LIMO toolbox: LInear Modeling of EEG data.

List files saved on the disk and their dimensions

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From 1st level analysis using mass univariate approach

In the subject analysis folder, the data, parameters and results are saved

<u>Yr.mat</u>: the EEG data from the .set reorganized to fit X, that is grouped by conditions if Cat $\sim =0$; dimension [electrodes x time frames x trials]

<u>Yhat.mat</u>: the predicted data; dimension [electrodes x time frames x trials]

<u>Beta.mat:</u> the beta values (parameter estimates); dimension [electrodes x time frames x number of parameters in the model (columns of X)]

Res.mat: the residuals (non modelled) data; dimension [electrodes x time frames x trials] R2.mat: the model fit statistic, i.e. percentage of variance explained; dimension [electrodes x time frames x $R^2/F/p$ values]

<u>Condition effect X:</u> refers to a factor effect in categorical designs; dimension [electrodes x time frames x F/p values).

<u>Interaction_effect_X:</u> refers to an interaction between factors, dimension [electrodes x time frames x F/p values).

<u>Covariate_effect_X</u>: refers to the effect of a continuous regressor, dimension [electrodes x time frames x F/p values).

<u>semi_partial_coef_X.mat:_refers</u> to the semi-partial coefficient of a factor (defined by LIMO.design.nb_conditions) or a covariate (defined in LIMO.design.nb_continuous), dimension [electrodes x frames x R2/F/p values].

<u>con_X.mat</u>: refers to a t contrast, dimension [electrodes x frames x cB/standard error/df/t/p values].

<u>ess X.mat</u>: refers to a F contrast, dimension [electrodes x frames x cB/standard error/df/t/p values].

--- adding bootstrap, in the H0 subfolder

<u>boot_table</u>: the resampling table used, dimension [number of trials x number of bootstraps] <u>HO_Betas</u>: dimension [electrodes x time frames x number of parameters in the model

(columns of X) x number of bootstraps]

<u>H0_R2</u>: dimension: [electrodes x time frames x R²/F/p values x number of bootstraps]

<u>H0_Condition_effect_X</u>: dimension [electrodes x time frames x F/p values x number of bootstraps]

<u>HO Interaction effect X</u>: dimension [electrodes x time frames x F/p values x number of bootstraps]

<u>H0_Covariate_effect_X</u>: dimension [electrodes x time frames x F/p values x number of bootstraps]

<u>H0_semi_partial_coef_X.mat:</u> dimension [electrodes x frames x R2/F/p values x number of bootstraps].

<u>H0 con X.mat</u>: dimension [electrodes x frames x cB/t/p values x number of bootstraps]. <u>H0 ess X.mat</u>: dimension [electrodes x frames x cB/F/p values x number of bootstraps].

--- adding tfce, in the TFCE subfolder

<u>tfce_R2</u>: <u>dimension</u>: <u>dimension</u> [electrodes x time frames]

tfce_Condition_effect_X: dimension [electrodes x time frames]

tfce_Interaction_effect_X: dimension [electrodes x time frames]

<u>tfce_Covariate_effect_X</u>: dimension [electrodes x time frames]

tfce semi_partial_coef_X.mat: dimension [electrodes x frames].

tfce_con_X.mat: dimension [electrodes x frames].

<u>tfce_ess_X.mat</u>: dimension [electrodes x frames].

In the H0 subfolder, the tfce score maps under H0

tfce_H0_R2: dimension [electrodes x time frames x number of bootstraps]

<u>tfce_H0_Condition_effect_X</u>: dimension [electrodes x time frames x number of bootstraps]

tfce H0 Interaction effect X: dimension [electrodes x time frames x number of bootstraps]

<u>tfce_H0_Covariate_effect_X</u>: dimension [electrodes x time frames x number of bootstraps]

tfce_H0_semi_partial_coef_X.mat: dimension [electrodes x frames x number of bootstraps].

tfce_H0_con_X.mat: dimension [electrodes x frames x number of bootstraps].

tfce_H0_ess_X.mat: dimension [electrodes x frames x number of bootstraps].

From 2nd level analysis using mass univariate approach

One sample t-test

<u>one_sample_ttest_parameter_X.mat</u>: returns the trimmed mean parameter values and associated statistics (dimensions electrodes x frames x 5). The last dimension codes mean values, standard error, degrees of freedom, t and p.

<u>HO one sample ttest parameter X.mat</u>: this file constrains the T and p values obtained under HO for each bootstrap (dimensions electrodes x frames x 2 x nboot)

<u>tfce_one_sample_ttest_parameter_X.mat</u>: tfce scores of the t-test (dimensions electrodes x frames)

<u>H0 tfce one sample ttest parameter X.mat</u>: tfce scores of the –test under H0 (dimensions electrodes x frames x nboot)