

# Manual for the PyMVPA wrapper tool

## Command:

```
fmriCmd.py dataset <filename>
    samples <filename>
    mask <filename>
    classifier <name> <options>
    [ detrend [ perchunk <value> model <name> polyord <value> ] ]
    [ zscore [ perchunk <value> pvoxel <value> targetdtype <type> ] ]
    [ select <list> ]
    [ crossvalidate { oddeven | {nfold <foldvalue> } | custom <list> } ]
    [ searchlight <radius> ]
    [ accuracy ]
    [ save <filename> ]
```

## Details:

This is a tool which takes input options for running PyMVPA classifiers

### Parameters:

#####

dataset <filename> - mandatory parameter  
- option to specify the dataset file (string)  
- <filename> is the NifTI filename with .nii.gz extension

#####

samples <filename> - mandatory parameter  
- option to specify the attributes file (string)  
- <filename> is the text filename with .txt extension

#####

mask <filename> - mandatory parameter  
- option to specify the mask file (string)  
- <filename> is the NifTI filename with .nii.gz extension (or .img file)

#####

classifier <name> <options> - mandatory parameter  
- option to specify the classifier (string)

- <name> is the classifier name. It takes the following values and options.

Note: Options are specified per classifier. For more details on the options, refer to the PyMVPA manuals (<http://pymvpa.org/modref.html#classifiers-and-errors>)

<options> takes the following parameters as input:

smlr lm <value>

lm <value> - optional parameter

- lambda penatly value (float)
- default: 0.1

knn k <value> voting <method>

<value> - optional parameter

- k value (int)
- default: 2

voting <method> - optional parameter

- voting method (string)
- default: weighted

svm kernel <kernel\_type>

kernel <kernel\_type> - optional parameter

- type of the kernel (string)
- default: linear

blr sigma\_noise <value>

sigma\_noise <value> - optional parameter

- sigma noise (float)
- default: 1.0

enet lm <value>

lm <value> - optional parameter

- lambda value (float)
- default: 1.0

glmnet

LinearCSVMC - SVM implementation with linear kernel

gpr kernel <kernel\_type>

<kernel\_type> - optional parameter

- kernel type (kernel object)
- default: None

lars model\_type <model>  
model\_type <model> - optional parameter  
- model type (string)  
- default: lasso

plr lm <value>  
lm <value> - optional parameter  
- lambda value (int)  
- default: 1

ridgereg lm <value>  
lm <value> - optional parameter  
- lambda value (float)  
- default: None

#####

detrend [ perchunk <value> model <name> polyord <value> ] ] - optional parameter  
(enabled by default)  
perchunk <value> - optional parameter  
- specifies whether data has to be handled in chunks (bool)  
- default: True

model <name> - optional parameter  
- name of the model (string)  
- default: linear

polyord <value> - optional parameter  
- order of the polynomial to remove from the data (int)  
- default: 3

#####

zscore [ perchunk <value> pervoxel <value> targetdtype <type> ] ] - optional parameter  
(enabled by default)

perchunk <value> - optional parameter  
- If perchunk is True samples within the same chunk are z-scored  
independent of samples from other chunks (bool)  
- default: True

pervoxel <value> - optional parameter  
- per voxel value (bool)  
- default: True

targetdtype <type> - optional parameter  
- specifies the target data type (string)  
- default: float64

#####

select <list> - optional parameter (disabled by default)

<list> - mandatory parameter  
- specify the labels to be selected for training in the form of a Python list  
- if select option is not specified, all labels will be selected by default

#####

crossvalidate { oddeven | {nfold <foldvalue> } | custom <list>} - optional parameter  
(disabled by default)

oddeven - optional parameter  
- odd even splitter (string)

nfold <foldvalue> - optional parameter  
- specifies the number of folds in Nfold splitter (int)  
- default: 1

custom <list> - optional parameter  
- specifies the custom splitting of the data set  
<list> - mandatory parameter to the option "custom"  
- specifies the list of runs/chunks to be selected for splitting

#####

searchlight <radius> - optional parameter (disabled by default)

<radius> - optional parameter  
- specifies the searchlight radius (float)  
- default: 4.0

#####

accuracy - optional parameter (disabled by default)  
- by default all the results return error value. In order to compute accuracy value, enable this option

save <filename> - optional parameter  
- filename to save the result (string)

- default: a file with extension .nii.gz is generated by appending current date and time to the string "file". eg: file200911162120.nii.gz

#####

Example:

```
python fmriCmd.py dataset data/bold.nii.gz samples data/attributes.txt mask data/  
mask.nii.gz classifier smlr select [1,2] crossvalidate oddeven
```