

Generating datatables from a directory tree

gen_group_command.py -command datatable ...

These details can be found from : **gen_group_command.py -help**

- or see: **gen_group_command.py -hweb**
- example section F : dataTable creation

Assume this study has 10 subjects, each with 2x2x3 betas, and the betas are from a design of (column headers and possible values):

- visit : **before** **after**
- color : **red** **green**
- task : **T1** **T2** **T3**

Enter the sample working directory and view the sample commands:

```
cd AFNI_data6/GroupAna_cases/gen_datatables
afni_open -e ggc_dt_commands.tcsh
tcsh ggc_dt_commands.tcsh
```

Example 1. each subject has one dataset with 12 volumes
(the tcsh script generates such a data tree)

```
gen_group_command.py          \
  -command datatable          \
  -dsets all_results/sub*.nii.gz \
  -dt_factor_list visit before after \
  -dt_factor_list color red green \
  -dt_factor_list task T1 T2 T3 \
  -subs_betas B_R_T1 B_R_T2 B_R_T3 \
               B_G_T1 B_G_T2 B_G_T3 \
               A_R_T1 A_R_T2 A_R_T3 \
               A_G_T1 A_G_T2 A_G_T3 \
  -write_script dt_1.txt
```

Example 2. like 1, but each volume is in a separate dataset
(the tcsh script generates such a data tree)

```
gen_group_command.py \
  -command datatable \
  -dt_factor_list visit before after \
  -dt_factor_list color red green \
  -dt_factor_list task T1 T2 T3 \
  -dsets all_results/data.B_R_T1/sub*.gz \
  -dsets all_results/data.B_R_T2/sub*.gz \
  -dsets all_results/data.B_R_T3/sub*.gz \
  -dsets all_results/data.B_G_T1/sub*.gz \
  -dsets all_results/data.B_G_T2/sub*.gz \
  -dsets all_results/data.B_G_T3/sub*.gz \
  -dsets all_results/data.A_R_T1/sub*.gz \
  -dsets all_results/data.A_R_T2/sub*.gz \
  -dsets all_results/data.A_R_T3/sub*.gz \
  -dsets all_results/data.A_G_T1/sub*.gz \
  -dsets all_results/data.A_G_T2/sub*.gz \
  -dsets all_results/data.A_G_T3/sub*.gz \
  -write_script dt_2.txt
```

Example 3. like 2, but include a table with subject attributes
(the tcsh script generates such a table)

- goal: combine an existing table with lines for the data volumes
- each subject line is duplicated for every data volume line
- the tcsh script generates **subject_attrs.tsv**

Subj	Group	ValA	ValB
sub-0044	G_0044	VA_0044	VB_0044
sub-0046	G_0046	VA_0046	VB_0046
sub-0049	G_0049	VA_0049	VB_0049
sub-0053	G_0053	VA_0053	VB_0053
sub-0060	G_0060	VA_0060	VB_0060
sub-0061	G_0061	VA_0061	VB_0061
sub-0064	G_0064	VA_0064	VB_0064
sub-0073	G_0073	VA_0073	VB_0073
sub-0075	G_0075	VA_0075	VB_0075
sub-0076	G_0076	VA_0076	VB_0076

For example, imagine:

Group : horse, rhino

ValA : age

ValB : grumpiness rating (beware correlation!)

Example 3. like 2, but include the exiting table via **-dt_tsv**

```
gen_group_command.py \
  -command datatable \
  -dt_tsv subject_attrs.tsv \
  -dt_factor_list visit before after \
  -dt_factor_list color red green \
  -dt_factor_list task T1 T2 T3 \
  -dsets all_results/data.B_R_T1/sub*.gz \
  -dsets all_results/data.B_R_T2/sub*.gz \
  -dsets all_results/data.B_R_T3/sub*.gz \
  -dsets all_results/data.B_G_T1/sub*.gz \
  -dsets all_results/data.B_G_T2/sub*.gz \
  -dsets all_results/data.B_G_T3/sub*.gz \
  -dsets all_results/data.A_R_T1/sub*.gz \
  -dsets all_results/data.A_R_T2/sub*.gz \
  -dsets all_results/data.A_R_T3/sub*.gz \
  -dsets all_results/data.A_G_T1/sub*.gz \
  -dsets all_results/data.A_G_T2/sub*.gz \
  -dsets all_results/data.A_G_T3/sub*.gz \
  -write_script dt_3.txt
```