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اصفهان



دانشگاه علوم پزشکی  
قزوین

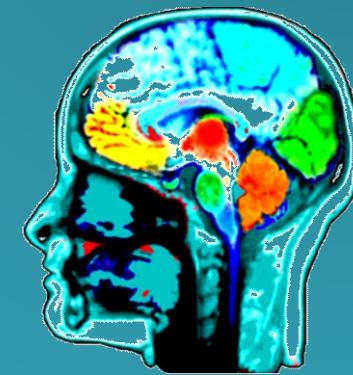


گروه تصویر پردازی و  
آلتایز تصاویر مغزی  
بیمارستان آمام خمینی  
تهران



# Workshop on Imaging and analysis of fMRI data

School of Advanced Technologies in Medicine  
13th Esfand 1399 - 12th Khordad 1400  
Isfahan, Iran



## Software Introduction

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Amirkabir university of technology  
[hakim@aut.ac.ir](mailto:hakim@aut.ac.ir)

# Content

Workshop-General Picture

AnyDesk

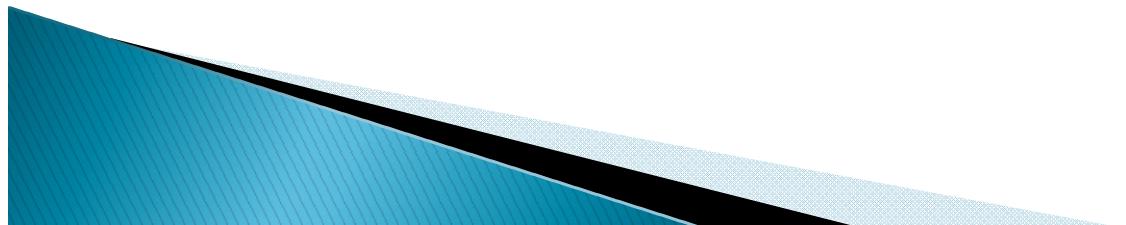
MATLAB

SPM

DPARSF

MRICro

Dcm2nii



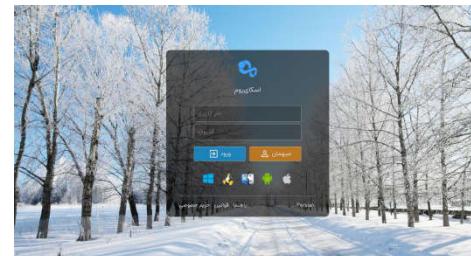
# Workshop-General Picture



Isfahan fMRI Workshop



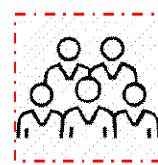
Skyroom



recording

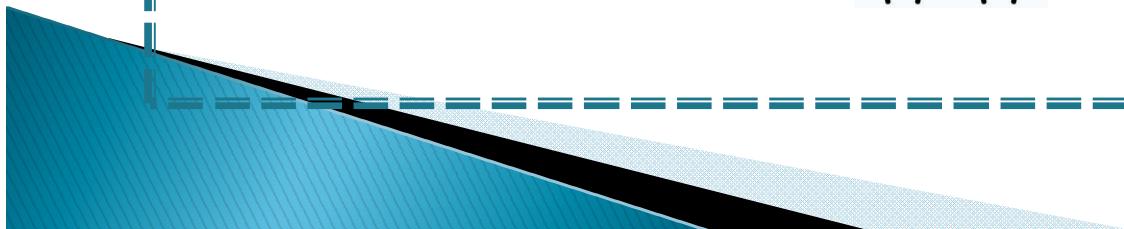


Troubleshooting

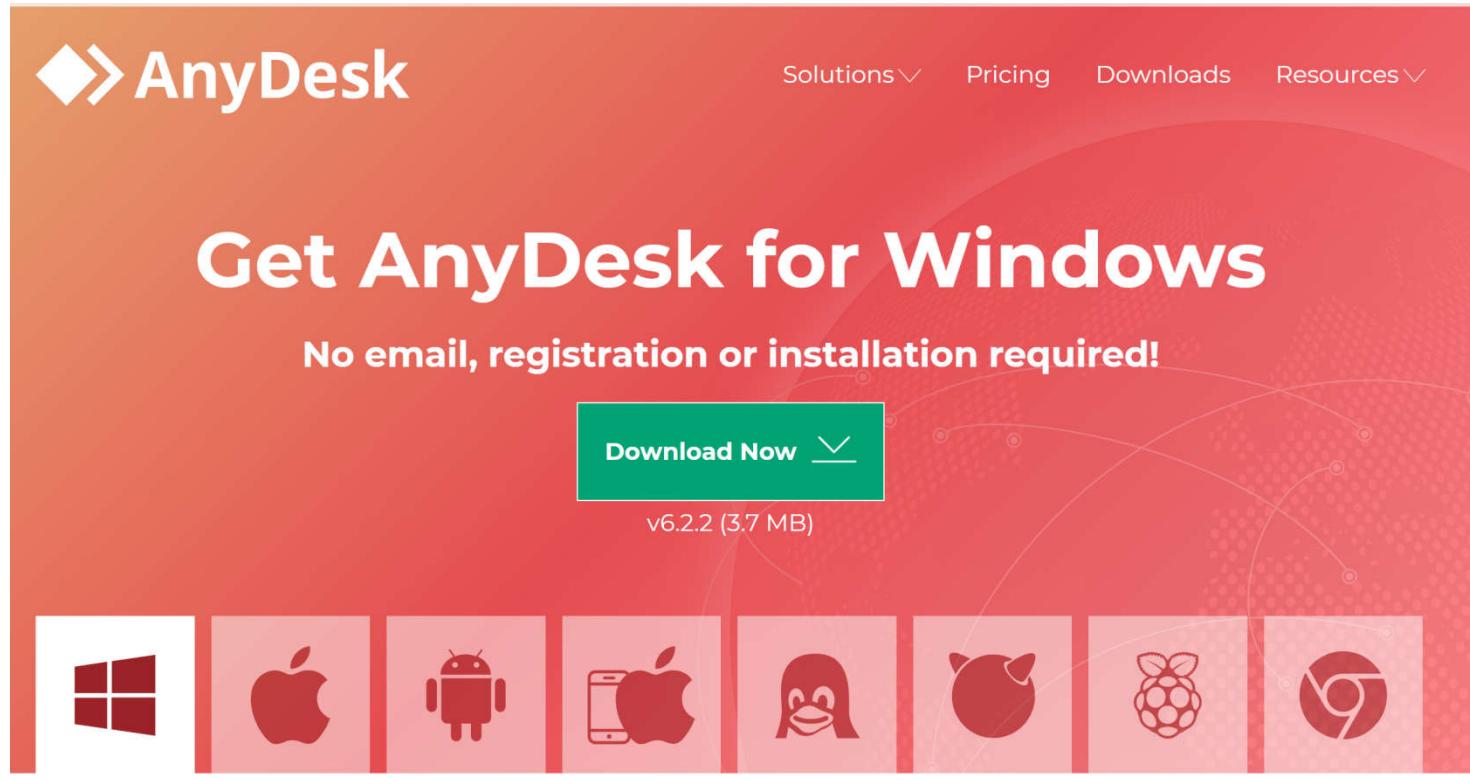


File sharing

Google  
Drive  
Website



# AnyDesk



The image shows the AnyDesk website's landing page for Windows. The background is a vibrant orange-red gradient with a subtle globe and network line pattern. At the top left is the AnyDesk logo (diamond icon and text). At the top right are navigation links: Solutions (with a dropdown arrow), Pricing, Downloads, and Resources (with a dropdown arrow). The main headline is "Get AnyDesk for Windows" in large white font. Below it is a sub-headline: "No email, registration or installation required!". A prominent green button with the text "Download Now" and a downward arrow is centered. To its right, the file size "v6.2.2 (3.7 MB)" is displayed. Below the download area is a row of eight icons representing supported operating systems: Windows, macOS, Android, iOS, Linux, FreeBSD, Raspberry Pi, and Chrome OS. Each icon has its name written below it in red text.

◆ AnyDesk

Solutions ▾ Pricing Downloads Resources ▾

## Get AnyDesk for Windows

No email, registration or installation required!

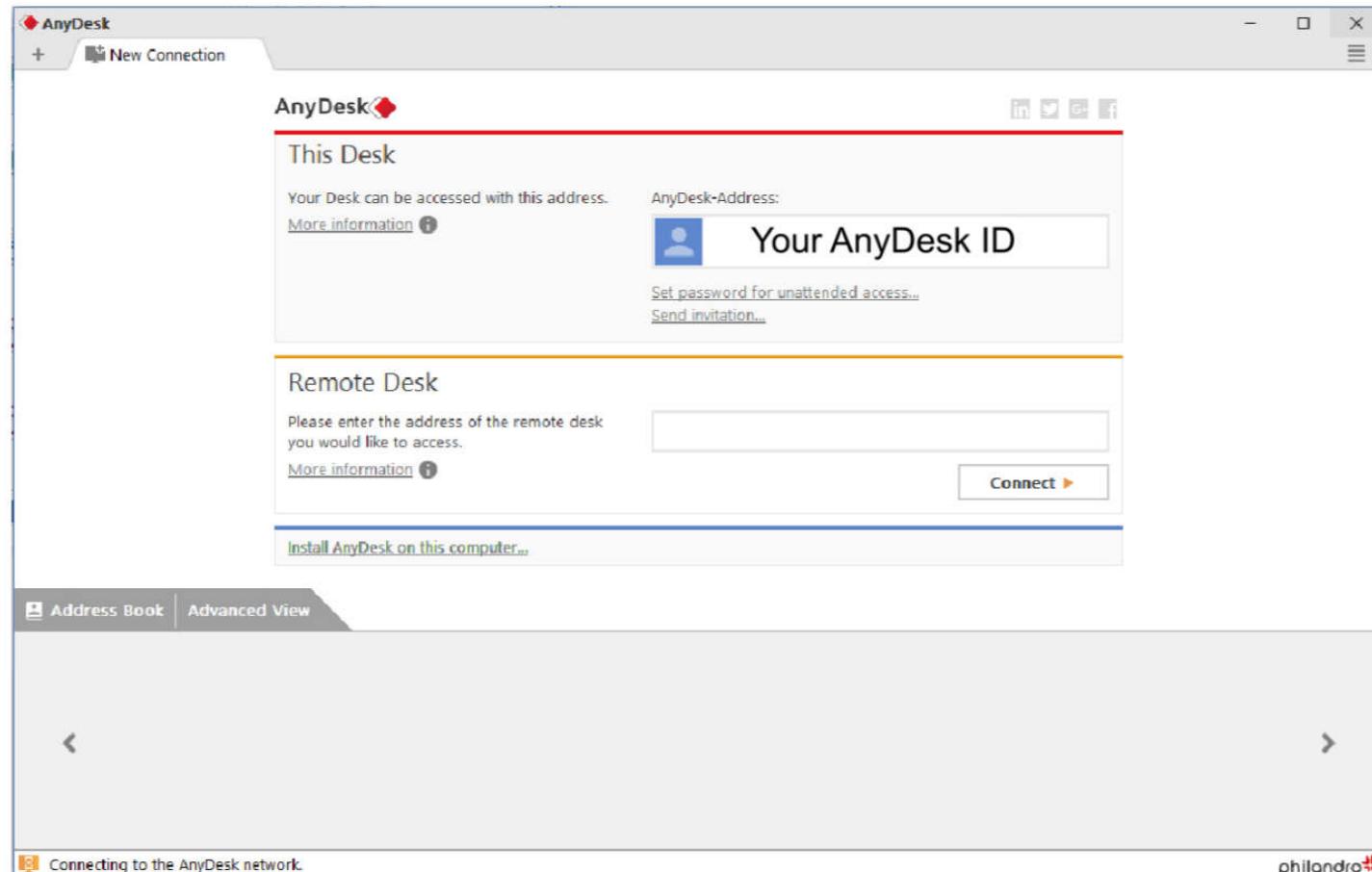
Download Now

v6.2.2 (3.7 MB)

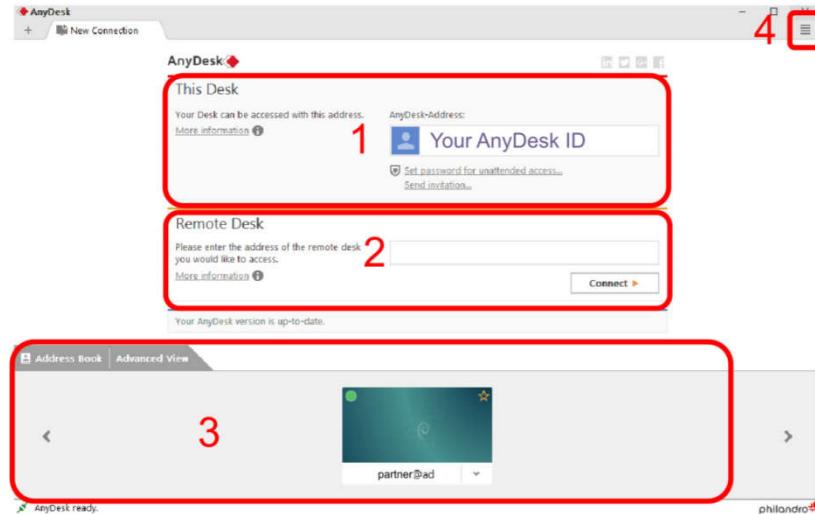
Windows macOS Android iOS Linux FreeBSD Raspberry Pi Chrome OS

<https://anydesk.com/en>

# AnyDesk window



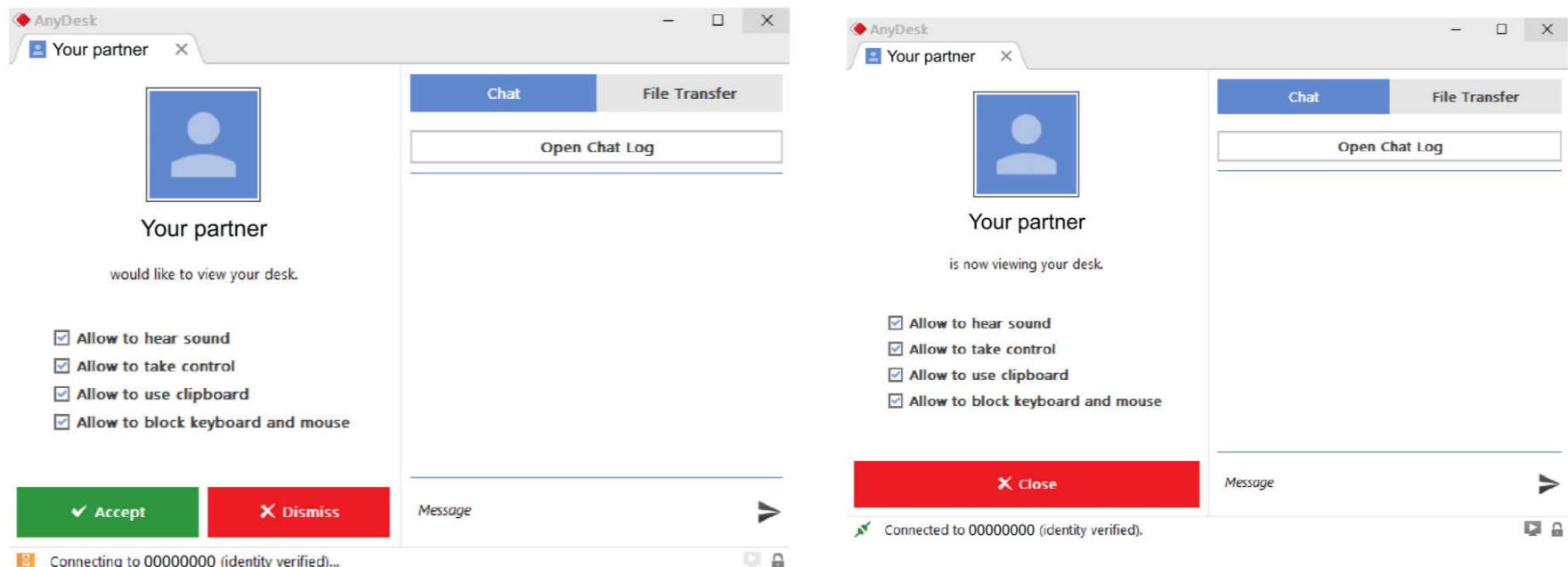
# AnyDesk window



1	<b>This Desk:</b> This is where you will see your AnyDesk address that you use to connect to other computers. Under the menu option "Settings", you can choose to display your AnyDesk address as an alias or a number [23].
2	<b>Remote Desk:</b> This is where you establish a connection to another computer. Simply enter the AnyDesk address of the remote computer into the field and click on "Connect".
3	<b>Quick dial:</b> The quick dial allows you to quickly access your previous connections. Simply click on a connection to enter the information directly into the field and establish a connection [8].
4	<b>Menu bar:</b> You will only be able to access the "General" menu here at first, including the settings [18]. During an active session, additional menus will become available.

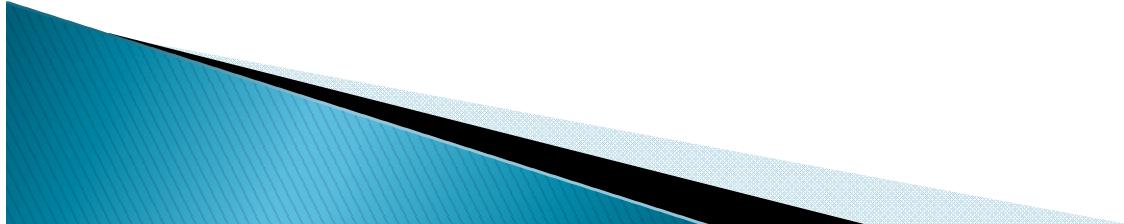
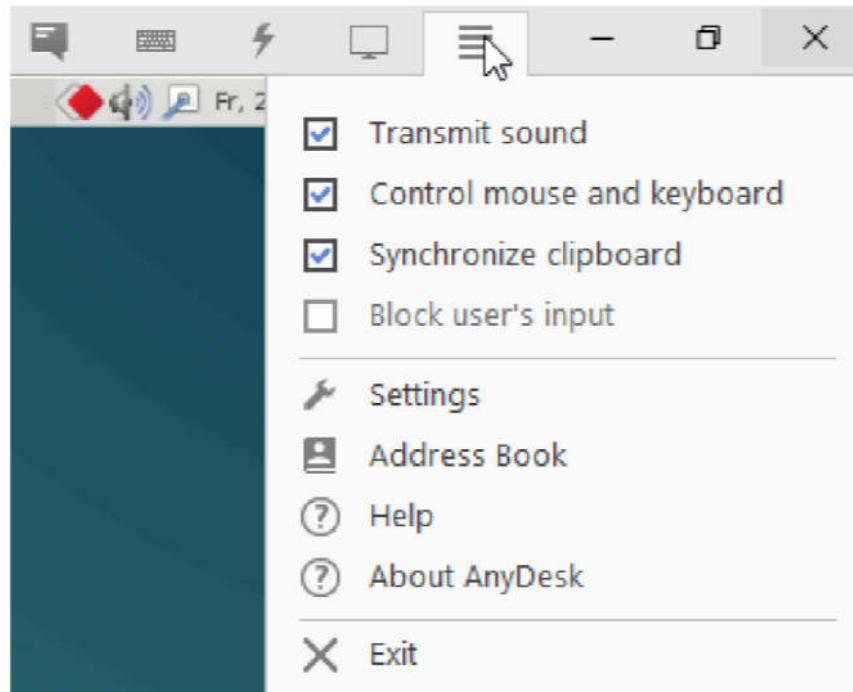
# AnyDesk

## Allowing access to your own desk



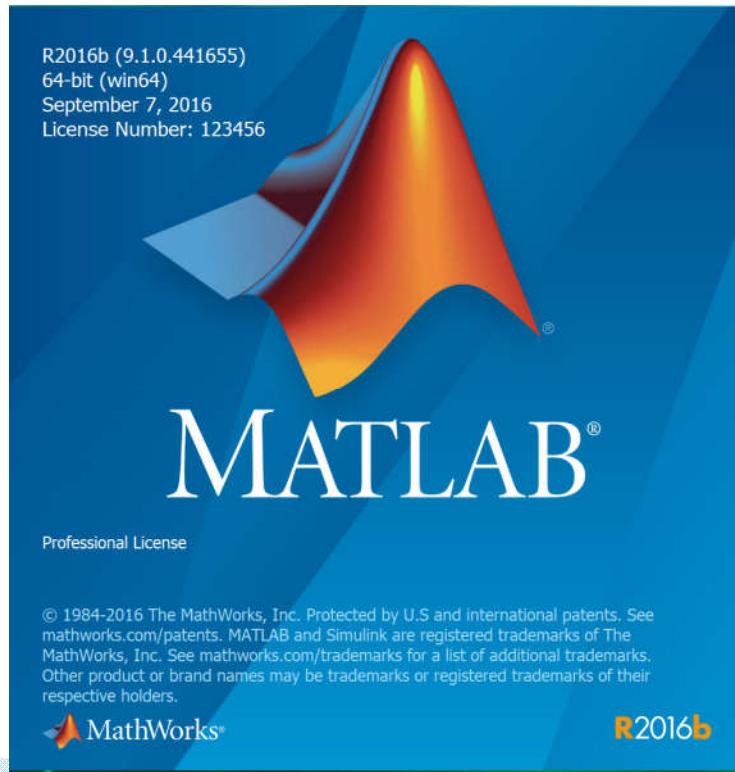
# AnyDesk-Menu

## "General" menu

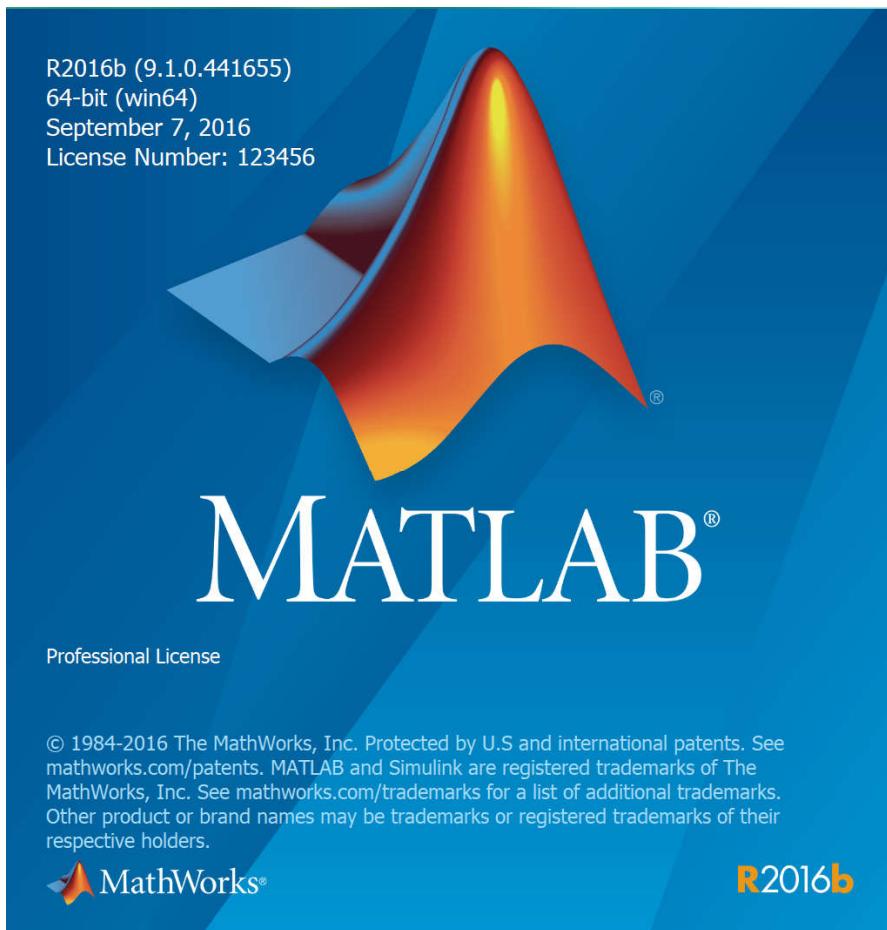


# MATLAB

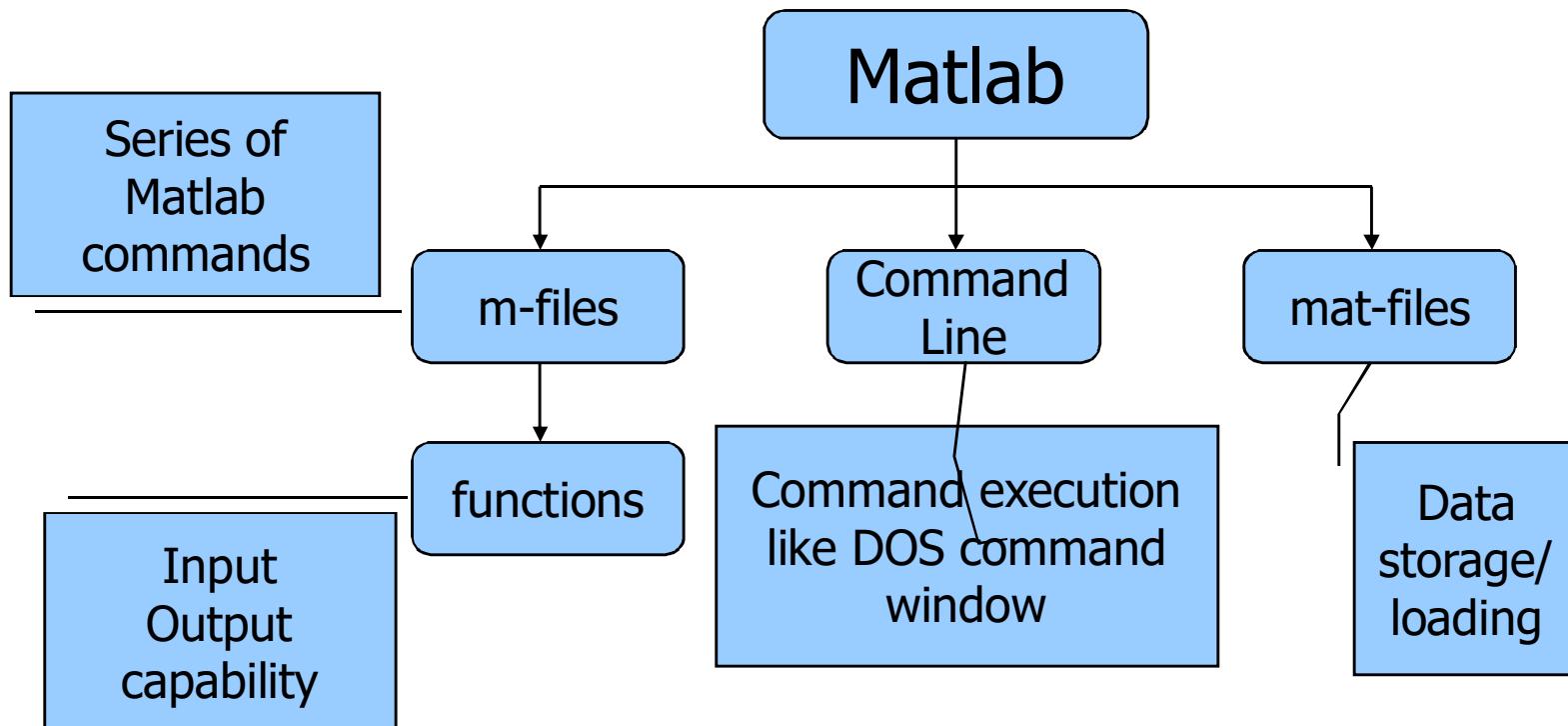
- ▶ Matlab = Matrix Laboratory
- ▶ Easy to learn
- ▶ Robust



# MATLAB-setup

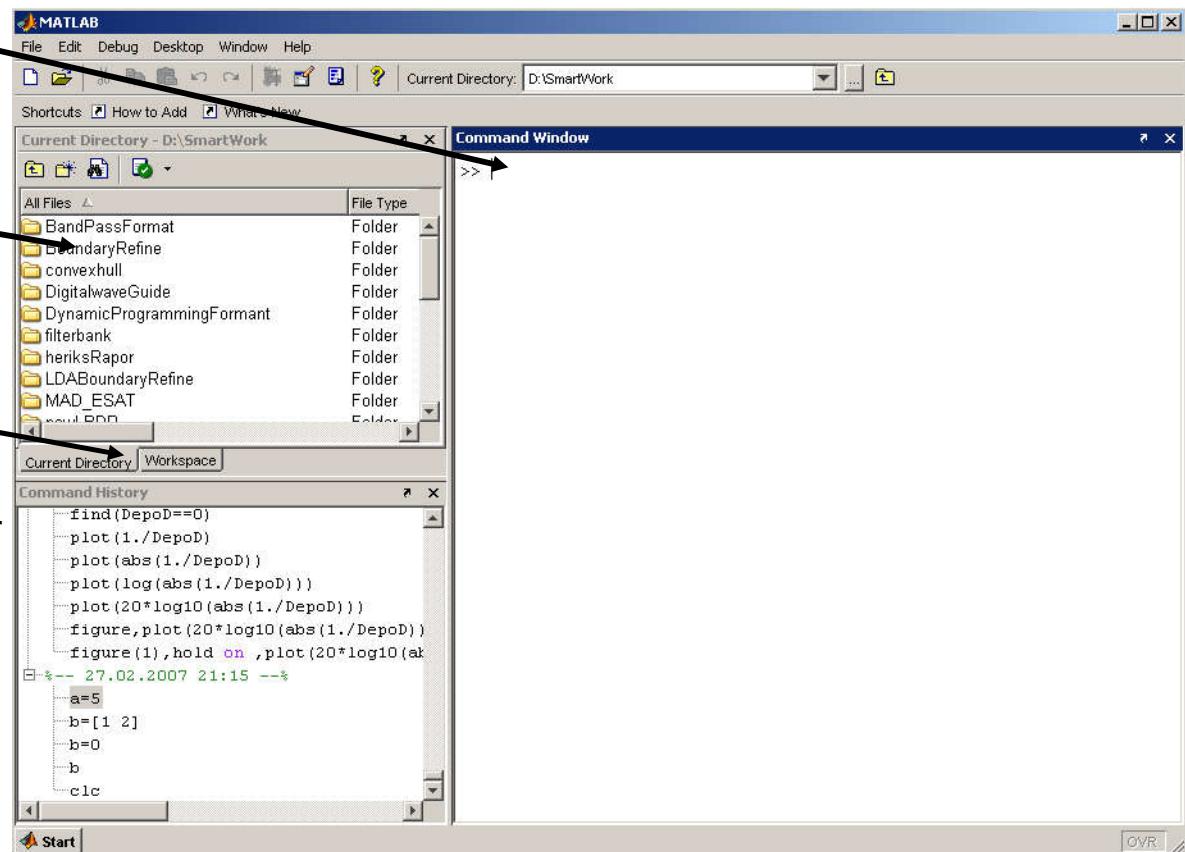


# MATLAB-General Picture



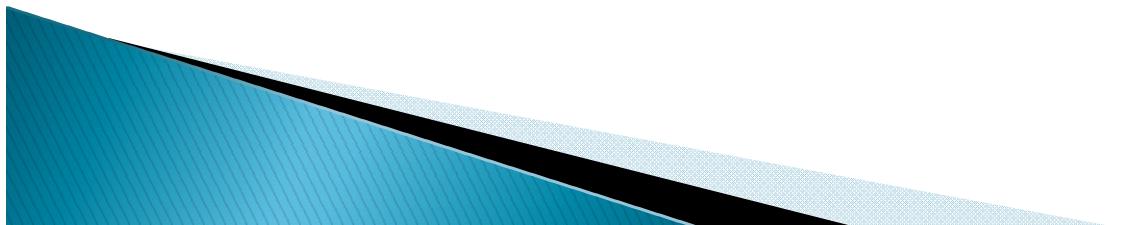
# MATLAB-General Picture

- ▶ **Command Window**
  - type commands
- ▶ **Current Directory**
  - View folders and m-files
- ▶ **Workspace**
  - View program variables
  - Double click on a variable to see it in the Array Editor
- ▶ **Command History**
  - view past commands
  - save a whole session using diary



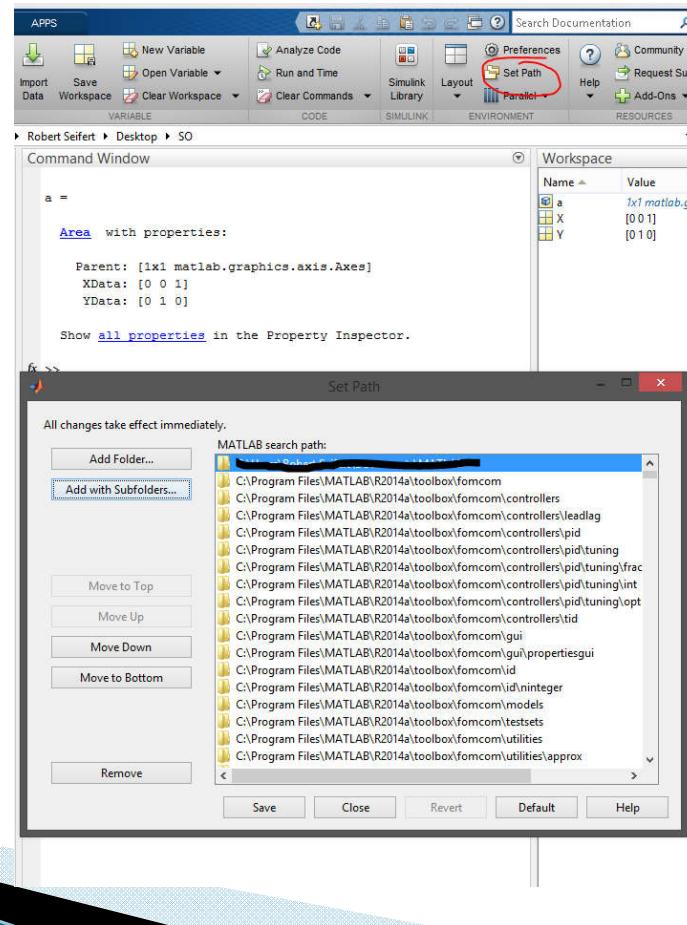
# Exercise

- ▶ Get Matlab version
  - Run “version()” expression in command line
- ▶ Set current directory to desktop folder



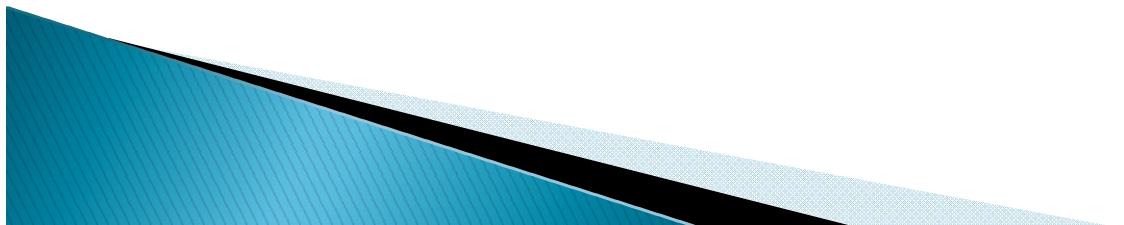
# Matlab - use other tools

- ▶ matlab search path



# Exercise

- ▶ Add path SPM toolbox to matlab path



# SPM–Dataset

University College London

Statistical Parametric Mapping(test hypotheses about functional imaging data)

fMRI, PET, SPECT, EEG and MEG.

## Requirements

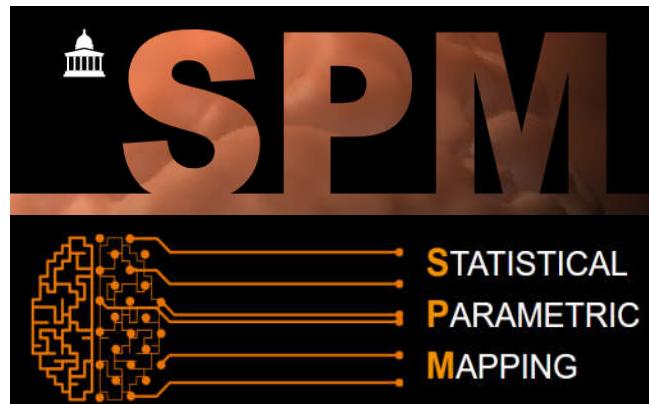
- MATLAB
- Support platforms(windows,Linux,MacOS)

## Standalone version

From 1998(SPM99)–2020(SPM12)

<https://www.fil.ion.ucl.ac.uk/spm/>

Copy left!



MATLAB Compatibility with SPM [edit]

Version	Release	Date	SPM12	SPM8	SPM5 [1]	SPM2	SPM99
9.8	R2020a	2020					
9.7	R2019b	2019					
9.6	R2019a	2019					
9.5	R2018b	2018					
9.4	R2018a	2018					
9.3	R2017b	2017					
9.2	R2017a	2017					
9.1	R2016b	2016					
9.0	R2016a	2016					
8.6	R2015b	2015					
8.5	R2015a	2015					
8.4	R2014b	2014					
8.3	R2014a	2014					
8.2	R2013b	2013					
8.1	R2013a	2013					
8.0	R2012b	2012					
7.14	R2012a	2012					
7.13	R2011b	2011					
7.12	R2011a	2011					
7.11	R2010b	2010					
7.10	R2010a	2010					
7.9	R2009b	2009					
7.8	R2009a	2009					
7.7	R2008b	2008					
7.6	R2008a	2008					
7.5	R2007b	2007					
7.4	R2007a	2007					
7.3	R2006b	2006					
7.2	R2006a	2006					
7.1	R2005b	2005					
7.0	R2005a	2005					
6.9	R2004b	2004					
6.8	R2004a	2004					
6.7	R2003b	2003					
6.6	R2003a	2003					
6.5	R2002b	2002					
6.4	R2002a	2002					
6.3	R2001b	2001					
6.2	R2001a	2001					
6.1	R2000b	2000					
6.0	R2000a	2000					
5.9	R1999b	1999					
5.8	R1999a	1999					



# SPM- Courses



Home Software Documentation Courses Email list Data sets Extensions

## SPM Courses

### London, May & October 2021

The FIL SPM course on using Statistical Parametric Mapping for neuroimaging is held each May and October as part of the Institute of Neurology's short course programme.

Since 2010, we provide two separate SPM courses reflecting the different imaging modalities. There is a three-day course on *SPM for EEG/MEG* followed by the long-established three-day course on *SPM for fMRI/VBM/PET*. Both courses are suitable for beginners and more advanced users. We advise students to gain at least some minimal familiarity with the methodology, for example, from reading introductory articles available from the SPM web page or by following data analysis examples in the SPM manual.

The next SPM course in London will take place in **May 2021** but the exact dates have not been finalised yet.

### Externally organised SPM courses

Other SPM courses are organised around the world at different times of year and in different languages. If you do organise one, please [let us know](#) so that it can be listed here.

Training programs at the Martinos Center, Boston, Massachusetts, USA:  
<http://www.martinos.org/training/courses>

TReNDS Center, Georgia State University, Atlanta, USA:



# SPM–Dataset

The screenshot shows the homepage of the SPM-Dataset website. At the top left is the SPM logo, which includes a stylized brain icon and the text "STATISTICAL PARAMETRIC MAPPING". To the right is a large orange banner with the letters "SPM" in white. Below the banner is a navigation bar with links: Home, Software, Documentation, Courses, Email list, Data sets, and Extensions. The main content area has a light gray background. A large orange sidebar on the left contains links to various datasets. The main text area starts with a section titled "Data sets and tutorials" and "Introduction". It then describes the available data sets for training and personal education. Below this, it details the fMRI: epoch dataset, mentioning block-design fMRI analysis, single-subject or 'first-level' analyses, and specific analyses like Auditory - single subject and Attention to Visual Motion - single subject. It also notes that the Attention to Visual Motion data includes instructions for Psychophysiological Interactions (PPIs) and Dynamic Causal Modelling (DCM). The sidebar lists datasets for fMRI: epoch, event-related, multi-subject, DCM, and EEG: MMN.

**SPM**

Home Software Documentation Courses Email list Data sets Extensions

## Data sets and tutorials

### Introduction

The following data sets are being made available for training and personal education and evaluation purposes. Those wishing to use these data for other purposes, including illustrations or evaluations of methods, should contact the Methods group at the Wellcome Centre for Human Neuroimaging.

A set of instructions showing how SPM can be used to analyse each data set are also provided. These tutorials show how one can use SPM to implement analyses of PET data, epoch or event-related fMRI data, and data from a group of subjects using Random effects analyses ([RFX](#)). They also cover more advanced topics such as Psychophysiological Interactions ([PPIs](#)) and Dynamic Causal Modelling ([DCM](#)).

### fMRI: epoch

The instructions accompanying these data sets show you how to implement a block-design fMRI analysis in SPM. They are both single-subject or 'first-level' analyses.

- Auditory - single subject
- Attention to Visual Motion - single subject

The instructions accompanying the Attention to Visual Motion data also show you how to use SPM to implement, for example, Psychophysiological Interactions ([PPIs](#)) and Dynamic Causal Modelling ([DCM](#)).

### fMRI: event-related

Home  
Software  
Documentation  
Courses  
Email list  
**Data sets**  
Introduction  
fMRI: epoch  
fMRI: event-related  
fMRI: multi-subject  
fMRI: DCM  
EEG: MMN

# SPM- Extensions

## Quick Links

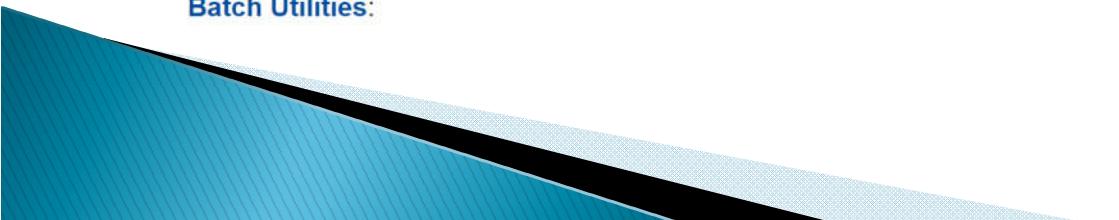
### Toolboxes:

[AAL](#) | [AAL2](#) | [AAL3](#) | [ACID](#) | [AICHA](#) | [ALI](#) | [ALVIN](#) | [AMAT](#) | [AnalyzeMovie](#) | [Anatomy](#) | [AQuA](#) | [ArtRepair](#) | [aslM](#) | [ASLtbx](#) | [at4fmri](#) | [aws4SPM](#) | [BAAD](#) | [BFAST3D](#) | [BrainNetViewer](#) | [Brainnetome](#) | [BRANT](#) | [BredeQuery](#) | [Bruker2nifti](#) | [bspmview](#) | [CAT](#) | [CCAfMRI](#) | [CLASS](#) | [Clinical](#) | [Complexity](#) | [conn](#) | [ConnExT](#) | [CPCA](#) | [DAiSS](#) | [DICOMCD\\_Import](#) | [Diffusion\\_II](#) | [DPABI](#) | [DPARSF](#) | [DRIFTER](#) | [EEGAnalyzer](#) | [EMS](#) | [ExtractVals](#) | [FASL](#) | [FAST](#) | [fECM](#) | [FDR](#) | [FieldMap](#) | [fieldmap\\_undistort](#) | [FieldTrip](#) | [fMRIPower](#) | [fOSA](#) | [gPPI](#) | [GraphVar](#) | [GridCAT](#) | [Grocer](#) | [HV](#) | [hMRI](#) | [IBASPM](#) | [iBrainAT](#) | [iBrainLT](#) | [IBZM\\_tool](#) | [ICN\\_Atlas](#) | [ImaGIN](#) | [INRIAlign](#) | [ISAS](#) | [lead-dbs](#) | [lesion\\_gnb](#) | [LI](#) | [LogTransform](#) | [MACS](#) | [Mantis](#) | [MARINA](#) | [MARS](#) | [MarsBar](#) | [MASCOI](#) | [mfBox](#) | [Masking](#) | [Masks](#) | [MEAW](#) | [MIP-C](#) | [MM](#) | [MP2RAGE](#) | [multifocal](#) | [MRTOOL](#) | [MRM](#) | [NIRS-SPM](#) | [NPBayes](#) | [NS](#) | [OCT](#) | [PETPVE12](#) | [pTFCE](#) | [Ortho](#) | [ppi\\_batch\\_hipp](#) | [PSPM](#) | [QModeling](#) | [REST](#) | [rfxplot](#) | [RobustWLS](#) | [rsHRF](#) | [SAFe](#) | [SAMIT](#) | [SCRalyze](#) | [SDM](#) | [SGTT](#) | [SimpleROIBuilder](#) | [SnPM](#) | [SpikeDet](#) | [spm\\_wavelet](#) | [SPMd](#) | [SPMMouse](#) | [SSM](#) | [STEM](#) | [SUIT](#) | [SurfRend](#) | [SwE](#) | [TDT](#) | [PhysIO](#) | [TOM](#) | [TFCE](#) | [UF2C](#) | [Unwarp2](#) | [VarTbx](#) | [VDB](#) | [Volumes](#) | [WBM](#) | [WSPM](#) | [WFU\\_PickAtlas](#) | [xjView](#) | [XMLTools](#) | [ASLtbx](#) | [BENtbx](#) | [SVRLSMtbx](#) | [GIFT](#)

### Utilities:

[AveLI](#) | [BrainMagix](#) | [Bruker2Analyze](#) | [CBMG-Tools](#) | [Design\\_Magic](#) | [dicom2nifti](#) | [DynPET](#) | [Easy\\_ROI](#) | [Easy\\_Volumes](#) | [FDRill](#) | [Fluctuation](#) | [fToolbelt](#) | [FAD](#) | [GA](#) | [GE2SPM](#) | [JG](#) | [L2S](#) | [log\\_roi\\_batch](#) | [LMGS](#) | [MatlabTFCE](#) | [Motion](#) | [mri\\_toolbox](#) | [MSU](#) | [Orth1](#) | [PCT](#) | [pvconv](#) | [r2agui](#) | [SEM](#) | [slice\\_overlay](#) | [TSDiffAna](#) | [TSU](#) | [iTT](#) | [VBMtools](#) | [VIS](#) | [visionToSPM](#) | [VoiTool](#)

### Batch Utilities:



# SPM– Methods



## Preprocessing

---

- Slice Timing
- Realignment
- Unwarp
- Coregistration
- Normalisation
- Segmentation
- Spatial smoothing

## Modelling

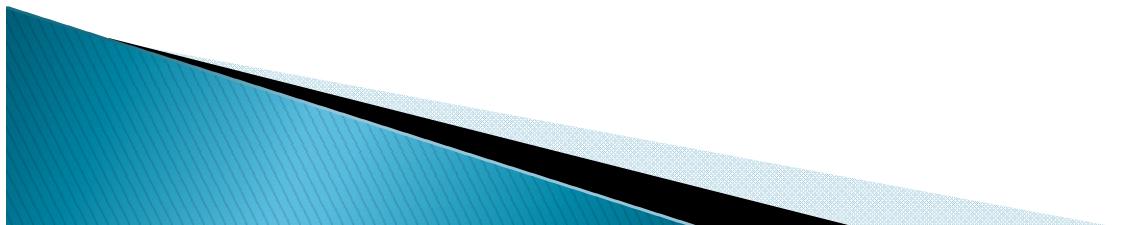
---

- Haemodynamic Response Function
- Basis functions
- General Linear Model
- Correlation and Regression
- Linear Hierarchical Models
- Covariance
- Autocorrelation
- Non-sphericity
- Session concatenation
- Group analysis

## Statistical Inference

---

- F and T tests
- Contrasts
- Inference
- Conjunction
- Power Analysis
- About Correlation
- Bayesian Inference
- Results Table
- Information to include in papers ↗
- Calculating Percentage signal change
- Comparing a single patient versus a group of controls
- Timeseries extraction

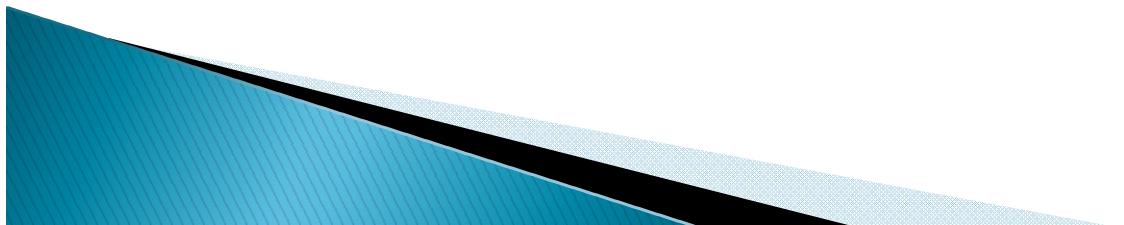


# SPM-Tools

## Other tools

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- [Cogent](#) a MATLAB-based stimulus presentation software
- [MRIcron](#) and [MRIcroGL](#) medical images viewers
- [MarsBaR](#) region of interest toolbox for SPM
- [SnPM](#) Statistical nonParametric Mapping
- [Anatomy](#) SPM Anatomy toolbox
- [AAL](#) Anatomical Automatic Labeling
- [WFU\\_PickAtlas](#) a region of interest toolbox for SPM based on the [Talairach Daemon Database](#)
- [Physiological noise correction](#)
- [Diffusion tools](#)



SPM



<https://www.fil.ion.ucl.ac.uk/spm/software/download/>

- ▶ website

## Download SPM

Please complete the following form to download a copy of the SPM software.

The requested SPM version is the only mandatory entry.

If you need further assistance with the installation of SPM, send an email to the [SPM manager](#) after having read the [installation documentation](#).

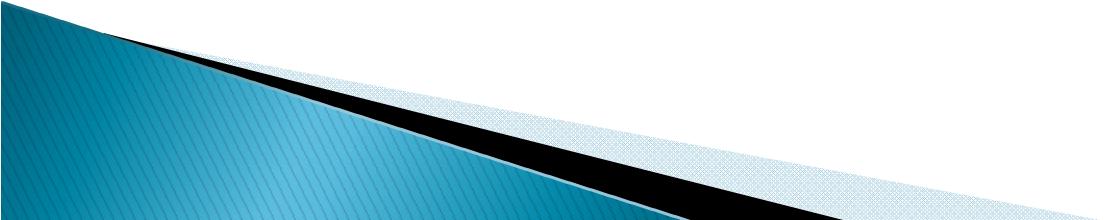
1 SPM Version

Select SPM version required:

Select

2 Additional Info

*It is helpful for us in developing and positioning SPM to know more about how you plan to use our software. All entries are optional.*



# Exercise

## ▶ Run SPM in matlab

### SPM12 [edit]

---

#### Installation [edit]

- Download [spm12.zip](#).
- Unzip `spm12.zip` in a folder of your choice, such as `C:\Users\login\Documents\MATLAB\spm12`.
- Start MATLAB and add SPM into your path, either using *File > Set Path > Add Folder...* or typing

```
>> addpath C:\Users\Login\Documents\MATLAB\spm12
```

in MATLAB's workspace.

- Launch SPM by typing

```
>> spm
```

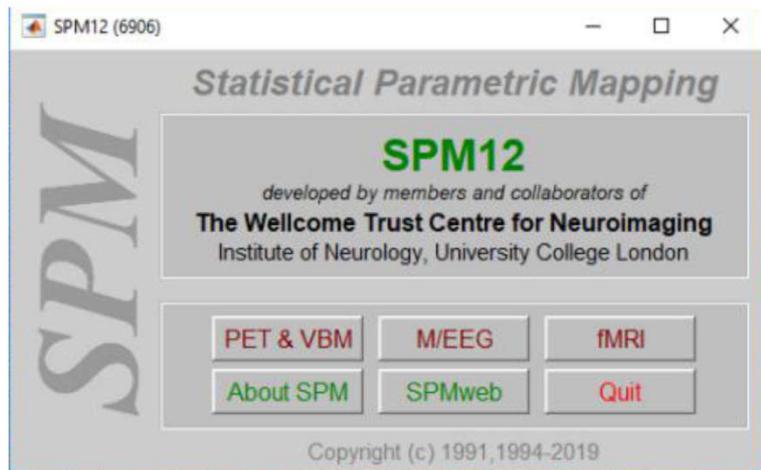
You might have to install the VC++ 2005 and 2008 Redistributable Packages (`vcredist_x86.exe`) from Microsoft:

- <http://support.microsoft.com/kb/2019667>



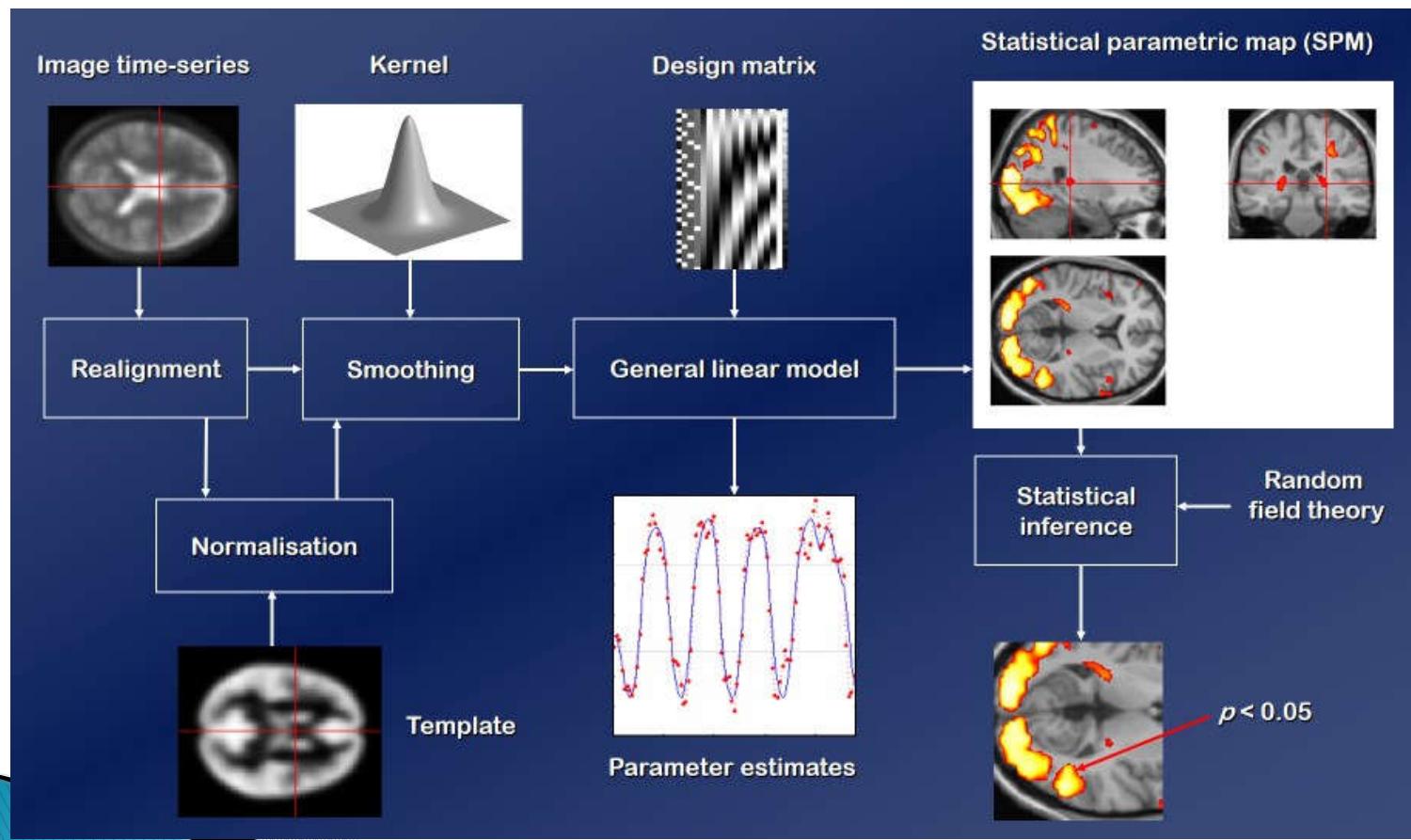
# SPM

## ▶ Window



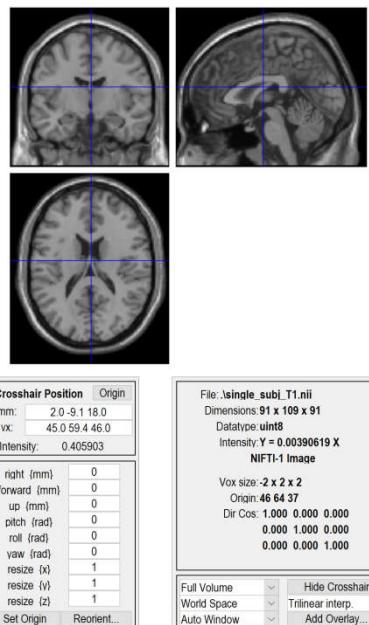
# SPM-Task fMRI

## ▶ Window



# Exercise

- ▶ Display MRI data in SPM
- ▶ Click Display
- ▶ go to spm folder
- ▶ Open the “\spm12\canonical\ single\_subj\_T1.nii”



# DPARSF



The R-fMRI Network

a network for supporting resting-state fMRI related studies.

<http://rfmri.org/DPARSF>

2010–2020

## Data Processing Assistant for Resting-State fMRI (DPARSF)

- Preprocessing data (slice timing, realign, normalize, smooth)
- Connectivity analysis( FC, ReHo, ALFF and fALFF)
- Post processing

## Requirements

- MATLAB
- SPM

frontiers in  
SYSTEMS NEUROSCIENCE

METHODS ARTICLE  
published: 14 May 2010  
doi: 10.3389/fnsys.2010.00013



## DPARSF: a MATLAB toolbox for “pipeline” data analysis of resting-state fMRI

Yan Chao-Gan\* and Zang Yu-Feng\*

State Key Laboratory of Cognitive Neuroscience and Learning, Beijing Normal University, Beijing, China

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Reviewed by:  
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Zang Yu-Feng, State Key Laboratory of Cognitive Neuroscience and Learning, Beijing Normal University, Beijing 100875, China.  
e-mail: zangyf@bnu.edu.cn

Resting-state functional magnetic resonance imaging (fMRI) has attracted more and more attention because of its effectiveness, simplicity and non-invasiveness in exploration of the intrinsic functional architecture of the human brain. However, user-friendly toolbox for “pipeline” data analysis of resting-state fMRI is still lacking. Based on some functions in Statistical Parametric Mapping (SPM) and Resting-State fMRI Data Analysis Toolkit (REST), we have developed a MATLAB toolbox called Data Processing Assistant for Resting-State fMRI (DPARSF) for “pipeline” data analysis of resting-state fMRI. After the user arranges the Digital Imaging and Communications in Medicine (DICOM) files and click a few buttons to set parameters, DPARSF will then give all the preprocessed (slice timing, realign, normalize, smooth) data and results for functional connectivity, regional homogeneity, amplitude of low-frequency fluctuation (ALFF), and fractional ALFF. DPARSF can also create a report for excluding subjects with excessive head motion and generate a set of pictures for easily checking the effect of normalization. In addition, users can also use DPARSF to extract time courses from regions of interest.

Keywords: data analysis, DPARSF, REST, resting-state fMRI, SPM

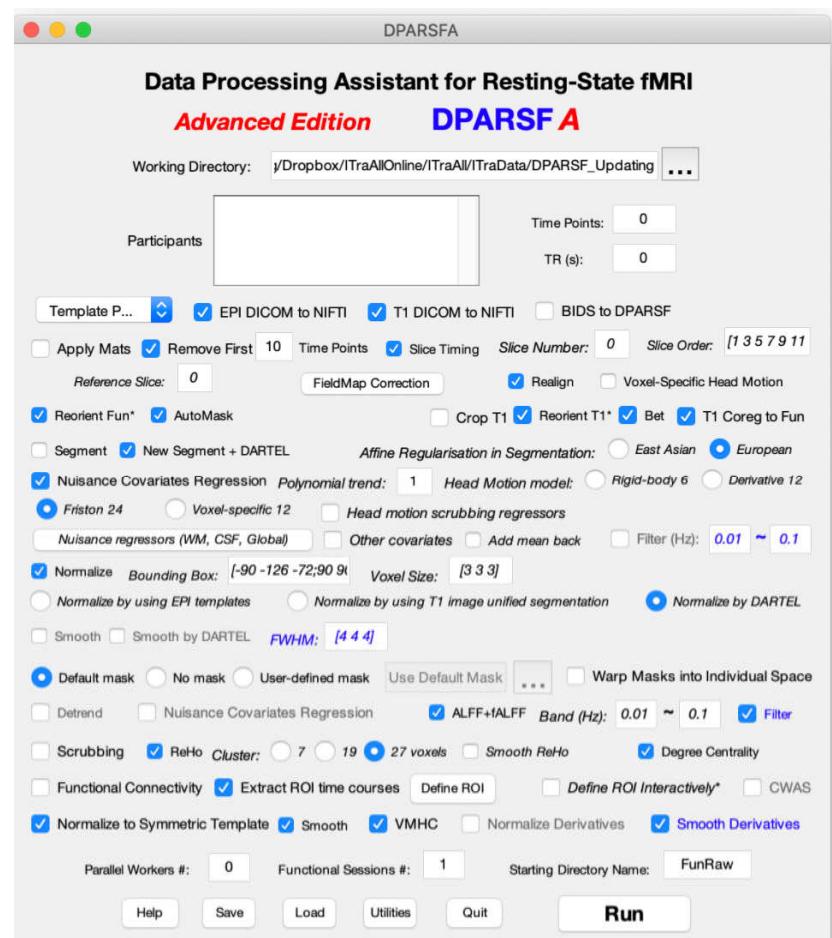
# DPARSF



## ▶ Download link

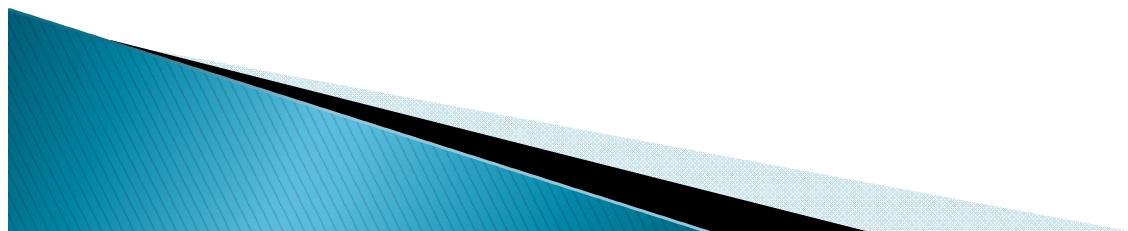
<http://rfmri.org/DPARSF>

## ▶ GUI



# Exercise

- ▶ Run Dparsf





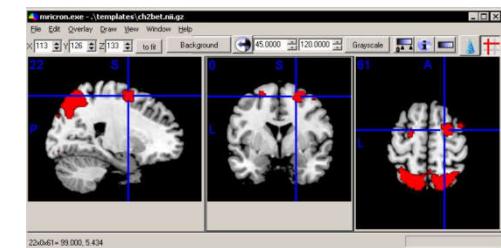
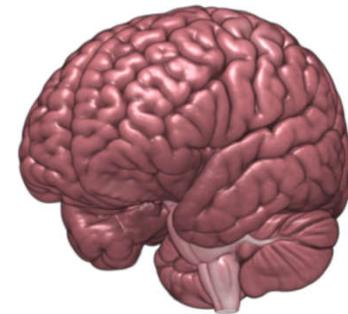
<https://people.cas.sc.edu/rorden/mricron/install.html>

load multiple layers of images

draw volumes of interest.

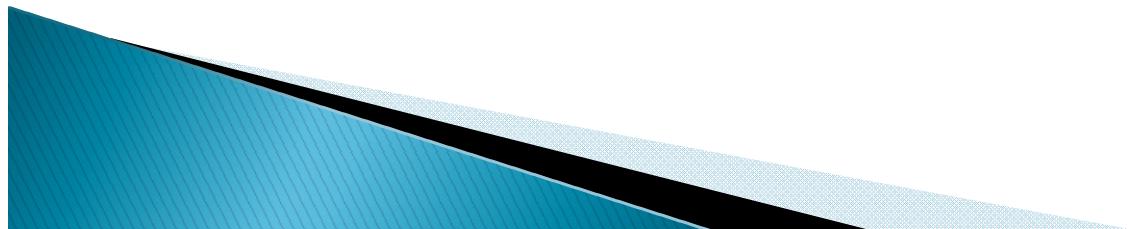
### New MRICronGL

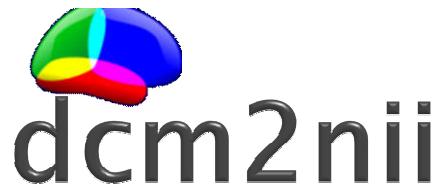
• <https://www.mccauslandcenter.sc.edu/mricrogl/home>



# Exercise

- ▶ Show the MRI data





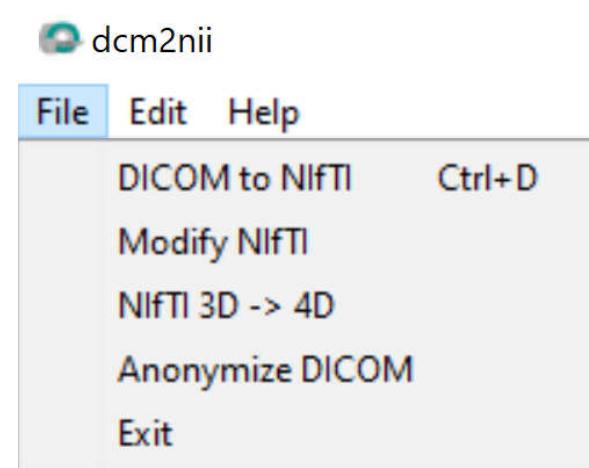
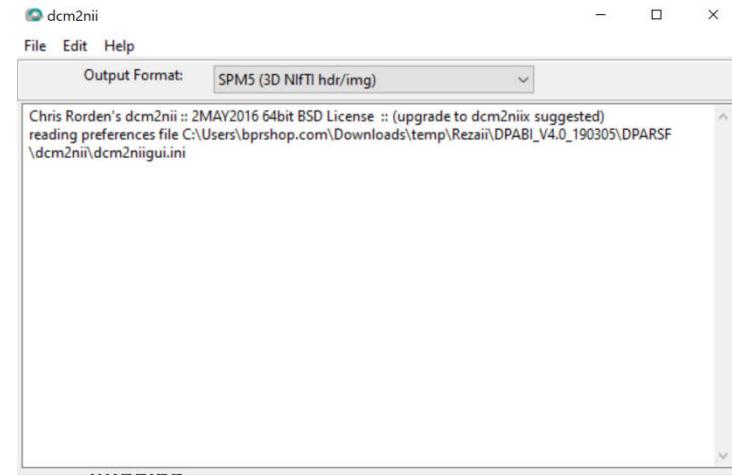
[people.cas.sc.edu/rorden/mricron/dcm2nii.html](http://people.cas.sc.edu/rorden/mricron/dcm2nii.html)

convert to the NIfTI format.

- NIfTI : Neuroimaging Informatics Technology Initiative
  - (a pair of files (hdr/img, compliant with most Analyze format viewers), or a single file (nii))

(Batch+gui)

Modify NIfTI

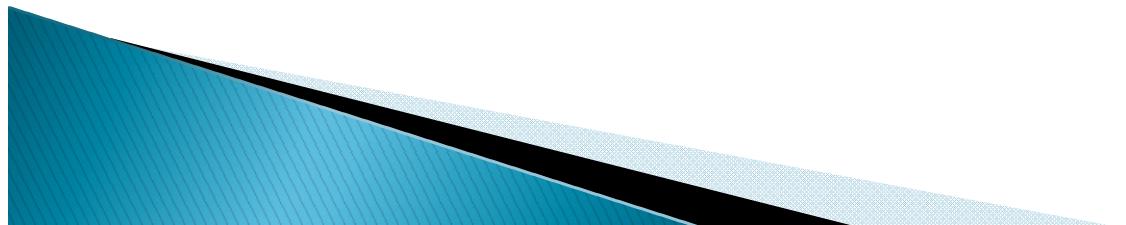


# Exercise

- ▶ Download data

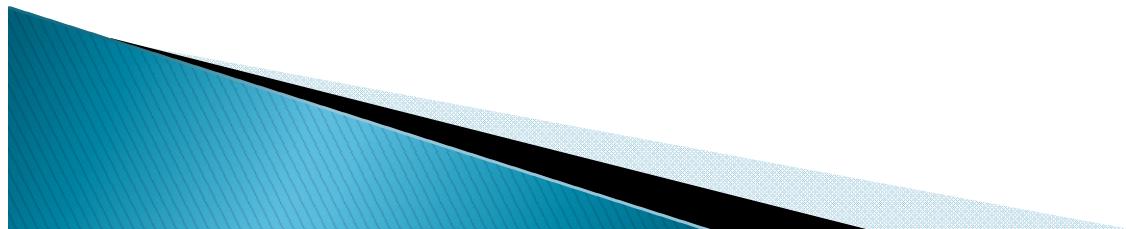
[https://wiki.idoimaging.com/index.php?title=Sample\\_Data](https://wiki.idoimaging.com/index.php?title=Sample_Data)

- ▶ Convert dcm file to nii



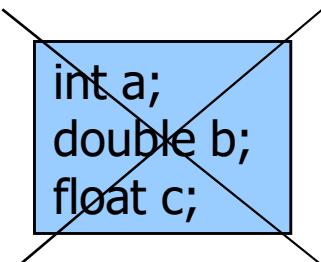
# Matlab Extra

»»



# Matlab- Variables

- ▶ No need for types. i.e.,



- ▶ All variables are created with double precision unless specified and they are matrices.

Example:  
    >>x=5;  
    >>x1=2;

- ▶ After these statements, the variables are  $1 \times 1$  matrices with double precision

# Matlab- Array, Matrix

- ▶ A vector     $x = [1 \ 2 \ 5 \ 1]$

```
x =  
     1    2    5    1
```

- ▶ Transpose  $y = x'$                  $y =$

```
1  
2  
5  
1
```

- ▶ A matrix : Rows are separated by semicolon

```
x = [1 2 3; 5 1 4; 3 2 -1]
```

```
x =  
     1    2    3  
     5    1    4  
     3    2   -1
```

# Matlab–Long Array, Matrix

►  $t = 1:10$

$t =$

1 2 3 4 5 6 7 8 9 10

►  $k = 2:-0.5:-1$

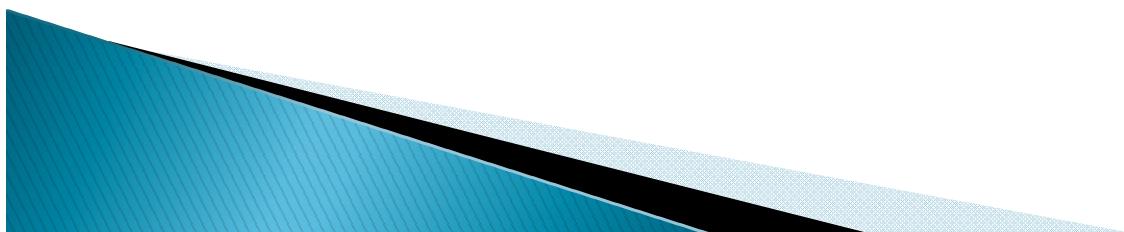
$k =$

2 1.5 1 0.5 0 -0.5 -1

►  $B = [1:4; 5:8]$

$x =$

1	2	3	4
5	6	7	8



# Matlab-Generating Vectors from functions

- ▶ `zeros(M,N)` MxN matrix of zeros

```
x = zeros(1, 3)
```

```
x =
```

```
0 0 0
```

- 
- ▶ `ones(M,N)` MxN matrix of ones

```
x = ones(1, 3)
```

```
x =
```

```
1 1 1
```

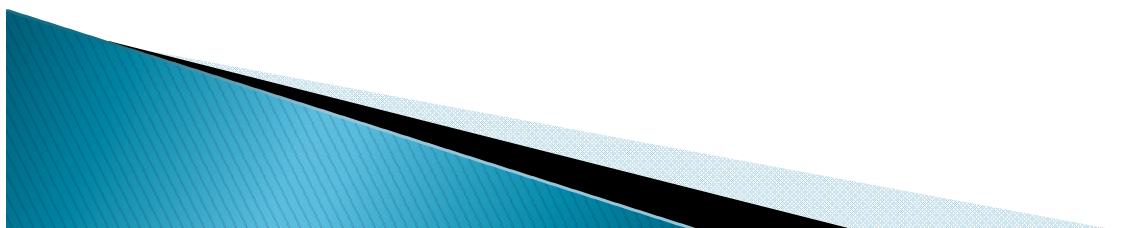
- 
- ▶ `rand(M,N)` MxN matrix of uniformly distributed random numbers on (0,1)

```
x = rand(1, 3)
```

```
x =
```

```
0.9501 0.2311
```

```
0.6068
```



# Matlab- Matrix Index

- ▶ The matrix indices begin from 1 (not 0 as in C)
- ▶ The matrix indices must be positive integer

Given:

```
A =
```

3	5	3
6	8	2
2	7	3

```
>> A(6)
```

```
ans =
```

```
7
```

```
>> A(3,2)
```

```
ans =
```

```
7
```

A(-2), A(0)

Error: ??? Subscript indices must either be real positive integers or logicals.

A(4,2)

Error: ??? Index exceeds matrix dimensions.

# Exercise

- ▶ Create A matrix

```
A =  
3 5 3  
6 8 2  
2 7 3
```

- ▶ Print the element in row 3 and column 2  $\gg A(3, 2)$

```
ans =
```

- ▶ Print all elements in row 2  $\gg A(2, :)$

```
7
```

```
ans =
```

```
6 8 2
```

- ▶ Print first and second elements in column 2  $\gg A(1:2, 2)$

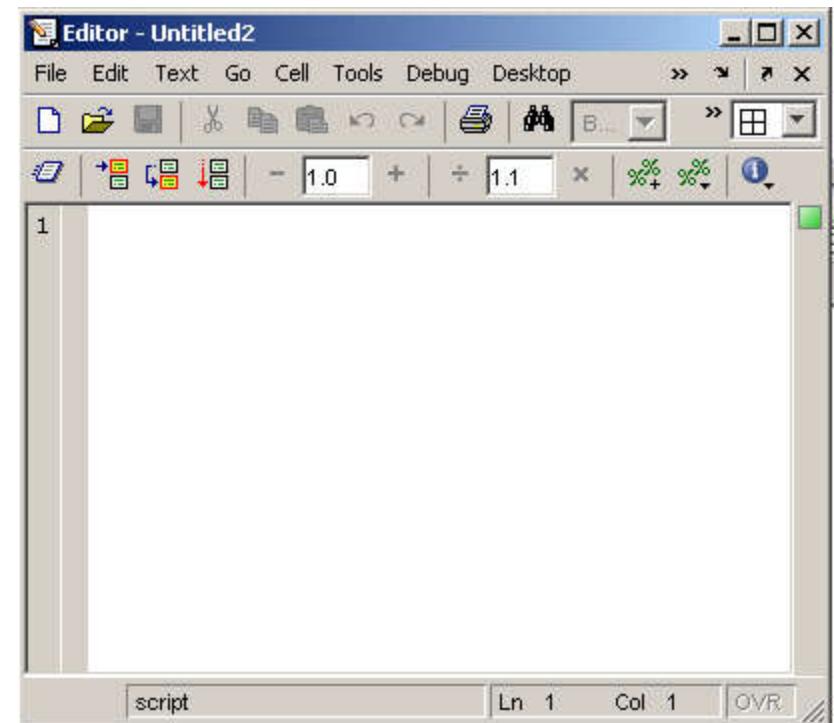
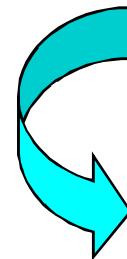
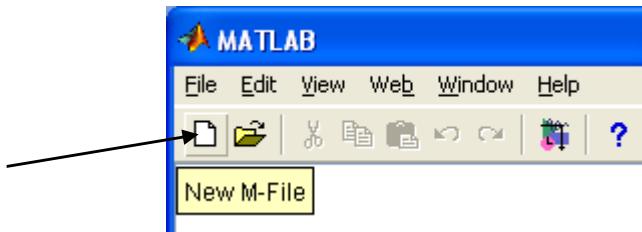
```
ans =
```

```
5  
8
```

Print the element in row 3 and column 2

# Use of M-File

Click to  
create a new  
M-File



- Extension “.m”
- A text file containing script or function or program to run

# Use of M-File

Save file as *Denem430.m*

```
Editor - D:\SmartWork\Denem430.m
File Edit Text Go Cell Tools Debug Desktop Window Help
Stack: Base
1 - x=linspace(0,4*pi,100);
2
3 - y=sin(x);
4 - y1=exp(-x/3);
5 - y2=y.*y1;
6
7 - figure(1)
8 - plot(y2)
9
10 - title('This is the sinus function')
11 - xlabel('x (secs)')
12 - ylabel('sin(x)exp(-x/3)')
13
```

If you include ";" at the end of each statement, result will not be shown immediately

# Exercise

- ▶ Create M file(test.m) plot line

```
a= (1:100);  
plot(a);
```

- ▶ Run test.m from command line



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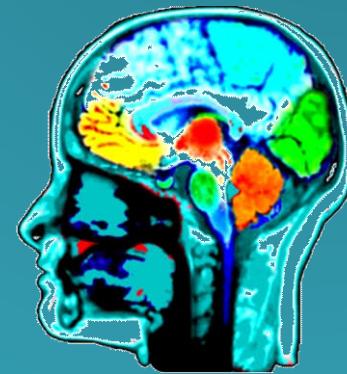


گروه تصویر پردازی و  
آلتایز تصاویر مغزی  
بیمارستان آمام خمینی  
تهران



# Workshop on Imaging and analysis of fMRI data

School of Advanced Technologies in Medicine  
13th Esfand 1399 - 12th Khordad 1400  
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## Software Introduction

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# Thanks