



ROAST: TES modeling made easy

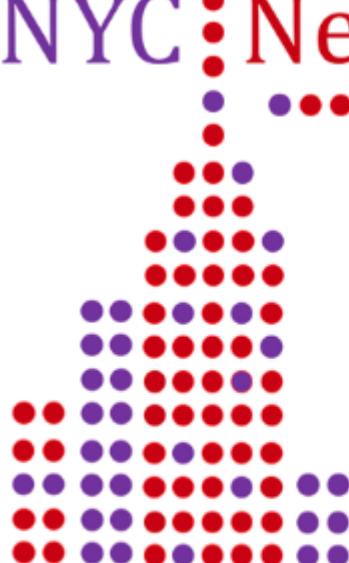
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A free, self-organized
community meeting



Agenda

- Talk & Demo (30 min)
- Q & A (30 min)

Quick links

- To get ROAST:

<https://www.parralab.org/roast/>

- Documentation:

<https://github.com/andypotatohy/roast>

- Mailing list:

roast-users@googlegroups.com

Overview

- `roast()` --- simulation
- `roast_target()` --- optimization / targeting
- `reviewRes()` --- review results
- Outputs of ROAST package
- Other issues

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Synopsis of roast()

- `roast(subj,recipe,varargin)`
- `subj` --- path/to/your/mri.nii
can be T1 or T2
use 'T2' option for T1+T2
- `recipe` --- montage for stimulation
{elec, current}
- `varargin` --- all the options
'Name', 'Value'

Examples – basic

- `roast('example/subject1.nii', {'F1',0.3,'P2',0.7,'C5',-0.6,'O2',-0.4},'simulationTag','basicDemo')`
- See capInfo.xls for all electrode layouts

Examples – cap type

- 'capType'

'1020' | '1010' (default) | '1005' | 'BioSemi' |
'EGI'

Examples – customized locations

- Use MRIcro to click for customized locations
- Record the voxel coordinates into a text file
- Call roast()

Examples – electrode shape

- 'elecType' -- the shape of electrode.

'disc' (default) | 'pad' | 'ring'

Examples – electrode size

- 'elecSize'
- disc: [radius height], default [6mm 2mm]
- pad: [length width height], default [50mm 30mm 3mm]
- ring: [innerRadius outerRadius height], default [4mm 6mm 2mm]

Examples – pad orientation

- 'elecOri' -- the orientation of pad electrode

'lr' (default) | 'ap' | 'si' | direction vector of the long axis

Examples – add T2

- 'T2'

[] (default) | file path to the T2 MRI

Examples – resampling and zero-padding

- 'resampling'
'on' | 'off' (default)
- 'zeroPadding'

Examples – mesh and conductivity control

- 'meshOptions'

meshOpt.distbound: default 0.3;

meshOpt.maxvol: default 10

- 'conductivities'

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Synopsis of roast_target()

- Generate lead field first:
`roast('example/MNI152_T1_1mm.nii','leadField','zeropadding',20,'simulationTag','MNI152leadField')`
- 74 electrodes placed (elec72.loc)
- Options frozen: 1010 system; disc electrode; 6mm radius, 2mm thick;
- `roast_target(subj,simTag,targetCoord,varargin)`
- subj --- same MRI file used for roast()
- simTag --- 'simulationTag' used when generating the lead field
- TargetCorod --- target locations in the brain
- varargin --- all the options
 'Name', 'Value'
- `roast_target('example/MNI152_T1_1mm.nii','MNI152leadField',[-48 -8 50],'targetingTag','basicDemo')`

Examples – split electrodes

- 'optType'

Max intensity: 'max-l1' (default) | 'max-l1per'

- 'elecNum'

Examples – intensity vs. focality

- 'optType'

Max intensity: 'max-l1' (default) | 'max-l1per'

Max focality: 'unconstrained-wls' | 'wls-l1' | 'wls-l1per' | 'unconstrained-lcmv' | 'lcmv-l1' | 'lcmv-l1per'

- 'k'

reduce 'k' to get more focality

increase 'k' to get more intensity

Examples – multi-focal targeting

- N-by-3 matrix
- Use 'wls-l1' with lower 'k' value

Examples – voxel coordinates

- Use MRIcro to click for voxel coordinates
- Call `roast_target()` with 'coordType','voxel'

Examples – define orientations

- 'orient'

'radial-in' (default) | 'radial-out' | 'right' | 'left' |
'anterior' | 'posterior' | 'right-anterior' | 'right-
posterior' | 'left-anterior' | 'left-posterior' |
'optimal' | orientation vector of your choice

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Synopsis of reviewRes()

- `reviewRes(subj,simTag,tissue,fastRender,tarTag)`
- `subj` --- path/to/your/mri.nii
- `simTag` --- tag of the simulation run in `roast()`
- `tissue` --- which tissue you want to visualize
- `fastRender` --- fast rendering or not
- `tarTag` --- tag of the targeting run in `roast_target()`

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Outputs of roast()

- subjName_roastLog
- Figures
- NifTI images: subjName_simulationTag_v.nii,
subjName_simulationTag_e.nii,
subjName_simulationTag_emag.nii
- Matlab file:
subjName_simulationTag_roastResult.mat
- Text files: subjName_simulationTag_v.pos,
subjName_simulationTag_e.pos

Outputs of roast_target()

- subjName_targetLog
- Figures
- Matlab file:
subjName_targetingTag_targetResult.mat

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Other issues

- Use of ‘New York Head’: memory hungry
- Use of RAS heads recommended
- Use of MNI coordinates recommended
- Use MRIcro instead of MRICron (my code showHeader() also good)

Original MRI

Original voxel space

To RAS;
Resample;
Zero-pad

Transformed MRI

Model voxel space

SPM computes

MNI (world) space

ROAST
operates
here (RAS)

Q & A