

# lme\_mods

Jacob Salminen

2024-04-19

## Packages & Setup

```
# install.packages(c("tidyverse", "purrr", "R.matlab", "readxl", "dplyr"))
library(readxl);
library(purrr)
library(tidyverse);
```

```
## -- Attaching core tidyverse packages ----- tidyverse 2.0.0 --
## v dplyr      1.1.4      v readr      2.1.5
## v forcats    1.0.0      v stringr   1.5.1
## v ggplot2    3.5.0      v tibble    3.2.1
## v lubridate  1.9.3      v tidyr     1.3.1
## -- Conflicts ----- tidyverse_conflicts() --
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()     masks stats::lag()
## i Use the conflicted package (<http://conflicted.r-lib.org/>) to force all conflicts to become errors
```

```
library(tibble)
library(knitr);
library(gtsummary)
library(kableExtra)
```

```
##
## Attaching package: 'kableExtra'
##
## The following object is masked from 'package:dplyr':
##
##     group_rows
```

```
library(lme4)
```

```
## Loading required package: Matrix
##
## Attaching package: 'Matrix'
##
## The following objects are masked from 'package:tidyr':
##
##     expand, pack, unpack
```

## GTSUMMARY THEME

```
# my_theme <-  
#   list(  
#     "tbl_summary-str:default_con_type" = "continuous2",  
#     "tbl_summary-str:continuous_stat" = c(  
#       "{median} ({p25} - {p75})",  
#       "{mean} ({sd})",  
#       "{min} - {max}"  
#     ),  
#     "tbl_summary-str:categorical_stat" = "{n} / {N} ({p}%)",  
#     "style_number-arg:big.mark" = "",  
#     "tbl_summary-fn:percent_fun" = function(x) style_percent(x, digits = 3)  
#   )  
# my_theme <-  
#   list()  
# gtsummary::set_gtsummary_theme(my_theme)  
gtsummary::set_gtsummary_theme(theme_gtsummary_journal("jama"))
```

```
## Setting theme 'JAMA'  
## Setting theme 'JAMA'
```

```
# reset_gtsummary_theme()
```

## load table

```
# excel_dir <- "M:/jsalminen/GitHub/par_EEGProcessing/src/_data/MIM_dataset/_studies/04162024_MIM_YAOAN8"  
excel_dir <- "M:/jsalminen/GitHub/par_EEGProcessing/src/_data/MIM_dataset/_studies/04232024_MIM_YAOAN89."  
eegt <- read_excel(excel_dir, sheet="Sheet1")
```

## get unique entries

```
clusters = unique(eegt$cluster_id);  
subjects = unique(eegt$subj_char);  
groups = unique(eegt$group_char);  
kin_measures = c('mean_APexc_COV', 'mean_APexc_mean', 'mean_MLexc_COV', 'mean_MLexc_mean', 'mean_StepDur', 'mean_StepDur_offset', 'mean_theta_avg_power', 'mean_alpha_avg_power', 'mean_beta_avg_power', 'mean_aperiodic_exp', 'mean_aperiodic_offset');  
eegt_measures = c('theta_avg_power', 'alpha_avg_power', 'beta_avg_power', 'aperiodic_exp', 'aperiodic_offset');
```

## get speeds only

```
eegt <- filter_at(eegt, vars('cond_char'), any_vars(. %in% c('0.25', '0.5', '0.75', '1.0')))  
flat_speeds = unique(eegt$cond_char)  
eegt$cond_char <- as.numeric(eegt$cond_char)  
eegt$speed_cond_num <- as.numeric(eegt$cond_char)  
eegt <- mutate(eegt, across(c('subj_char'), factor))
```

Cluster	H1004_COV				H1007_COV				H1009_COV				H1010_COV				H1011_COV				H1012_COV				H1013_COV				H1014_COV				H1015_COV				H1016_COV				H1017_COV				H1018_COV				H1019_COV				H1020_COV				H1021_COV				H1022_COV				H1023_COV				H1024_COV				H1025_COV				H1026_COV				H1027_COV				H1028_COV				H1029_COV				H1030_COV				H1031_COV				H1032_COV				H1033_COV				H1034_COV				H1035_COV				H1036_COV				H1037_COV				H1038_COV				H1039_COV				H1040_COV				H1041_COV				H1042_COV				H1043_COV				H1044_COV				H1045_COV				H1046_COV				H1047_COV				H1048_COV				H1049_COV				H1050_COV				H1051_COV				H1052_COV				H1053_COV				H1054_COV				H1055_COV				H1056_COV				H1057_COV				H1058_COV				H1059_COV				H1060_COV				H1061_COV				H1062_COV				H1063_COV				H1064_COV				H1065_COV				H1066_COV				H1067_COV				H1068_COV				H1069_COV				H1070_COV				H1071_COV				H1072_COV				H1073_COV				H1074_COV				H1075_COV				H1076_COV				H1077_COV				H1078_COV				H1079_COV				H1080_COV				H1081_COV				H1082_COV				H1083_COV				H1084_COV				H1085_COV				H1086_COV				H1087_COV				H1088_COV				H1089_COV				H1090_COV				H1091_COV				H1092_COV				H1093_COV				H1094_COV				H1095_COV				H1096_COV				H1097_COV				H1098_COV				H1099_COV				H1100_COV				H1101_COV				H1102_COV				H1103_COV				H1104_COV				H1105_COV				H1106_COV				H1107_COV				H1108_COV				H1109_COV				H1110_COV				H1111_COV				H1112_COV				H1113_COV				H1114_COV				H1115_COV				H1116_COV				H1117_COV				H1118_COV				H1119_COV				H1120_COV				H1121_COV				H1122_COV				H1123_COV				H1124_COV				H1125_COV				H1126_COV				H1127_COV				H1128_COV				H1129_COV				H1130_COV				H1131_COV				H1132_COV				H1133_COV				H1134_COV				H1135_COV				H1136_COV				H1137_COV				H1138_COV				H1139_COV				H1140_COV				H1141_COV				H1142_COV				H1143_COV				H1144_COV				H1145_COV				H1146_COV				H1147_COV				H1148_COV				H1149_COV				H1150_COV				H1151_COV				H1152_COV				H1153_COV				H1154_COV				H1155_COV				H1156_COV				H1157_COV				H1158_COV				H1159_COV				H1160_COV				H1161_COV				H1162_COV				H1163_COV				H1164_COV				H1165_COV				H1166_COV				H1167_COV				H1168_COV				H1169_COV				H1170_COV				H1171_COV				H1172_COV				H1173_COV				H1174_COV				H1175_COV				H1176_COV				H1177_COV				H1178_COV				H1179_COV				H1180_COV				H1181_COV				H1182_COV				H1183_COV				H1184_COV				H1185_COV				H1186_COV				H1187_COV				H1188_COV				H1189_COV				H1190_COV				H1191_COV				H1192_COV				H1193_COV				H1194_COV				H1195_COV				H1196_COV				H1197_COV				H1198_COV				H1199_COV				H1200_COV				H1201_COV				H1202_COV				H1203_COV				H1204_COV				H1205_COV				H1206_COV				H1207_COV				H1208_COV				H1209_COV				H1210_COV				H1211_COV				H1212_COV				H1213_COV				H1214_COV				H1215_COV				H1216_COV				H1217_COV				H1218_COV				H1219_COV				H1220_COV				H1221_COV				H1222_COV				H1223_COV				H1224_COV				H1225_COV				H1226_COV				H1227_COV				H1228_COV				H1229_COV				H1230_COV				H1231_COV				H1232_COV				H1233_COV				H1234_COV				H1235_COV				H1236_COV				H1237_COV				H1238_COV				H1239_COV				H1240_COV				H1241_COV				H1242_COV				H1243_COV				H1244_COV				H1245_COV				H1246_COV				H1247_COV				H1248_COV				H1249_COV				H1250_COV				H1251_COV				H1252_COV				H1253_COV				H1254_COV				H1255_COV				H1256_COV				H1257_COV				H1258_COV				H1259_COV				H1260_COV				H1261_COV				H1262_COV				H1263_COV				H1264_COV				H1265_COV				H1266_COV				H1267_COV				H1268_COV				H1269_COV				H1270_COV				H1271_COV				H1272_COV				H1273_COV				H1274_COV				H1275_COV			
	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value	Beta (95% CrI)	p-value	q-value	h-value</																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																				

<sup>2</sup> False discovery rate correction for multiple testing

<sup>2</sup> False discovery rate correction for multiple testing<sup>2</sup> False discovery rate correction for multiple testing<sup>2</sup> False discovery rate correction for multiple testing<sup>2</sup> False discovery rate correction for multiple testing<sup>2</sup> False discovery rate correction for multiple testing<sup>2</sup> False discovery rate correction for multiple testing<sup>2</sup> False discovery rate correction for multiple testing