetImage
                                                                          setSize
                        distractor
                                        distractorImage posture
                                                                                          rt
## [13] resp
                        cresp
## <0 rows> (or 0-length row.names)
#... GET RID OF PRACTICE TRIALS
merged.VS.data <- merged.VS.data[!merged.VS.data$blockType=="practice",]</pre>
#.... CHECK TRIALS PER CONDITION
ftable(posture+target+distractor+setSize~sj, merged.VS.data)
```

##		posture	SITTING								STANDING							
##		target	h				s				h				s			
##		${\tt distractor}$	е		u		е		u		е		u		е		u	
##		setSize	4	8	4	8	4	8	4	8	4	8	4	8	4	8	4	8
##	sj																	
##	1		16	16	16	16	16	16	16	16	16	16	16	16	16	16	16	16
##	2		16	16	16	16	16	16	16	16	16	16	16	16	16	16	16	16
##	3		16	16	16	16	16	16	16	16	16	16	16	16	16	16	16	16
##	4		16	16	16	16	16	16	16	16	16	16	16	16	16	16	16	16
##	5			16							16	16	16	16	16	16	16	16
##				16								16						
##				16								16						
##				16								16						
##	_			16								16						
##				16								16						
##				16								16						
##				16								16						
##				16								16						
##				16								16						
##				16								16						
##				16								16						
##				16								16						
##				16								16						
##				16								16						
##	20		16	16	16	16	16	16	16	16	16	16	16	16	16	16	16	16

```
## 21
                     16 16 16 16 16 16 16
                                                  16 16 16 16 16 16 16
## 22
                                                  16 16 16 16 16 16 16
                     16 16 16 16 16 16 16
                     16 16 16 16 16 16 16
## 23
                                                  16 16 16 16 16 16 16 16
## 24
                                                  16 16 16 16 16 16 16
                     16 16 16 16 16 16 16
## 25
                     16 16 16 16 16 16 16
                                                  16 16 16 16 16 16 16 16
## 26
                    16 16 16 16 16 16 16 16
                                                  16 16 16 16 16 16 16 16
## 27
                    16 16 16 16 16 16 16 16
                                                  16 16 16 16 16 16 16
## 28
                    16 16 16 16 16 16 16 16
                                                  16 16 16 16 16 16 16 16
## 29
                    16 16 16 16 16 16 16 16
                                                  16 16 16 16 16 16 16
## 30
                    16 16 16 16 16 16 16 16
                                                  16 16 16 16 16 16 16 16
## 31
                    16 16 16 16 16 16 16 16
                                                  16 16 16 16 16 16 16
## 32
                    16 16 16 16 16 16 16
                                                  16 16 16 16 16 16 16 16
                     16 16 16 16 16 16 16
## 33
                                                  16 16 16 16 16 16 16
## 34
                    16 16 16 16 16 16 16 16
                                                  16 16 16 16 16 16 16
## 35
                    16 16 16 16 16 16 16
                                                  16 16 16 16 16 16 16
## 36
                    16 16 16 16 16 16 16
                                                  16 16 16 16 16 16 16
## 37
                    16 16 16 16 16 16 16 16
                                                  16 16 16 16 16 16 16
## 38
                    16 16 16 16 16 16 16
                                                  16 16 16 16 16 16 16
## 39
                    16 16 16 16 16 16 16 16
                                                  16 16 16 16 16 16 16 16
## 40
                    16 16 16 16 16 16 16
                                                  16 16 16 16 16 16 16
## 41
                    16 16 16 16 16 16 16
                                                  16 16 16 16 16 16 16
## 42
                    16 16 16 16 16 16 16 16
                                                  16 16 16 16 16 16 16 16
## 43
                    16 16 16 16 16 16 16 16
                                                  16 16 16 16 16 16 16 16
## 44
                    16 16 16 16 16 16 16 16
                                                  16 16 16 16 16 16 16
## 45
                    16 16 16 16 16 16 16
                                                  16 16 16 16 16 16 16 16
## 46
                    16 16 16 16 16 16 16 16
                                                  16 16 16 16 16 16 16
## 47
                     16 16 16 16 16 16 16
                                                  16 16 16 16 16 16 16 16
## 48
                     16 16 16 16 16 16 16
                                                  16 16 16 16 16 16 16
## 49
                     16 16 16 16 16 16 16 16
                                                  16 16 16 16 16 16 16 16
## 50
                     16 16 16 16 16 16 16
                                                  16 16 16 16 16 16 16
#... UNLIKE THE STROOP, PARTICIPANTS WERE ALLOWED TO TAKE LONGER THAN 1500MS BUT WERE GIVEN A WARNING
#... TRIALS LONGER THAN 1500 MS will be considered errors (i.e., they will be dropped in RT but kept in
#... Set values in the ac column to 0 on trials where a response is > = 1500
#...check that only experimental trials are left
unique(merged.VS.data$blockType)
## [1] "experimental"
write.table(merged.VS.data, file = "results/exp3_merged_vs_data.txt", row.names = F)
#...count trials
totalTrialsVS = dim(merged.VS.data)[1]
observationDataVS = data.frame(ftable(blockType~sj, merged.VS.data))[,c(1,3)]
#...qet the number of extreme trials <100 - anticipatory or fast responses
merged.VS.data= merged.VS.data[!merged.VS.data$rt<=100,]</pre>
validRTTrialsVS = dim(merged.VS.data)[1]
observationDataVS$validTrials = data.frame(ftable(blockType~sj, merged.VS.data))[,c(3)]
print(paste("percent invalid trials = ", ((totalTrialsVS-validRTTrialsVS)/totalTrialsVS)*100))
## [1] "percent invalid trials = 0"
```

```
#...get the number of time outs
timeOutVS = merged.VS.data %>% filter(rt >= 1500)
dim(timeOutVS)[1]
```

# ## [1] 168

# ftable(posture~setSize, timeOutVS)

# ftable(posture~sj, timeOutVS)

##		posture	SITTING	STANDING
##	sj			
##	2		9	12
##	3		2	0
##	4		6	2
##	5		3	1
##	6		0	1
##	7		1	2
##	9		4	6
##	10		7	3
##	11		3	2
##	12		2	7
##	13		4	1
##	15		1	2
##	17		2	0
##	18		0	1
##	19		3	2
##	20		1	2
##	22		2	2
##	24		0	2
##	25		2	4
##	27		0	2
##	28		0	1
##	29		5	6
##	30		2	0
##	31		3	0
##	33		0	1
##	35		3	2
##	36		0	1
##	37		0	1
##	39		2	3
##	40		3	3
##	41		0	2
##	42		2	1
##	44		4	2
##	45		3	0
##	46		3	0

```
## 47
                    0
                             1
## 48
                    1
                             0
                    3
                             2
## 49
## 50
                    2
                             0
ftable(blockType~setSize, timeOutVS)
##
           blockType experimental
## setSize
## 4
                               45
## 8
                              123
merged.VS.data.conventional <- merged.VS.data %>% filter(rt < 1500)
write.csv(merged.VS.data.conventional,
          file = "results/exp3 merged vs data conv.csv",
          row.names = F)
#...this code changes the 1550ms+ trials into errors
merged.VS.data.orig <- merged.VS.data
merged.VS.data$ac[merged.VS.data$rt>=1500] = 0
vsCorrect = merged.VS.data[merged.VS.data$ac ==1,]
errorsRemovedVS = dim(vsCorrect)[1]
observationDataVS$correctTrials = data.frame(ftable(blockType~sj, vsCorrect))[,c(3)]
trimInfo = data.frame(totalTrialsVS, validRTTrialsVS, errorsRemovedVS)
head(trimInfo)
   totalTrialsVS validRTTrialsVS errorsRemovedVS
## 1
             12800
                             12800
                                              12397
######################################
#...CHECK 20% CRITERION
######################################
observationDataVS$percentLoss = ((observationDataVS$Freq-observationDataVS$correctTrials)/observationDa
sum(observationDataVS$percentLoss>20)
## [1] O
#...None!
#...RUN TRIMMING PROCEDURE
tempList = pjRecursiveTrim2(vsCorrect, #...dataset
                            "rt", #...dependent variables
                            c("sj",
                              "cb",
                              "setSize",
                              "posture")) #.independent variables
trimmedData=tempList[[1]]
```

```
totalN = tempList[[2]]
rejected = tempList[[3]]
percentTrimmed = tempList[[4]]
Ncells = tempList[[5]]
print(paste("Percent of outliers removed: ",round(percentTrimmed,3)))
## [1] "Percent of outliers removed: 1.339"
#...get the trimming info
output.out= data.frame(totalN, rejected,percentTrimmed,Ncells)
head(output.out)
   totalN rejected percentTrimmed Ncells
## 1 12397
               166
                         1.339034
#...get mean error data
vsPE = plyr::ddply(merged.VS.data,
                .(sj,cb,setSize, posture),
                summarise,
                meanPE = 100 - (mean(ac)*100))
head(vsPE)
   sj cb setSize posture meanPE
## 1 1 1 4 SITTING 0.0000
## 2 1 1
              4 STANDING 0.0000
## 3 1 1
             8 SITTING 0.0000
## 4 1 1
              8 STANDING 0.0000
## 5 2 1
              4 SITTING 4.6875
## 6 2 1
              4 STANDING 4.6875
vsRT = plyr::ddply(trimmedData,
                .(sj, cb, setSize, posture),
                summarise,
                meanRT = mean(rt))
#...combine the RT and error data
vsCombined = cbind(vsRT,meanPE =vsPE$meanPE)
str(vsCombined)
## 'data.frame':
                  200 obs. of 6 variables:
          : int 1111222333...
## $ sj
## $ cb : int 1 1 1 1 1 1 1 1 1 ...
## $ setSize: int 4 4 8 8 4 4 8 8 4 4 ...
## $ posture: chr "SITTING" "STANDING" "SITTING" "STANDING" ...
## $ meanRT : num 677 593 736 620 792 ...
## $ meanPE : num 0 0 0 0 4.69 ...
#...set as factors
vsCombined$sj = factor(vsCombined$sj)
vsCombined$cb = factor(vsCombined$cb)
```

```
vsCombined$setSize = factor(vsCombined$setSize)
vsCombined$postureFactor = factor(vsCombined$posture)
summary(vsCombined$cb)
```

```
## 1 2
## 100 100
```

#### Reaction time results

```
##
                  Effect DFn DFd
                                         SSn
                                                    SSd
                                                                               p p<.05
## 1
             (Intercept) 1 49 1.084958e+08 1189588.17 4.469020e+03 8.326740e-50
            postureFactor 1 49 2.052064e+04 153738.29 6.540411e+00 1.369090e-02
## 2
                  setSize 1 49 3.574624e+05 46863.03 3.737628e+02 1.414816e-24
## 3
## 4 postureFactor:setSize 1 49 2.246613e+01 35654.35 3.087534e-02 8.612429e-01
##
             ges
## 1 9.870285e-01
## 2 1.418774e-02
## 3 2.004492e-01
## 4 1.575613e-05
```

```
## [1] "(Intercept): F(1, 49) = 4469.02, MSE = 24277.31, p = 0, partialEtaSq = 0.9892"
## [2] "postureFactor: F(1, 49) = 6.54, MSE = 3137.516, p = 0.014, partialEtaSq = 0.1178"
## [3] "setSize: F(1, 49) = 373.763, MSE = 956.388, p = 0, partialEtaSq = 0.8841"
## [4] "postureFactor:setSize: F(1, 49) = 0.031, MSE = 727.64, p = 0.861, partialEtaSq = 6e-04"
```

```
#...CALCULATE THE BAYES FACTORS FOR THE RT ANALYSIS
bfValues3 = anovaBF(meanRT~setSize*postureFactor+sj,
                 data = vsCombined,
                 whichRandom = "sj",
                 method="laplace")
bfValues3
## Bayes factor analysis
## -----
## [1] setSize + sj
                                                        : 2.916459e+26 ±NA%
## [2] postureFactor + sj
                                                        : 1.51507
## [3] setSize + postureFactor + sj
                                                        : 1.321058e+28 ±NA%
## [4] setSize + postureFactor + setSize:postureFactor + sj : 2.585184e+27 ±NA%
##
## Against denominator:
   meanRT ~ sj
## ---
## Bayes factor type: BFlinearModel, JZS
\#... get the Bayes factor for the Null Interaction
bfValues3[3]/bfValues3[4]
## Bayes factor analysis
## [1] setSize + postureFactor + sj : 5.110113 \pmNA%
## Against denominator:
## meanRT ~ setSize + postureFactor + setSize:postureFactor + sj
## Bayes factor type: BFlinearModel, JZS
1/(bfValues3[3]/bfValues3[4])
## Bayes factor analysis
## -----
## [1] setSize + postureFactor + setSize:postureFactor + sj : 0.1956904 ±NA%
## Against denominator:
## meanRT ~ setSize + postureFactor + sj
## ---
## Bayes factor type: BFlinearModel, JZS
# GET DIFFERENCE SCORES - SEARCH RATE
wideData = dcast(vsCombined, #the name of the dataframe you want to reshape
               sj+cb #row variables
               ~posture+setSize, #row variables ~ column variables
               value.var = "meanRT")
head(wideData)
```

```
sj cb SITTING_4 SITTING_8 STANDING_4 STANDING_8
## 1 1 1 676.5238 735.5397
                                593.1129 619.6406
## 2 2 1 792.4590 931.9474
                                815.7213
                                          993.5000
## 3 3 1 721.2787 827.2222
                                654.1639 774.5238
## 4 4 1 695.7119 741.9298
                                660.2632
                                         653.0172
## 5 5 1 693.6034 839.2903
                                705.1967
                                          759.7419
## 6 6 1 625.3750 694.0484
                                592.4531
                                          687.5645
wideData$sittingEffect = (wideData$SITTING_8-wideData$SITTING_4)/4
wideData$standingEffect = (wideData$STANDING_8-wideData$STANDING_4)/4
wideData$interaction = wideData$sittingEffect - wideData$standingEffect
searchratestand = mean(wideData$standingEffect) #...search rate in standing condition
searchratesit = mean(wideData$sittingEffect) #...search rate in the sitting condition
searchratestand
## [1] 21.30589
searchratesit
## [1] 20.97073
#One-sample t-tests
t.test(wideData$standingEffect)
##
##
  One Sample t-test
## data: wideData$standingEffect
## t = 16.69, df = 49, p-value < 2.2e-16
## alternative hypothesis: true mean is not equal to 0
## 95 percent confidence interval:
## 18.74050 23.87127
## sample estimates:
## mean of x
## 21.30589
t.test(wideData$sittingEffect)
##
## One Sample t-test
## data: wideData$sittingEffect
## t = 13.055, df = 49, p-value < 2.2e-16
## alternative hypothesis: true mean is not equal to 0
## 95 percent confidence interval:
## 17.74261 24.19884
## sample estimates:
## mean of x
## 20.97073
```

```
##
## Exact binomial test
##
## data: length(wideData$interaction[wideData$interaction >= 0]) and length(unique(vsCombined$sj))
## number of successes = 25, number of trials = 50, p-value = 1
## alternative hypothesis: true probability of success is not equal to 0.5
## 95 percent confidence interval:
## 0.355273 0.644727
## sample estimates:
## probability of success
## 0.5
```

#### Percent error results

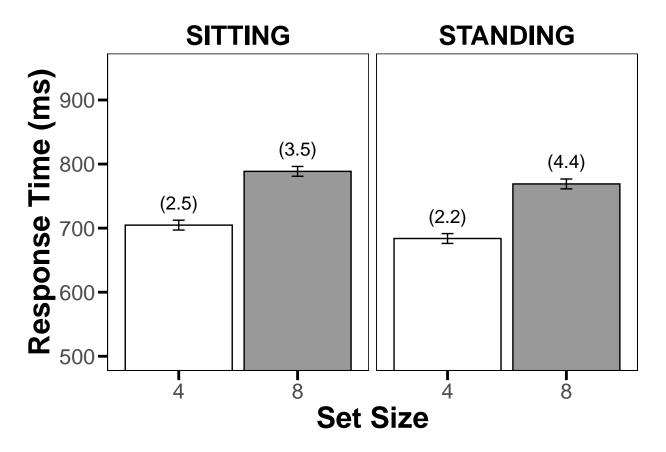
```
## $ANOVA
##
                    Effect DFn DFd
                                           SSn
                                                     SSd
                                                                               p p<.05
## 1
               (Intercept) 1 49 1982.531738 1158.9478 83.8209098 3.463466e-12
                                                                                      * 0.504342884
## 2
            postureFactor
                                49
                                      3.527832 227.7954 0.7588554 3.879351e-01
                                                                                        0.001807368
                             1
                   setSize
                             1 49
                                    129.504395 343.5181 18.4727266 8.162026e-05
                                                                                      * 0.062324860
## 4 postureFactor:setSize
                                     20.520020 218.1274 4.6096032 3.676850e-02
                                                                                      * 0.010422027
                             1 49
## $aov
##
## Call:
## aov(formula = formula(aov_formula), data = data)
##
## Grand Mean: 3.148438
##
## Stratum 1: sj
##
## Terms:
                   Residuals
## Sum of Squares
                    1158.948
## Deg. of Freedom
                          49
##
## Residual standard error: 4.863332
##
```

```
## Stratum 2: sj:postureFactor
##
## Terms:
##
                  postureFactor Residuals
## Sum of Squares
                        3.52783 227.79541
## Deg. of Freedom
                              1
## Residual standard error: 2.156128
## 1 out of 2 effects not estimable
## Estimated effects are balanced
## Stratum 3: sj:setSize
## Terms:
##
                   setSize Residuals
## Sum of Squares 129.5044 343.5181
## Deg. of Freedom
                        1
##
## Residual standard error: 2.647749
## 1 out of 2 effects not estimable
## Estimated effects are balanced
## Stratum 4: sj:postureFactor:setSize
## Terms:
                  postureFactor:setSize Residuals
## Sum of Squares
                        20.52002 218.12744
## Deg. of Freedom
## Residual standard error: 2.109877
## Estimated effects are balanced
err.VS.MSE <- errModelVS$ANOVA$SSd/errModelVS$ANOVA$DFd
exp3 ANOVA PE uncon <-
paste(errModelVS$ANOVA$Effect,": F(",
      errModelVS$ANOVA$DFn,
      errModelVS$ANOVA$DFd,
      ") = ",
     round(errModelVS$ANOVA$F,3),
      ", MSE = ",
     round(err.VS.MSE,3),
      ", p = ",
     round(errModelVS$ANOVA$p,3),
      ", partialEtaSq = ",
     round(errModelVS$ANOVA$SSn/(errModelVS$ANOVA$SSn+errModelVS$ANOVA$SSd),4),sep="")
write.csv(exp3_ANOVA_PE_uncon, "results/exp3_ANOVA_PE_uncon.csv")
wideData = dcast(vsCombined, #the name of the dataframe you want to reshape
                 sj+cb #row variables
                 ~posture+setSize, #row variables ~ column variables
                value.var = "meanPE")
```

```
head(wideData)
    sj cb SITTING_4 SITTING_8 STANDING_4 STANDING_8
## 1 1 1 0.0000
                     0.0000
                                 0.0000
                                            0.0000
## 2 2 1
            4.6875 10.9375
                                 4.6875
                                           15.6250
## 3 3 1 1.5625
                      1.5625
                                 0.0000
                                            0.0000
## 4 4 1 6.2500 10.9375
                                 4.6875
                                            6.2500
## 5 5 1 3.1250
                                            1.5625
                     3.1250
                                 1.5625
## 6 6 1 0.0000
                    1.5625
                                 0.0000
                                            1.5625
wideData$sittingEffect = (wideData$SITTING_8-wideData$SITTING_4)/4
wideData$standingEffect = (wideData$STANDING 8-wideData$STANDING 4)/4
wideData$interaction = wideData$sittingEffect - wideData$standingEffect
searchratestand = mean(wideData$standingEffect) #...search rate in standing condition
searchratesit = mean(wideData$sittingEffect) #...search rate in the sitting condition
searchratestand
## [1] 0.5625
searchratesit
## [1] 0.2421875
#One-sample t-tests
t.test(wideData$standingEffect)
##
## One Sample t-test
##
## data: wideData$standingEffect
## t = 4.0858, df = 49, p-value = 0.0001623
## alternative hypothesis: true mean is not equal to 0
## 95 percent confidence interval:
## 0.2858399 0.8391601
## sample estimates:
## mean of x
     0.5625
##
t.test(wideData$sittingEffect)
##
## One Sample t-test
## data: wideData$sittingEffect
## t = 2.4588, df = 49, p-value = 0.01752
## alternative hypothesis: true mean is not equal to 0
## 95 percent confidence interval:
## 0.04424588 0.44012912
## sample estimates:
## mean of x
## 0.2421875
```

```
#...CALCULATE THE BAYES FACTORS FOR THE RT ANALYSIS
bfValues3.error = anovaBF(meanPE~setSize*postureFactor+sj,
                  data = vsCombined,
                  whichRandom = "sj",
                  method="laplace")
bfValues3.error
## Bayes factor analysis
## -----
                                                           : 4888.005 ±NA%
## [1] setSize + sj
## [2] postureFactor + sj
                                                           : 0.1823325 ±NA%
## [3] setSize + postureFactor + sj
                                                           : 937.7872 ±NA%
## [4] setSize + postureFactor + setSize:postureFactor + sj : 997.3734 ±NA%
## Against denominator:
## meanPE ~ sj
## ---
## Bayes factor type: BFlinearModel, JZS
#...get the Bayes factor for the Null Interaction
bfValues3.error[3]/bfValues3.error[4]
## Bayes factor analysis
## [1] setSize + postureFactor + sj : 0.9402568 ±NA%
## Against denominator:
## meanPE ~ setSize + postureFactor + setSize:postureFactor + sj
## Bayes factor type: BFlinearModel, JZS
1/(bfValues3.error[3]/bfValues3.error[4])
## Bayes factor analysis
## -----
## [1] setSize + postureFactor + setSize:postureFactor + sj : 1.063539 ±NA%
##
## Against denominator:
## meanPE ~ setSize + postureFactor + sj
## Bayes factor type: BFlinearModel, JZS
Make plots for visual search
```

```
mat=TRUE,
                   digits = 1)
graphRT3 = graphRT3[,c("group1","group2","mean","se")]
graphPE3 = graphPE3[,c("group1","group2","mean","se")]
names(graphRT3) = c("posture", "setSize", "mean", "se")
names(graphPE3) = c("posture", "setSize", "mean", "se")
graphRT3$posture = str_to_upper(graphRT3$posture)
#..calculate the within subjects confidence intervals based on loftus and masson
#..the confidence intervals are based on the interaction term.
graphRT3$se = sqrt((rt.VS.MSE[4])/length(unique(vsCombined$sj)))
graphPE3$se= sqrt((err.VS.MSE[4])/length(unique(vsCombined$sj)))
#..calculate the within subjects confidence intervals based on loftus and masson
#..the confidence intervals are based on the interaction term.
critT3 = qt(p=.025,df=length(unique(vsCombined$sj))-2,lower.tail =FALSE)
#---add the min and max for the confidence intervals
graphRT3$min = graphRT3$mean - (graphRT3$se*critT3)
graphRT3$max = graphRT3$mean + (graphRT3$se*critT3)
####GET AC DATA FROM twoAnimalWordsPRPac.R
graphRT3$ac = paste("(",format(round(graphPE3$mean,digits=1),nsmall = 1),")",sep="")
head(graphRT3)
##
       posture setSize mean
                                se
                                       min
                                               max
## X11 SITTING 4 704.7 3.814813 697.0298 712.3702 (2.5)
## X12 STANDING
                 4 683.8 3.814813 676.1298 691.4702 (2.2)
                  8 788.6 3.814813 780.9298 796.2702 (3.5)
## X13 SITTING
## X14 STANDING
                 8 769.0 3.814813 761.3298 776.6702 (4.4)
graphRT3$vAdj = 35 #down
graphRT3$vAdj[graphRT$setSize=="incongruent"]=35 #up
graphRT3$hAdj = 0 #right
#graphRT$hAdj[graphRT$posture=="SITTING"]=-60 #left
graphRT3$congruency = factor(graphRT3$setSize,labels = c("4","8"))
interactionPlot3 <- ggplot(graphRT3, aes(setSize, mean, group=posture)) +</pre>
 theme(legend.position = "none")+
 scale_fill_manual(values=c("#FFFFFF","#999999","#FFFFFF","#999999"))+
 coord cartesian(ylim=c(500,950),expand=TRUE)+
 scale_y_continuous(breaks = round(seq(500, 950, by = 100),0))+
```



```
ggsave(interactionPlot3,
    file = "results/plots/fig4_exp3_visual_search_interaction_plot.pdf",
    units = "in",
    width = 8.5,
    height = 5,
    dpi = 600)
```

# Reproduce results from Smith et al.

```
# Data source: The data from all the experiments are available at:
# http://rabrams.net under the Resources tab.
### Experiment 1 (Stroop)
#load acc data
Smith_Exp1_acc <- read_excel("data/smith_data.xlsx",</pre>
                             sheet = "Exp1Acc",
                             n_max = 14) #Sample size in Smith
#load rt data
Smith_Exp1_rt <- read_excel("data/smith_data.xlsx",</pre>
                            sheet = "Exp1RT",
                            n \max = 14
#Restructure from wide to narrow, using tidyr
Smith_Exp1_acc_narrow <- Smith_Exp1_acc %>%
 pivot_longer(cols = sit_neut:sta_con, names_to = "condition", values_to = "acc") %%
  separate(col = condition, into = c("posture", "con"))
Smith_Exp1_rt_narrow <- Smith_Exp1_rt %>%
  pivot_longer(cols = sit_neut:sta_con, names_to = "condition", values_to = "rt") %>%
  separate(col = condition, into = c("posture", "con"))
Smith_Exp1 <- merge(Smith_Exp1_acc_narrow, Smith_Exp1_rt_narrow)</pre>
Smith_exp1_anova_acc <- aov_ez(data = Smith_Exp1,</pre>
                                dv = 'acc',
                                id = 'subj'
                                within = c('posture', 'con'),
                                anova_table = list(es = "pes", correction = "none"),
                                type = 3)
kable(nice(Smith_exp1_anova_acc), caption = "ANOVA results for Smith Exp 1 - accuracy")
```

Table 2: ANOVA results for Smith Exp 1 - accuracy

Effect	df	MSE	F	pes	p.value
posture	1, 13	3.19	0.51	.038	.488
con	2, 26		3.76 *	.224	.037
posture:con	2, 26		1.47	.101	.250

Table 3: ANOVA results for Smith Exp 1 - RT

Effect	df	MSE	F	pes	p.value
posture	1, 13 2, 26	816.34 150.32		.007 .210	.768 .047
posture:con	$\frac{2}{2}, \frac{20}{26}$		4.73 *	.267	.018

```
### Experiment 2 (Task-switching)
#load acc data
Smith_Exp2_acc <- read_excel("data/smith_data.xlsx",</pre>
                              sheet = "Exp2Acc",
                              n \max = 30)
#load rt data
Smith_Exp2_rt <- read_excel("data/smith_data.xlsx",</pre>
                             sheet = "Exp2RT",
                             n_max = 30)
#Restructure from wide to narrow, using tidyr
Smith_Exp2_acc_narrow <- Smith_Exp2_acc %>%
  pivot_longer(cols = sit_congruent_noswitch:stand_incongruent_switch,
               names_to = "condition", values_to = "acc") %>%
  separate(col = condition, into = c("posture", "con", "switch"))
Smith_Exp2_rt_narrow <- Smith_Exp2_rt %>%
  pivot_longer(cols = sit_congruent_noswitch:stand_incongruent_switch,
               names_to = "condition", values_to = "rt") %>%
  separate(col = condition, into = c("posture", "con", "switch"))
Smith_Exp2 <- merge(Smith_Exp2_acc_narrow, Smith_Exp2_rt_narrow)</pre>
Smith_exp2_anova_acc <- aov_ez(data = Smith_Exp2,</pre>
                                dv = 'acc',
                                id = 'subj',
                                within = c('posture', 'con', 'switch'),
                                anova_table = list(es = "pes", correction = "none"),
                                type = 3)
kable(nice(Smith_exp2_anova_acc), caption = "ANOVA results for Smith Exp 2 - accuracy")
```

Table 4: ANOVA results for Smith Exp 2 - accuracy

Effect	$\mathrm{d}\mathrm{f}$	MSE	F	pes	p.value
posture	1, 29	0.00	2.86	.090	.101
con	1, 29	0.00	67.40 ***	.699	<.001
switch	1, 29	0.00	62.94 ***	.685	<.001
posture:con	1, 29	0.00	1.68	.055	.205
posture:switch	1, 29	0.00	5.54 *	.160	.026
con:switch	1, 29	0.00	23.34 ***	.446	<.001
posture:con:switch	1, 29	0.00	0.50	.017	.484

Table 5: ANOVA results for Smith Exp 2 - RT

Effect	df	MSE	F	pes	p.value
posture	1, 29	0.02	0.03	.001	.856
con	1, 29	0.00	40.95 ***	.585	<.001
switch	1, 29	0.00	115.10 ***	.799	<.001
posture:con	1, 29	0.00	0.49	.017	.489
posture:switch	1, 29	0.00	0.10	.004	.751
con:switch	1, 29	0.00	4.77 *	.141	.037
posture:con:switch	1, 29	0.00	0.67	.023	.420

```
### Experiment 3 (Visual Search)
#load acc data
Smith_Exp3_acc <- read_excel("data/smith_data.xlsx",</pre>
                              sheet = "Exp3Acc",
                              n_{max} = 12) \%
  select(subj:sit8)
#load rt data
Smith_Exp3_rt <- read_excel("data/smith_data.xlsx",</pre>
                             sheet = "Exp3RT",
                             n_{max} = 12)\%>\%
  select(subj:sit8)
#Restructure from wide to narrow, using tidyr
Smith_Exp3_acc_narrow <- Smith_Exp3_acc %>%
  pivot_longer(cols = stand4:sit8, names_to = "condition", values_to = "acc") %>%
  separate(col = condition, into = c("posture", "set.size"), sep = -1)
Smith_Exp3_rt_narrow <- Smith_Exp3_rt %>%
  pivot_longer(cols = stand4:sit8, names_to = "condition", values_to = "rt") %>%
  separate(col = condition, into = c("posture", "set.size"), sep = -1)
Smith_Exp3 <- merge(Smith_Exp3_acc_narrow, Smith_Exp3_rt_narrow)</pre>
Smith_exp3_anova_acc <- aov_ez(data = Smith_Exp3,
                                dv = 'acc'
                                id = 'subj',
                                within = c('posture', 'set.size'),
                                anova_table = list(es = "pes", correction = "none"),
                                type = 3)
kable(nice(Smith_exp3_anova_acc), caption = "ANOVA results for Smith Exp 3 - accuracy")
```

Table 6: ANOVA results for Smith Exp 3 - accuracy

Effect	df	MSE	F	pes	p.value
posture	1, 11	4.61	0.76	.065	.401
set.size	1, 11	1.75	3.44 +	.238	.090
posture:set.size	1, 11	1.38	7.96 *	.420	.017

Table 7: ANOVA results for Smith Exp 3 - RT

Effect	df	MSE	F	pes	p.value
posture	1, 11	2323.81	0.23	.021	.639
set.size		473.24	81.88 ***	.882	<.001
posture:set.size		298.96	5.91 *	.350	.033

# Overall summary plots: Smith and replication

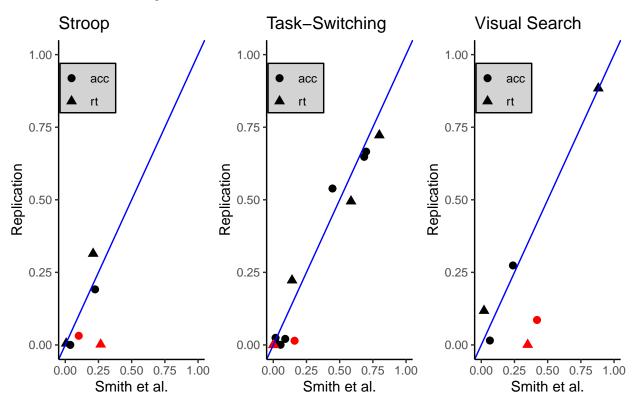
```
smith_anovas <- lst(Smith_exp1_anova_acc$anova_table,</pre>
                    Smith_exp1_anova_rt$anova_table,
                    Smith_exp2_anova_acc$anova_table,
                    Smith_exp2_anova_rt$anova_table,
                    Smith_exp3_anova_acc$anova_table,
                    Smith_exp3_anova_rt$anova_table)
repl_anovas <- lst(aov_ez(data = stroopCombined,</pre>
                           dv = "meanPE",
                           id = "sj",
                           within = c("posture", "congruency"),
                           type = 3,
                           anova_table = list(es = "pes")),
                   aov_ez(data = stroopCombined,
                           dv = "meanRT",
                           id = "sj",
                           within = c("posture", "congruency"),
                           type = 3,
                           anova_table = list(es = "pes")),
                   accModelTS,
                   rtModelTS,
                   aov_ez(data = vsCombined,
                           dv = "meanPE",
                           id = "sj",
```

```
within = c("postureFactor", "setSize"),
                          type = 3,
                          anova_table = list(es = "pes")),
                   aov ez(data = vsCombined,
                          dv = "meanRT",
                          id = "sj",
                          within = c("postureFactor", "setSize"),
                          type = 3,
                          anova_table = list(es = "pes")))
for (i in 1:6){
  smith anovas[[i]] <- smith anovas[[i]] %>%
   rownames_to_column() %>%
   as.data.frame() %>%
   rowwise() %>%
   mutate(LL = get.ci.partial.eta.squared(F, `num Df`, `den Df`, conf.level = 0.9)$LL,
           UL = get.ci.partial.eta.squared(F, `num Df`, `den Df`, conf.level = 0.9)$UL)
  repl_anovas[[i]] <- repl_anovas[[i]]$anova_table %>%
   rownames_to_column() %>%
   as.data.frame() %>%
   rowwise() %>%
   mutate(LL = get.ci.partial.eta.squared(F, `num Df`, `den Df`, conf.level = 0.9)$LL,
           UL = get.ci.partial.eta.squared(F, `num Df`, `den Df`, conf.level = 0.9)$UL)
}
###Exp1 (Stroop)
smith.stroop <- smith_anovas[[1]] %>%
 ungroup() %>%
  bind_rows(smith_anovas[[2]]) %>%
  select(Effect = rowname, pes, LL, UL) %>%
  mutate(dv = rep(c("acc","rt"), each = 3), col = rep(c("black","black","red"),2))
repl.stroop <- repl_anovas[[1]] %>%
  ungroup() %>%
  bind_rows(repl_anovas[[2]]) %>%
  select(Effect = rowname, pes, LL, UL) %>%
  mutate(dv = rep(c("acc","rt"), each = 3), col = rep(c("black","black","red"),2),
         Effect = smith.stroop$Effect)
stroop.effects <- merge(smith.stroop, repl.stroop,</pre>
                        by = c("Effect", "dv"), suffixes = c("Smith", "Replication"))
stroop.plot \leftarrow ggplot(data = stroop.effects, aes(x = pesSmith, y = pesReplication, shape = dv)) +
  geom_point(size = 2.5, col = stroop.effects$colSmith) +
  xlim(0, 1.00) +
  ylim(0, 1.00) +
  geom_abline(slope = 1, intercept = 0, col = "blue") +
  theme_classic() +
  theme(legend.position = c(0.2, 0.85),
```

```
legend.background = element_rect(colour = "black",
                                         linetype = "solid",
                                         fill = "lightgray"),
        legend.title = element_blank(),
        legend.margin=margin(-3,5,0,0)) +
  labs(y = "Replication", x = "Smith et al.", title = "Stroop")
###Exp2 (Task-switching)
smith.ts <- smith_anovas[[3]] %>%
  ungroup() %>%
  bind_rows(smith_anovas[[4]]) %>%
  select(Effect = rowname, pes, LL, UL) %>%
  mutate(dv = rep(c("acc","rt"), each = 7),
         col = rep(c("black","black","black","black","red","black","black"),2))
repl.ts <- repl_anovas[[3]] %>%
  ungroup() %>%
  bind_rows(repl_anovas[[4]]) %>%
  select(Effect = rowname, pes, LL, UL) %>%
  mutate(dv = rep(c("acc","rt"), each = 7),
         col = rep(c("black","black","black","red","black","black"),2),
         Effect = smith.ts$Effect)
ts.effects <- merge(smith.ts, repl.ts,</pre>
                        by = c("Effect","dv"), suffixes = c("Smith", "Replication"))
ts.plot \leftarrow ggplot(data = ts.effects, aes(x = pesSmith, y = pesReplication, shape = dv)) +
  geom_point(size = 2.5, col = ts.effects$colSmith) +
  xlim(0, 1.00) +
  ylim(0, 1.00) +
  geom_abline(slope = 1, intercept = 0, col = "blue") +
  theme_classic() +
  theme(legend.position = c(0.2, 0.85),
        legend.background = element_rect(colour = "black",
                                         linetype = "solid",
                                         fill = "lightgray"),
        legend.title = element_blank(),
        legend.margin=margin(-3,5,0,0)) +
  labs(y = "Replication", x = "Smith et al.", title = "Task-Switching")
###Exp3 (Visual Search)
smith.vs <- smith_anovas[[5]] %>%
  ungroup() %>%
 bind_rows(smith_anovas[[6]]) %>%
  select(Effect = rowname, pes, LL, UL) %>%
 mutate(dv = rep(c("acc","rt"), each = 3),
         col = rep(c("black", "black", "red"), 2))
repl.vs <- repl_anovas[[5]] %>%
  ungroup() %>%
  bind_rows(repl_anovas[[6]]) %>%
```

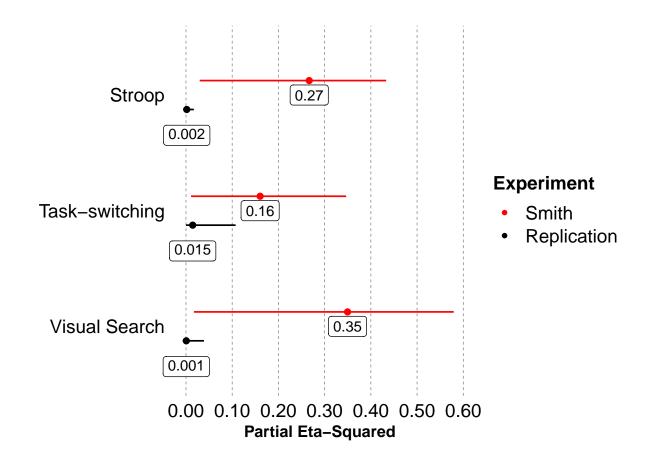
```
select(Effect = rowname, pes, LL, UL) %>%
  mutate(dv = rep(c("acc","rt"), each = 3),
         col = rep(c("black", "black", "red"), 2),
         Effect = smith.vs$Effect)
vs.effects <- merge(smith.vs, repl.vs,</pre>
                        by = c("Effect","dv"), suffixes = c("Smith",
                                                              "Replication"))
vs.plot \leftarrow ggplot(data = vs.effects, aes(x = pesSmith, y = pesReplication, shape = dv)) +
  geom_point(size = 2.5, col = vs.effects$colSmith) +
 xlim(0, 1) +
 ylim(0, 1) +
  geom_abline(slope = 1, intercept = 0, col = "blue") +
  theme_classic() +
  theme(legend.position = c(0.2, 0.85),
        legend.background = element_rect(colour = "black",
                                          linetype = "solid",
                                          fill = "lightgray"),
        legend.title = element_blank(),
        legend.margin=margin(-3,5,0,0)) +
 labs(y = "Replication", x = "Smith et al.", title = "Visual Search")
all.plot <- plot_grid(stroop.plot, ts.plot, vs.plot, ncol = 3)</pre>
title <- ggdraw() +
 draw_label(
    "Effect Size Comparisons",
   fontface = 'bold',
   x = 0,
   hjust = 0
  ) +
 theme(
    # add margin on the left of the drawing canvas,
    # so title is aligned with left edge of first plot
    plot.margin = margin(0, 0, 0, 7)
 )
all.plot <- plot_grid(</pre>
 title, all.plot,
 ncol = 1,
 # rel_heights values control vertical title margins
 rel_heights = c(0.1, 1)
all.plot
```

# **Effect Size Comparisons**



```
ggsave(all.plot,
       file = "results/plots/supp_all_effects_plot.pdf",
       units = "in",
       width = 9.5,
       height = 4.50,
       dpi = 600)
### Forest plot
#Graph comparison of key effects for all three experiments
forest.colors <- c("black", "red")</pre>
forest.data <- data.frame(Experiment = rep(c("Smith", "Replication"), 3),</pre>
                            name = rep(c("Stroop", "Task-switching", "Visual Search"), each = 2),
                            dv = rep(c("rt", "acc", "rt"), each = 2),
                            pes = numeric(6),
                            LL = numeric(6),
                            UL = numeric(6))
forest.data[1,4:6] \leftarrow smith\_anovas[[2]][3,c(6,8,9)]
forest.data[2,4:6] \leftarrow repl_anovas[[2]][3,c(6,8,9)]
forest.data[3,4:6] <- smith_anovas[[3]][5,c(6,8,9)]
forest.data[4,4:6] \leftarrow repl_anovas[[3]][5,c(6,8,9)]
forest.data[5,4:6] \leftarrow smith_anovas[[6]][3,c(6,8,9)]
forest.data[6,4:6] \leftarrow repl_anovas[[6]][3,c(6,8,9)]
```

```
forest.comp <- mod.forestplot(df = forest.data,</pre>
                              estimate = pes,
                              ci.lower = LL,
                              ci.upper = UL,
                              colour = Experiment,
                              xlab = "Partial Eta-Squared"
  scale_color_manual(values = forest.colors) +
  scale_x_continuous(labels = label_number(accuracy = 0.01), breaks = seq(0.00, 0.60, 0.10)) +
  #Too busy w/numbers for effects?
  geom_label(data = subset(forest.data, Experiment == "Smith"),
                           aes(label = round(pes, digits = 2))) +
  geom_label(data = subset(forest.data, Experiment == "Replication"),
                           aes(label = round(pes, digits = 3)),
                          vjust = 2.50) +
   theme(axis.text=element_text(size=14),
      panel.background = element_rect(fill = "white", colour = "white", size = 1),
      axis.title = element_text(size=12, face="bold"),
      strip.text = element_text(size = 12, face = "bold",colour = "black", angle = 0),
      legend.text = element_text(size = 14),
      legend.title = element_text(size= 14, face = "bold")) +
  coord_cartesian(clip="off") #Disable clipping to draw outside plot area
## Warning: Using 'size' aesthetic for lines was deprecated in ggplot2 3.4.0.
## i Please use 'linewidth' instead.
## Scale for colour is already present.
## Adding another scale for colour, which will replace the existing scale.
forest.comp
## Warning: Using the 'size' aesthietic with geom_segment was deprecated in ggplot2 3.4.0.
## i Please use the 'linewidth' aesthetic instead.
```



## [1] 0.6181838 9.0496059 0.1801278

```
#Average proportion is 3.28%
mean(prop.effects)
```

## [1] 3.282639

# Appendix: Analysis with conventional dropping of timeout trials (only affects PE analysis)

## Experiment 1: Stroop

```
e1.conv.PE = plyr::ddply(mergedStroopData.conventional,
                     .(sj, cb, congruency, posture),
                     summarise,
                     meanPE = 100 - (mean(ac)*100))
e1.conv.PE$posture = factor(e1.conv.PE$posture)
e1.conv.PE$congruency = factor(e1.conv.PE$congruency)
e1.conv.PE$sj = factor(e1.conv.PE$sj)
e1.conv.errmodel <- ezANOVA(e1.conv.PE,</pre>
                   dv = .(meanPE),
                   wid=.(sj),
                   within=.(posture,congruency),
                   detailed=TRUE,
                   type=3,
                   return aov = TRUE)
e1.conv.MSE = e1.conv.errmodel$ANOVA$SSd/e1.conv.errmodel$ANOVA$DFd
exp1_ANOVA_PE_con <-
paste(e1.conv.errmodel$ANOVA$Effect,": F(",
      e1.conv.errmodel$ANOVA$DFn,
     e1.conv.errmodel$ANOVA$DFd,
      ") = "
     round(e1.conv.errmodel$ANOVA$F,3),
      ", MSE = ",
     round(e1.conv.MSE,3),
      ", p = ",
      round(e1.conv.errmodel$ANOVA$p,3),
      ", partialEtaSq = ",
     round(e1.conv.errmodel$ANOVA$SSn/(e1.conv.errmodel$ANOVA$SSn+e1.conv.errmodel$ANOVA$SSd),4),
      sep="")
exp1_ANOVA_PE_con
## [1] "(Intercept): F(1, 49) = 54.682, MSE = 69.145, p = 0, partialEtaSq = 0.5274"
## [2] "posture: F(1, 49) = 0.028, MSE = 14.823, p = 0.868, partialEtaSq = 6e-04"
## [3] "congruency: F(2, 98) = 8.813, MSE = 9.641, p = 0, partialEtaSq = 0.1524"
## [4] "posture:congruency: F(2, 98) = 1.533, MSE = 5.865, p = 0.221, partialEtaSq = 0.0303"
write.csv(exp1_ANOVA_PE_con, "results/exp1_ANOVA_PE_con.csv")
#BF for errors
e1.conv.BF = anovaBF(meanPE~congruency*posture+sj,
```

```
data = e1.conv.PE,
                         whichRandom = "sj",
                         method="laplace")
e1.conv.BF
## Bayes factor analysis
## -----
## [1] congruency + sj
                                                     : 136.8983 ±NA%
## [2] posture + sj
                                                     : 0.1188895 ±NA%
## [3] congruency + posture + sj
                                                     : 16.39088 ±NA%
## [4] congruency + posture + congruency:posture + sj : 2.363896 ±NA%
##
## Against denominator:
## meanPE ~ sj
## ---
## Bayes factor type: BFlinearModel, JZS
#...get the Bayes factor for the Null Interaction
e1.conv.BF[3]/e1.conv.BF[4]
## Bayes factor analysis
## -----
## [1] congruency + posture + sj : 6.933838 \pm NA\%
## Against denominator:
## meanPE ~ congruency + posture + congruency:posture + sj
## ---
## Bayes factor type: BFlinearModel, JZS
1/(e1.conv.BF[3]/e1.conv.BF[4])
## Bayes factor analysis
## [1] congruency + posture + congruency:posture + sj : 0.1442203 ±NA%
## Against denominator:
## meanPE ~ congruency + posture + sj
## Bayes factor type: BFlinearModel, JZS
Experiment 2: Stroop
```

## 'switchTrialType'. You can override using the '.groups' argument.

```
e2.conv.PE$posture = factor(e2.conv.PE$posture)
e2.conv.PE$congruentTrialType = factor(e2.conv.PE$congruentTrialType)
e2.conv.PE$participant = factor(e2.conv.PE$participant)
e2.conv.PE$switchTrialType = factor(e2.conv.PE$switchTrialType)
e2.conv.errmodel <- ezANOVA(e2.conv.PE,
                  dv = .(meanPE),
                   wid=.(participant),
                   within=.(posture,congruentTrialType, switchTrialType),
                   detailed=TRUE,
                   type=3,
                   return_aov = TRUE)
## Warning: Collapsing data to cell means. *IF* the requested effects are a subset of the full design,
## you must use the "within_full" argument, else results may be inaccurate.
e2.conv.MSE = e2.conv.errmodel$ANOVA$SSd/e2.conv.errmodel$ANOVA$DFd
exp2_ANOVA_PE_con <-
paste(e2.conv.errmodel$ANOVA$Effect,": F(",
      e2.conv.errmodel$ANOVA$DFn,
     e2.conv.errmodel$ANOVA$DFd,
     ") = ",
     round(e1.conv.errmodel$ANOVA$F,3),
      ", MSE = ",
     round(e2.conv.MSE,3),
      ", p = ",
     round(e2.conv.errmodel$ANOVA$p,3),
      ", partialEtaSq = ",
     round(e2.conv.errmodel$ANOVA$SSn/(e2.conv.errmodel$ANOVA$SSn+e2.conv.errmodel$ANOVA$SSd),4),
      sep="")
exp2_ANOVA_PE_con
## [1] "(Intercept): F(1, 50) = 54.682, MSE = 124.788, p = 0, partialEtaSq = 0.8018"
## [2] "posture: F(1, 50) = 0.028, MSE = 77.648, p = 0.425, partialEtaSq = 0.0128"
## [3] "congruentTrialType: F(1, 50) = 8.813, MSE = 76.694, p = 0, partialEtaSq = 0.6854"
## [4] "switchTrialType: F(1, 50) = 1.533, MSE = 21.856, p = 0, partialEtaSq = 0.6665"
## [5] "posture:congruentTrialType: F(1, 50) = 54.682, MSE = 38.811, p = 0.939, partialEtaSq = 1e-04"
## [6] "posture:switchTrialType: F(1, 50) = 0.028, MSE = 18.608, p = 0.243, partialEtaSq = 0.0272"
## [7] "congruentTrialType:switchTrialType: F(1, 50) = 8.813, MSE = 13.78, p = 0, partialEtaSq = 0.560
## [8] "posture:congruentTrialType:switchTrialType: F(1, 50) = 1.533, MSE = 13.555, p = 0.491, partial
write.csv(exp2_ANOVA_PE_con, "results/exp2_ANOVA_PE_con.csv")
#BF for errors
e2.conv.BF = anovaBF(meanPE~congruentTrialType*posture+participant,
                          data = e2.conv.PE,
```

## Warning: data coerced from tibble to data frame

whichRandom = "participant",

method="laplace")

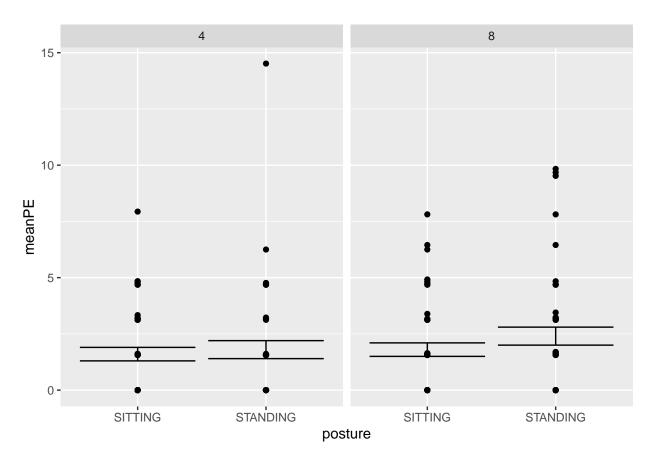
```
e2.conv.BF
## Bayes factor analysis
## -----
## [1] posture + participant
                                                                               : 0.06825202
                                                                                               ±NA%
## [2] congruentTrialType + participant
                                                                                : 7.659849e+119 \pm NA\%
## [3] posture + congruentTrialType + participant
                                                                               : 5.611788e+118 ±NA%
\#\# [4] posture + congruentTrialType + posture:congruentTrialType + participant : 1.187856e+117 \pmNA%
## Against denominator:
   meanPE ~ participant
## ---
## Bayes factor type: BFlinearModel, JZS
#...get the Bayes factor for the Null Interaction
e2.conv.BF[3]/e2.conv.BF[4]
## Bayes factor analysis
## [1] posture + congruentTrialType + participant : 47.24301 ±NA%
## Against denominator:
##
   meanPE ~ posture + congruentTrialType + posture:congruentTrialType + participant
## ---
## Bayes factor type: BFlinearModel, JZS
1/(e2.conv.BF[3]/e2.conv.BF[4])
## Bayes factor analysis
## [1] posture + congruentTrialType + posture:congruentTrialType + participant : 0.02116715 ±NA%
## Against denominator:
   meanPE ~ posture + congruentTrialType + participant
## ---
## Bayes factor type: BFlinearModel, JZS
Experiment 3: Visual Search
e3.conv.PE = plyr::ddply(merged.VS.data.conventional,
                     .(sj, setSize, posture),
                     summarise,
                     meanPE = 100 - (mean(ac)*100))
```

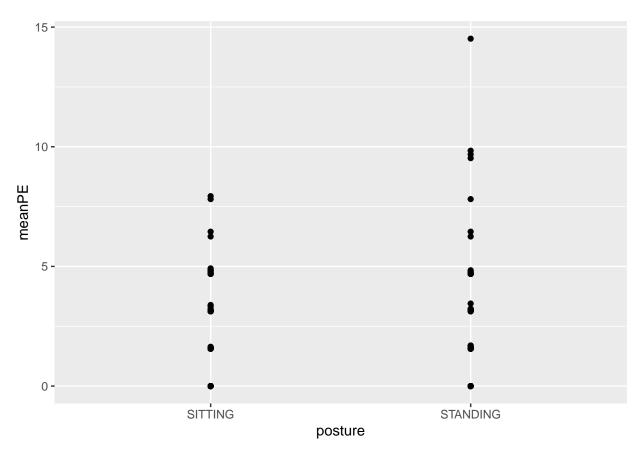
e3.conv.PE\$posture = factor(e3.conv.PE\$posture)
e3.conv.PE\$setSize = factor(e3.conv.PE\$setSize)

e3.conv.PE\$sj = factor(e3.conv.PE\$sj)

e3.conv.errmodel <- ezANOVA(e3.conv.PE,

```
dv = .(meanPE),
                   wid=.(sj),
                   within=.(posture, setSize),
                   detailed=TRUE,
                   type=3,
                   return_aov = TRUE)
e3.conv.MSE = e3.conv.errmodel$ANOVA$SSd/e3.conv.errmodel$ANOVA$DFd
exp3_ANOVA_PE_con <-
paste(e3.conv.errmodel$ANOVA$Effect,": F(",
      e3.conv.errmodel$ANOVA$DFn,
     e3.conv.errmodel$ANOVA$DFd,
     ") = ",
     round(e3.conv.errmodel$ANOVA$F,3),
      ", MSE = ",
     round(e3.conv.MSE,3),
      ", p = ",
     round(e3.conv.errmodel$ANOVA$p,3),
      ", partialEtaSq = ",
     round(e3.conv.errmodel$ANOVA$SSn/(e3.conv.errmodel$ANOVA$SSn+e3.conv.errmodel$ANOVA$SSd),4),
      sep="")
exp3_ANOVA_PE_con
## [1] "(Intercept): F(1, 49) = 55.236, MSE = 12.588, p = 0, partialEtaSq = 0.5299"
## [2] "posture: F(1, 49) = 3.311, MSE = 2.43, p = 0.075, partialEtaSq = 0.0633"
## [3] "setSize: F(1, 49) = 2.312, MSE = 3.628, p = 0.135, partialEtaSq = 0.0451"
## [4] "posture:setSize: F(1, 49) = 0.623, MSE = 2.794, p = 0.434, partialEtaSq = 0.0125"
write.csv(exp3_ANOVA_PE_con, "results/exp3_ANOVA_PE_con.csv")
graphPE3.con = describeBy(e3.conv.PE$meanPE,
                     list(e3.conv.PE$posture, e3.conv.PE$setSize),
                     mat=TRUE,
                     digits = 1)
graphPE3.con = graphPE3.con[,c("group1", "group2", "mean", "sd", "se")]
names(graphPE3.con) = c("posture", "setSize", "meanPE", "sd", "se")
e3.PE.conv.plot <-
ggplot(data = e3.conv.PE, aes(x = posture,
                             y = meanPE)) +
  geom_point() +
  facet_wrap(~setSize) +
  geom_errorbar(data = graphPE3.con, aes(ymin = meanPE - se, ymax = meanPE + se))
e3.PE.conv.plot
```





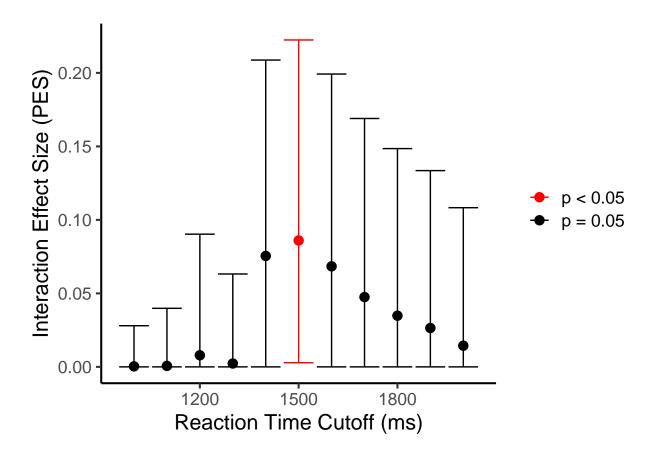
```
#hist(e3.conv.PE$meanPE)
#BF for errors
e3.conv.BF = anovaBF(meanPE~setSize*posture+sj,
                         data = e3.conv.PE,
                         whichRandom = "sj",
                         method="laplace")
e3.conv.BF
## Bayes factor analysis
## [1] setSize + sj
                                               : 0.5212586 ±NA%
## [2] posture + sj
                                               : 0.4937759 ±NA%
## [3] setSize + posture + sj
                                              : 0.2659863 ±NA%
## [4] setSize + posture + setSize:posture + sj : 0.06740099 \pm NA\%
## Against denominator:
   meanPE ~ sj
## ---
## Bayes factor type: BFlinearModel, JZS
\#... get the Bayes factor for the Null Interaction
e3.conv.BF[3]/e3.conv.BF[4]
```

## Bayes factor analysis

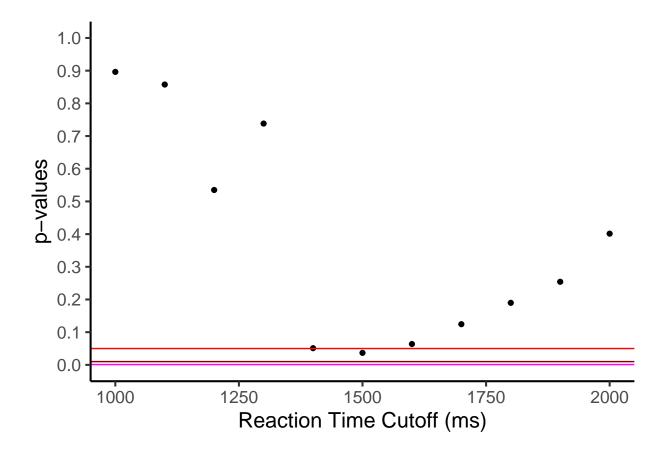
```
## [1] setSize + posture + sj : 3.946327 \pm NA\%
## Against denominator:
## meanPE ~ setSize + posture + setSize:posture + sj
## ---
## Bayes factor type: BFlinearModel, JZS
1/(e3.conv.BF[3]/e3.conv.BF[4])
## Bayes factor analysis
## -----
## [1] setSize + posture + setSize:posture + sj : 0.2534002 ±NA%
## Against denominator:
## meanPE ~ setSize + posture + sj
## ---
## Bayes factor type: BFlinearModel, JZS
Effect of different RT cutoffs on PE interaction
#compare different criteria for removing long trials
criteria \leftarrow seq(1000, 2000, by = 100)
ncriteria <- length(criteria)</pre>
criteria.dat <- merged.VS.data.orig</pre>
for (i in criteria){
  criteria.dat <- criteria.dat %>%
    mutate("drop{i}" := ifelse(rt >= i | ac == 0, 0, 1))
}
vsPEcriteria = criteria.dat %>%
  group_by(sj, cb, setSize, posture) %>%
  summarize(across(starts_with("drop"), ~ 100 - (mean(.x)*100)))
## 'summarise()' has grouped output by 'sj', 'cb', 'setSize'. You can override using the '.groups'
## argument.
#...set as factors
vsPEcriteria$sj = factor(vsPEcriteria$sj)
vsPEcriteria$cb = factor(vsPEcriteria$cb)
vsPEcriteria$setSize = factor(vsPEcriteria$setSize)
vsPEcriteria$postureFactor = factor(vsPEcriteria$posture)
criteria.results <- data.frame(cutoff = criteria,</pre>
                               es = numeric(ncriteria),
                                p.vals = numeric(ncriteria),
                                ci.LL = numeric(ncriteria),
                               ci.UL = numeric(ncriteria))
for (i in 1:length(criteria)){
```

## -----

```
tempcol <- paste0("drop",criteria[i])</pre>
  tempmod <- aov_ez(data = vsPEcriteria,</pre>
                                dv = tempcol,
                                id = 'sj',
                                within = c('posture', 'setSize'),
                                anova_table = list(es = "pes", correction = "none"),
                                type = 3)
  criteria.results$es[i] <- tempmod$anova_table$pes[3]</pre>
  criteria.results$p.vals[i] <- tempmod$anova_table$`Pr(>F)`[3]
  criteria.results$ci.LL[i] <- get.ci.partial.eta.squared(F.value=tempmod$anova_table$F[3],
                                                     df1=tempmod$anova_table$`num Df`[3],
                                                     df2 = tempmod$anova_table$`den Df`[3])$LL
  criteria.results$ci.UL[i] <- get.ci.partial.eta.squared(F.value=tempmod$anova_table$F[3],</pre>
                                                     df1=tempmod$anova_table$`num Df`[3],
                                                     df2 = tempmod$anova_table$`den Df`[3])$UL
}
criteria.results$sig <- ifelse(criteria.results$p.vals < 0.05, "p < 0.05", "p \u2265 0.05")
exp3_rt_cutoff <-
ggplot(criteria.results, aes(criteria, es, col = sig)) +
  geom_point(size = 3) +
  geom_errorbar(aes(ymin = ci.LL, ymax = ci.UL)) +
  scale_color_manual(values = c("red","black")) +
  theme_classic(base_size = 16) +
  theme(legend.title = element_blank()) +
  labs(y = "Interaction Effect Size (PES)", x = "Reaction Time Cutoff (ms)")
exp3_rt_cutoff
```



```
ggsave(exp3_rt_cutoff,
       file = "results/plots/supp_exp3_RT_cutoff.pdf",
       units = "in",
       width = 9,
       height = 6,
       dpi = 600)
#Not in the supp: Distribution of p-values
exp3_rt_cutoff.pvals <-</pre>
ggplot(criteria.results, aes(criteria, p.vals)) +
  geom_point(size = 1.5) +
  scale_color_manual(values = c("red","black")) +
  theme_classic(base_size = 16) +
  theme(legend.title = element_blank()) +
  scale_y_continuous(limits = c(0, 1), breaks = seq(0, 1, 0.1)) +
  labs(y = "p-values", x = "Reaction Time Cutoff (ms)")
exp3_rt_cutoff.pvals + geom_hline(yintercept = 0.05, color = "red") +
                       geom_hline(yintercept = 0.01, color = "darkred") +
                       geom_hline(yintercept = 0.001, color = "magenta")
```



#plot(criteria.results\$p.vals, criteria.results\$es)