

## Software resources

Edinburgh  
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# PRoNTTo

- Software
- Conclusion

## Existing stuff

### Princeton Multi-Voxel Pattern Analysis (MVPA) Toolbox

(<http://code.google.com/p/princeton-mvpa-toolbox/>)

- (+) Matlab based, AFNI/BV oriented but some SPM5 compatibility,
- (-) only(?) neural network classification, only(?) fMRI data, last version from 2009.

### Multivariate Pattern Analysis with Python – PyMVPA

(<http://www.pymvpa.org/>)

- (+) looks powerful with loads of tools,
- (-) Python based, OS specific (Linux).

### Brain Voyager MVPA – BV-MVPA (<http://www.brainvoyager.com/bvqx/>)

- (+) search-light and SVM,
- (-) proprietary code, fMRI only, no way of adding own machine

## OUR GOAL

*“develop a toolbox based on machine learning techniques for the analysis of neuroimaging data”*

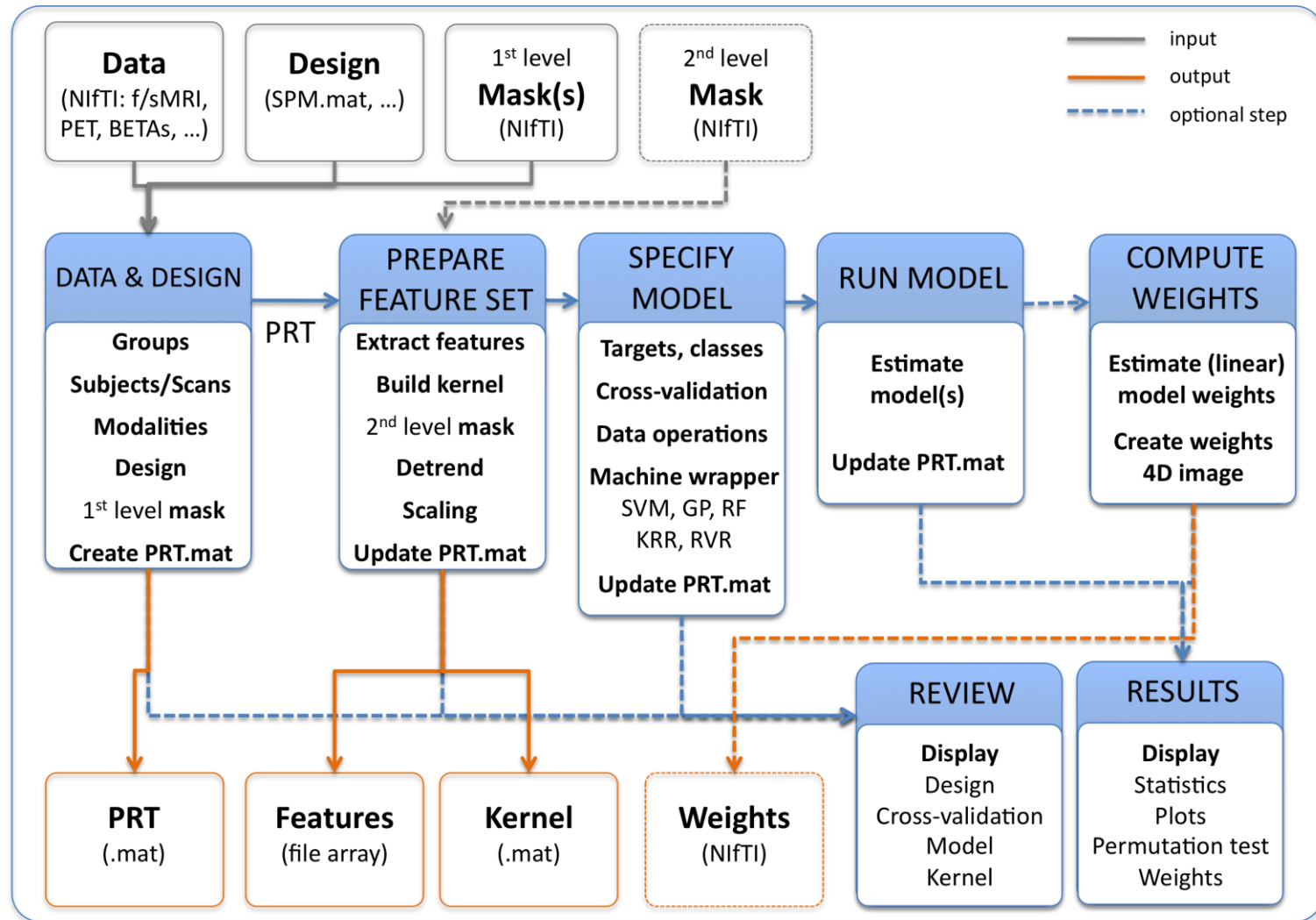
BUT

Matlab based, easy to use,  
multiple modalities (fMRI/sMRI/PET),  
various machines, modular code

**➔ PRoNTTo**

# PRoNTo

## PRoNTo FRAMEWORK



## Requirements

- A recent version of Matlab
- SPM8: Wellcome Trust Centre for Neuroimaging, University College London, UK.

<http://www.fil.ion.ucl.ac.uk/spm/software/spm8/>

Some code in C/C++

Available for: Windows XP (32 bits), Windows 7 (64 bits), Mac OS 10, *some* Linux flavours (32/64 bits).

If more “exotic”  $\Rightarrow$  compile for your OS !!!

## For users

### User Interface

Easy to use  
Click buttons

### Matlab Batch

Very efficient  
Can be saved  
and copied  
Compatible  
with SPM

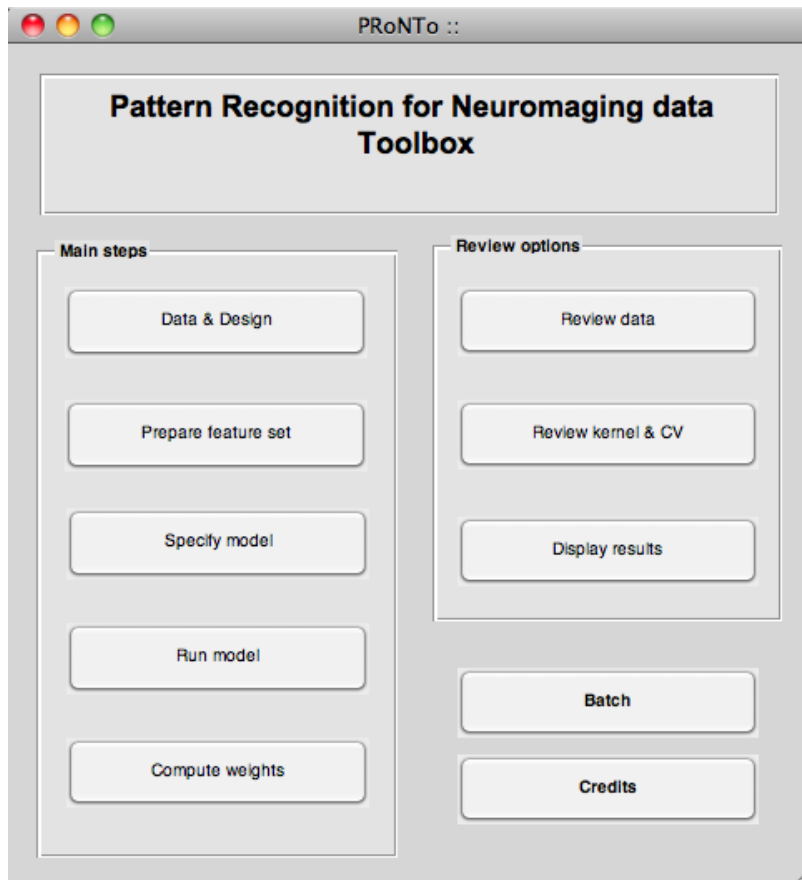
### Functions

User-specific  
analysis  
More  
programming  
skills required

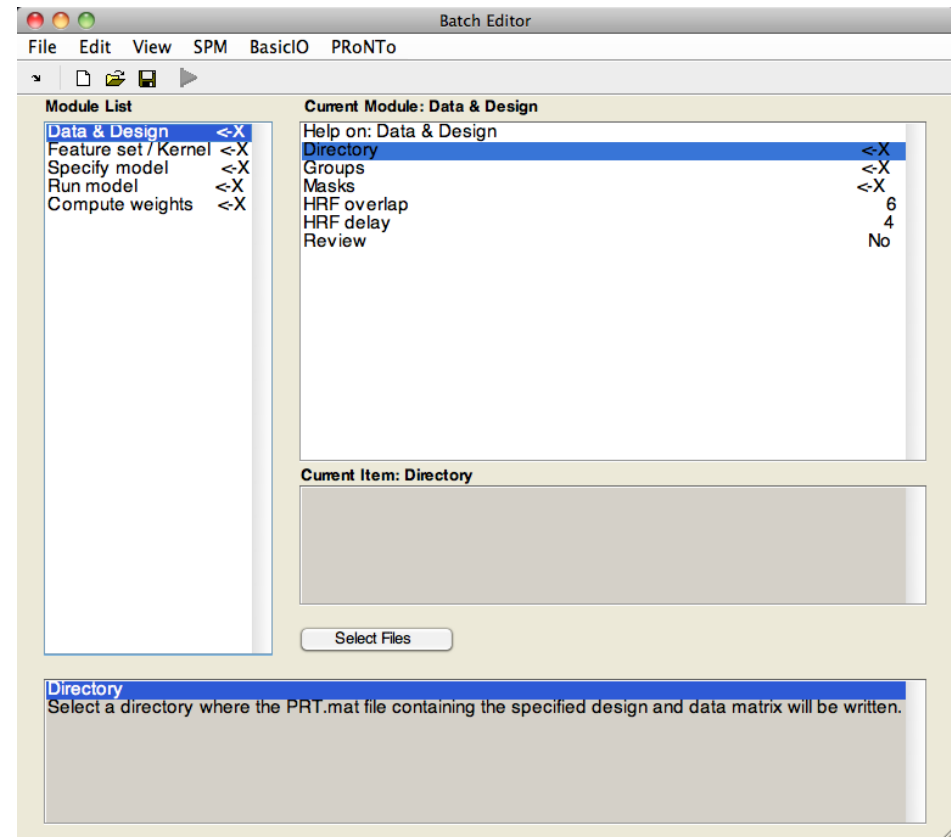
# PRoNTTo

## User point of view

### User Interface



### Matlab Batch





# PRoNTTo

For developers

## User Interface

Specific GUI  
Batch system  
Script

## Machine learning

Features  
Kernel  
Model  
Training  
Validation

Wrapper

## Machines

Classification  
(SVM, GPC, RF)  
Regression  
(KRR, RVR)

## Machine Learning tools, classification

- Support Vector Machines (SVM)

LibSVM implementation,

<http://www.csie.ntu.edu.tw/~cjlin/libsvm>

Nello Cristianini and John Shawe-Taylor. An introduction to support Vector Machines: and other kernel-based learning methods. Cambridge University Press, New York, NY, USA, 2000.

- Gaussian Process Classifier (GPC)

GPLML implementation,

<http://www.gaussianprocess.org/gpml/code/matlab/doc/>

Carl Edward Rasmussen and Christopher K. I. Williams. Gaussian Processes for Machine Learning. Adaptive Computation and Machine Learning. the MIT Press, 2006.

## Machine Learning tools, regression

- Kernel Ridge Regression (KRR)

John Shawe-Taylor and Nello Cristianini. Kernel Methods for Pattern Analysis. Cambridge University Press, 2004.

- Relevance Vector Regression (RVR)

Michael E. Tipping. Sparse bayesian learning and the relevance vector machine. Journal of Machine Learning Research, 1:211(244), 2001.

- Gaussian Process Regression (GPC)

GPLML implementation,

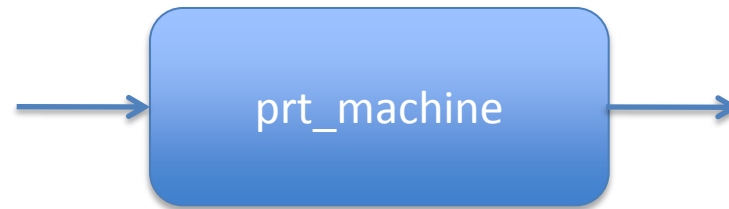
## Developer point of view

### Structure containing:

- Data/Kernel
- Labels
- Etc

### Structure containing:

- Function name (machine)
- Arguments



### Structure containing:

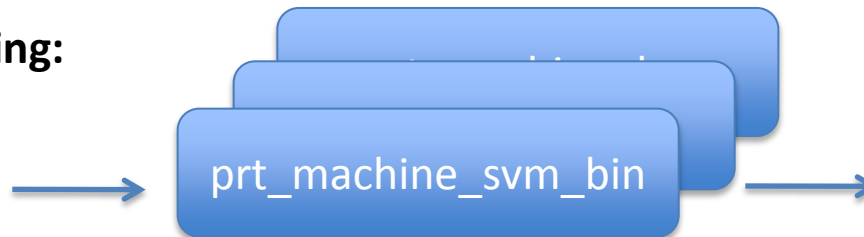
- Predictions
- Coefficients/Weights
- etc

### Machine Library

(classification and regression models)

### Structure containing:

- Data/Kernel
- Labels
- Etc

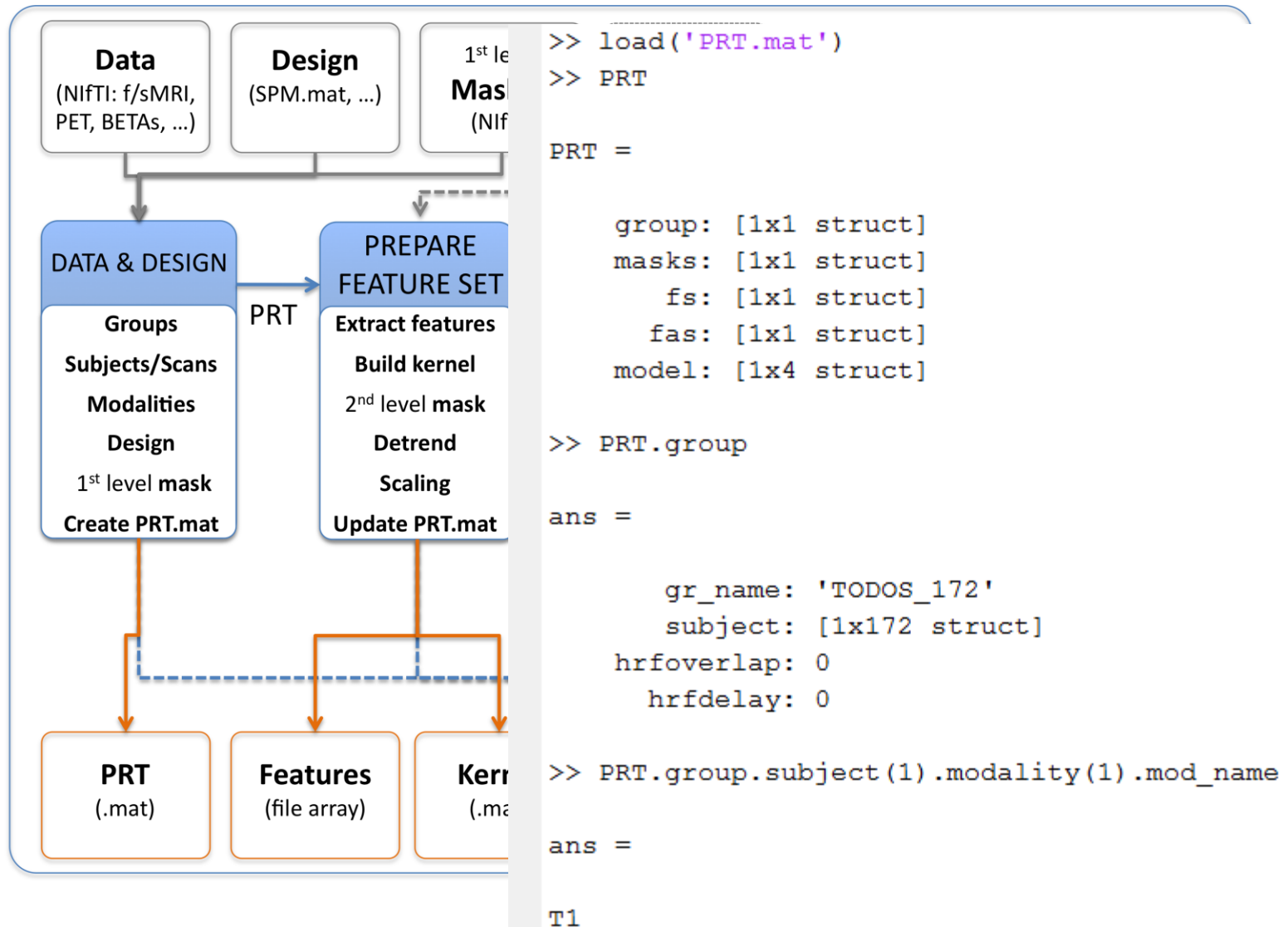


### Structure containing:

- Predictions
- Coefficients/Weights

# PRoNTo

## PRoNTo FRAMEWORK



PRT	group	gr_name				
		subject	subj_name()			
			modality()	mod_name		
				TR		
				scans		
				design	conds	cond_name()
						onsets()
						durations()
						rt_trial()
						scans()
						blocks()
						discardedscans()
						hrfdiscardedscans()
					stats	overlap
						goodscans
						discscans
						meanovl
						stdovl
						mgoodovl
						sgoodovl
						goodovl
					TR	
					unit	
					covar	

# P<sub>R</sub>oN<sub>T</sub>o

PRT	mask	mod_name				
		fname				
	fs	fs_name				
		k_file				
		id_col_names				
		fas		im		
				ifa		
		modality		mod_name		
				Detrend		
				param_dt		
				mode		
				idfeat_fas		
				normalise		type
						scaling
		id mat				

# PRoNTTo

fas	mod_name	
	Dat	
	Detrend	
	param_dt	
	hdr	fname
		dim
		mat
		pinfo
		dt
		n
		descrip
		private
	idfeat_img	



PRT

model

model_name()				
input()	use_kernel			
	type			
	machine	function		
		Args		
	class	class_name		
		group()	gr_name()	
			subj	num()
				modality
	fs	fs_name		
	samp_idx			
	Targets			
	targ_allscans			
	cv_mat			
operations				
cv_type				
output()	fold	targets()		
		predictions()		
		stats()	con_mat	
			acc	
			c_acc	
			b_acc	
			acc_lb	
			acc_ub	
		func_val()		
		type()		
		alpha()		
		b()		
	totalSV()			
	stats	con_mat		
		acc		
		c_acc		
		b_acc		
acc_lb				
acc_ub				

# PRoNTo

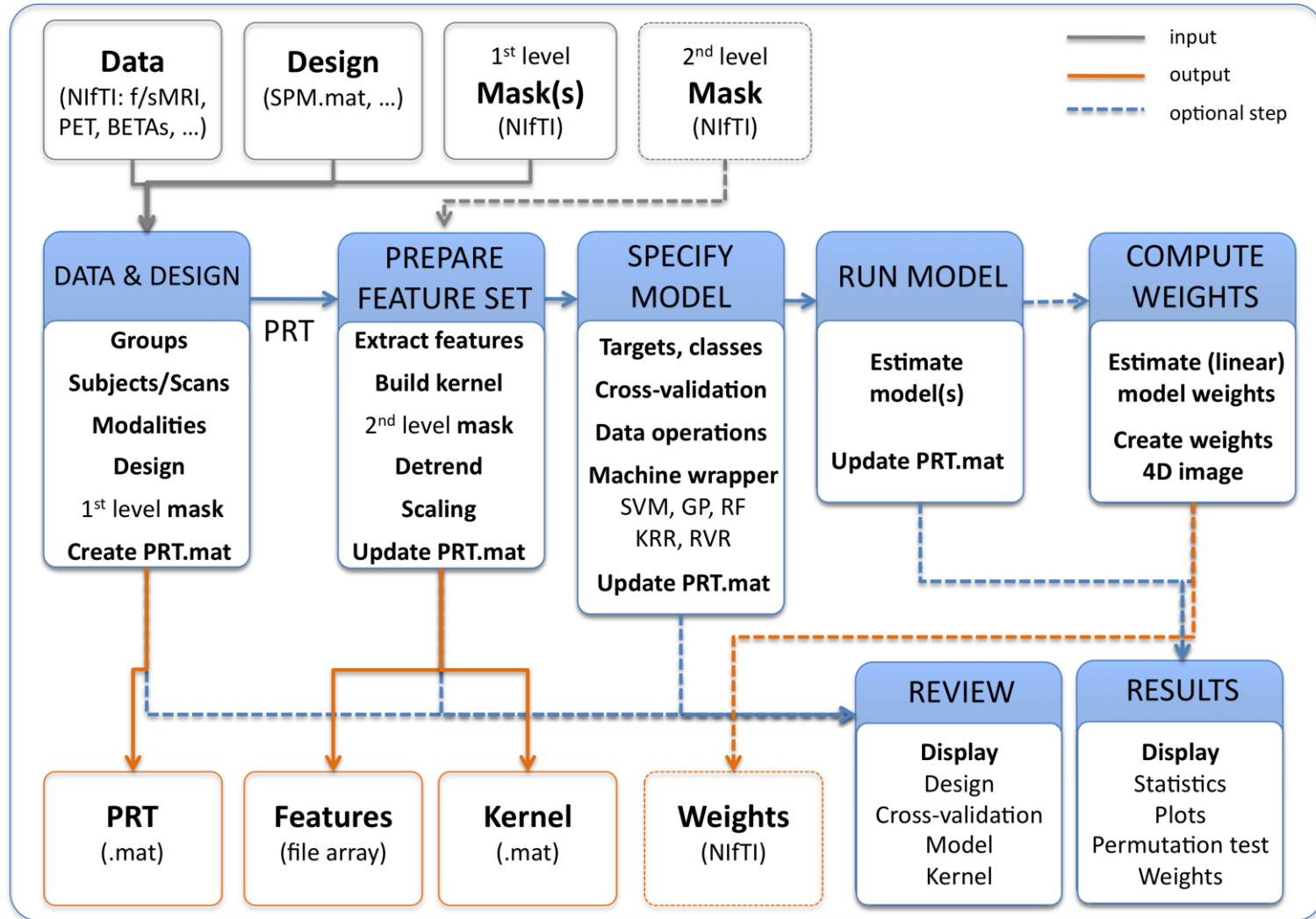
## Data format:

- Input images in NIfTI (Neuroimaging Informatics Technology Initiative), <http://nifti.nimh.nih.gov/>
- Other format could be supported but this would require coding the I/O routine...

Note: sometimes it's simpler to turn your data into a NIfTI 'image'

# PRoNTo

## PRoNTo FRAMEWORK




# Download

## Available here:

<http://www.mlnl.cs.ucl.ac.uk/pronto/>

PATTERN RECOGNITION FOR NEUROIMAGING TOOLBOX (PRoNTO)


UCL

**GO**
[UCL Home » MLNL » PRoNTo](#)

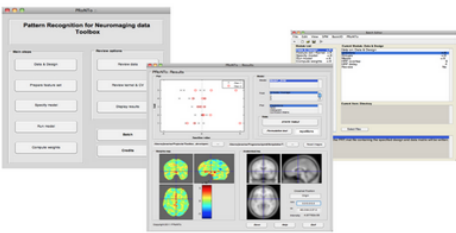
### PRoNTo Menu

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## Pattern Recognition for Neuroimaging Toolbox (PRoNTo)

**PRoNTo** is a toolbox based on machine learning techniques for the analysis of neuroimaging data. The development of the toolbox has been supported by the **PASCAL Harvest** framework and **The Wellcome Trust**.

PRoNTo aims to facilitate the interaction between the machine learning and neuroimaging communities. The toolbox provides a variety of tools for the neuroscience and clinical neuroscience communities, enabling these communities to ask new questions that cannot be easily investigated using existing statistical analysis tools. On the other hand the machine learning community can easily contribute to the toolbox with novel published machine learning algorithms. The toolbox code is distributed for free, but as copyright software under the terms of the GNU General Public License as published by the Free Software Foundation.



### Latest news

- 11/04/2012 - **PRoNTo** (including documentation) will be officially released in May 2012: [download here](#)
- 11/04/2012 - **PRoNTo** team will run a course in London, May 2012: [more info here](#)

## Credits

The development of PRoNTTo was possible with the financial and logistic support of:

- PASCAL Harvest Programme  
(<http://www.pascal-network.org/>)
- the Department of Computer Science, University College London  
(<http://www.cs.ucl.ac.uk>);
- the Wellcome Trust;
- PASCAL2 (<http://www.pascal-network.org/>);
- the Fonds de la Recherche Scientifique-FNRS, Belgium  
(<http://www.fnrs.be>);
- The Foundation for Science and Technology, Portugal  
(<http://www.fct.pt>);
- Swiss National Science Foundation (PP00P2-123438) and Center for Biomedical Imaging (CIBM) of the EPFL and Universities and Hospitals of Lausanne and Geneva.

# PRoNTTo



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