## Steps to Run a Complete FDA with a Continuous Predictor

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Below we will define the steps required to complete an FDA with our data

- 1. Load the Actigraphy package:
- > library(Actigraphy)
- 2. Read in the activity and covariate data and store them as the datasets covariate and activity:
- data(clinic\_29pt\_ahi)
- > data(act\_29pt)
- > covariate <- clinic\_29pt\_ahi</pre>
- > activity <- act\_29pt
  - 3. Match the data in the activity and covariate datasets:
- > matchid <- fda.matchid(activity, covariate, "contin")</pre>
  - 4. Smooth the data and plot it:
  - L is the number of entries in the activity file
- > L <- nrow(activity)
- > FDinterest <- fda.smoothdata(matchid)</pre>
- > ts.plot(predict(FDinterest\$fd\$fd, 1:L), main="Smoothed Activity Data")
  - 5. Apply FLM to the data with the flm\_cate function:
- > geftFDcont <- flm\_cate(FDinterest)</pre>
  - 6. Set up the plot parameters for the next step:
  - xlim is a vector of length 2 that contains the X-axis boundries based on the time span of the activity data
  - ylim is a vector of length 2 that contains the Y-axis boundries based on the results on the flm\_cate function
  - lb is a vector of labels for the X-axis
  - xat is a vector of positions for the labels (lb) on the X-axis
  - legendx is the x-axis position of the left edge of the legend box

• legendy is the y-axis position of the upper edge of the legend box

7. Plot results from FLM results and F-test if indicated (with the continuous flm plotting function):

```
cont.flm.results <- cont_flm_plot(FDinterest, matchid, geftFDcont, xlim, ylim, T
```

## Notes:

- Columns in Activity file data MUST represent subjects
- Covariate file MUST only contain 2 columns; subject identifier (id) and one covariate (numeric or factor). If user is interested in 2 or more covariates, they should be put into separate datasets and analyzed separately.
- Subject identifier should be the same in both Activity (first column) and Covariate (row names) files