# RapidPT: A MATLAB toolbox for fast and efficient Permutation Tests

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http://felipegb94.github.io/RapidPT/

### The Project - Overview

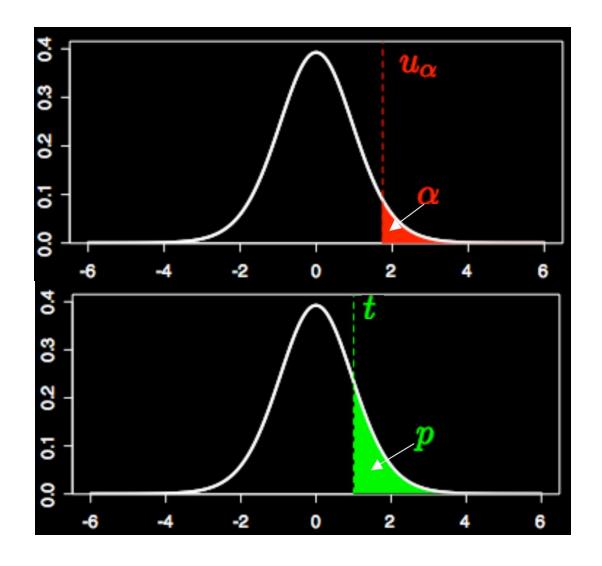
- RapidPT is a MATLAB toolbox for fast, scalable, and reliable permutation testing.
  - Fast/Scalable: Speedups of 20x 80x when compared to a simple permutation testing implementation. 2x 6x when compared to the state of the art, Statistical NonParametric Mapping (SnPM).
  - **Reliable:** Hundreds of validations runs against SnPM on neuroimaging datasets ranging from 20-400 subjects.
  - Easy to use: No setup, unlike hardware-based procedures. Will run fast on any modern machine.

## Hypothesis Testing

- Idea: Calculate the probability that your claim/hypothesis is true.
- Question: Does the data display any interesting activity?
  - In neuroimaging Voxel-wise differences between rest vs. activation? Healthy vs. Diseased?
- Two Possibilities:
  - Null Hypothesis vs. Alternate Hypothesis
  - H<sub>0</sub> vs. H<sub>a</sub>

### Hypothesis Testing - Procedure

- 1. Choose appropriate test statistic (t-test, mean difference, etc).
- 2. State  $\rm H_0$  and  $\rm H_a$
- 3. Construct the null distribution for the test statistic. This is the distribution of the statistic given that H<sub>0</sub> is true. This can be done analytically in some cases.
- 4. Compute the t-statistic with the given data.
- 5. Calculate the probability of observing such a statistic given that  $H_0$  is true (p-value).
- 6. Accept or reject  $H_0$ .



### Hypothesis Testing - Types of Error

#### **Actual Situation "Truth"** Ho True H<sub>o</sub> False **Incorrect Decision Correct Decision** Do Not Type II Error Reject Ho $1 - \alpha$ **Incorrect Decision Correct Decision** Rejct Ho Type I Error 1 - β $\alpha$

$$\alpha = P(Type\ I\ Error)$$
  $\beta = P(Type\ II\ Error)$ 

### Multiple Testing Problem

• FWER: Probability of making at least Type I Error (False Positive).

P(Making at least 1 error in m tests) = 1 - (1 - 
$$\alpha$$
)  
P(Making at least 1 error in m tests) = 1 - (1 -  $\alpha$ )<sup>m</sup>

- For  $m = 100 \rightarrow FWER = 0.99$ .
- For a functional neuroimaging dataset we would do >100,000 tests. One for every voxel.
- Hence, we want to somehow control for the FWER and keep it under a certain probability  $\alpha_0$ .

### Controlling FWER

#### Parametric Methods

- Assumptions about the data and its distribution.
- **Bonferroni** Conservative. Simply set  $\alpha_0 = \alpha/v$  (v = number of tests).
- Random Fields Theory Estimate number of activation areas, effectively reducing the number of tests.

### Nonparametric/Resampling Methods

- Permutation Testing Empirically estimate the null distribution of the test statistics.
  - Exact control of the FWER.

### **Permutation Testing**

• Idea: If the two groups do not differ, then I can permute the group/class labels and end up with approximately same set of t statistics.

#### Procedure:

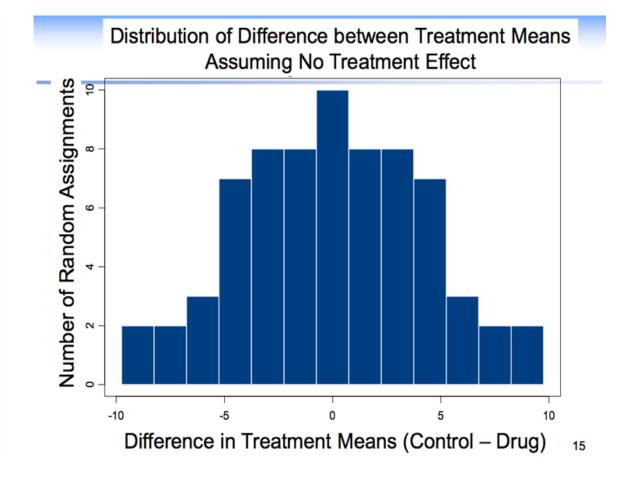
- 1. Re-label images (permute the labels of the images).
- 2. Compute test statistic for each voxel
- 3. Repeat N times.
- After the procedure is done we will have the exact null distribution for each voxel, and we can proceed from step 4 of hypothesis testing.

### Permutation Testing – Example, Single Test

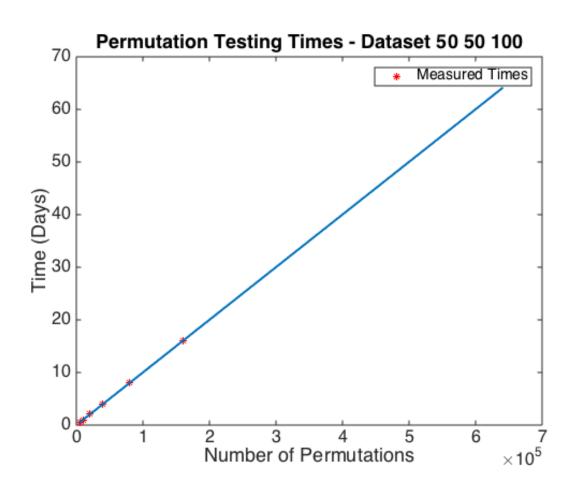
	Control				Drug			
Expression	9	12	14	17	18	21	23	26
Average	13				22			

Rearrangement of data Random Difference Assignment in Averages Drug 23 9.0 8.5 13 14 15 69 -8.5 14 70 -9.0

8 choose 4 = 70 possible permutations



### Computational Issues with Permutation Tests



#### **Memory Issues**

- Permutation Testing Matrix is very large
  - Let v be the number of voxels (around 5e5)
  - Let T be the number of permutations (10,000)
  - Let P be the permutation testing matrix (v x
     T)
  - Memory for that matrix is around 40 gigabytes
    - Note MATLAB crashes at some point when dealing with matrices that big.

### Motivation: We need a lot of permutations...

#### • 1 test

- 10,000 permutations  $\rightarrow$  The smallest possible p-value is  $\frac{1}{10,000}$ .
- Therefore the uncertainty near alpha = 0.01 is ±0.0001 OR ±1%

#### • 2 tests

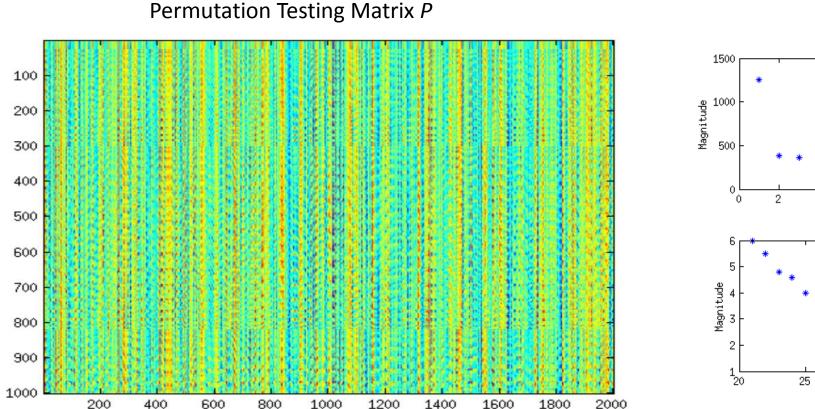
- 10,000 permutations  $\rightarrow$  The smallest possible p-value is  $\frac{1}{10,000}$  in each test.
- Now we have 2 opportunities for a type I error
- Therefore the uncertainty near alpha 0.01 in the worst case is ±0.0002 or 2%

#### • 500,000 tests

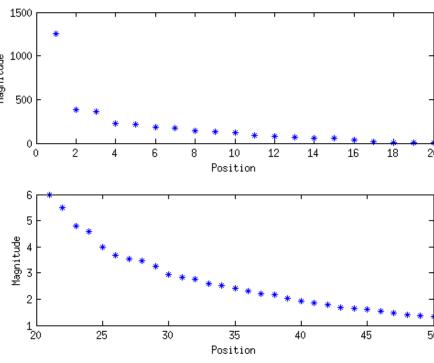
- 10,000 permutations  $\rightarrow$  The smallest possible p-value is  $\frac{1}{10,000}$  in each test.
- Therefore the uncertainty in the worst case scenario is 100%

### Idea: Look at the structure of P

- P is "highly structured" A combination of low-rank signal and high-rank residual
- Example: MRI data 100 healthy vs. non-healthy. v = 1,000, T = 2,000



#### Singular values of P



### Core of RapidPT

- Many columns in P look similar to othe columns as well as many rows look similar to other rows.
- If we compute a small number of entries of *P* we should be able to estimate the rest of it.
- Mathematically,

$$P = UW + S$$

- U is the low-rank basis of P.
- W is the coefficient matrix
- S is a high-rank random residual (noise).
- How many entries? In our experiments, subsampling <1% was enough</li>

### RapidPT Algorithm

#### Algorithm 3 Efficient Permutation Testing Algorithm

```
procedure RapidPT(data, nGroup1, numPermutations, numTrainingSamples, samplin-
   gRate)
   ▶ Training
       P\_exact \leftarrow PermTest(data, nGroup1, numTrainingSamples)
       U \leftarrow EstimateBasis(P\_exact)
       W \leftarrow EstimateCoeffMat(U, P\_exact)
       S \leftarrow ApproximateNoise(U, P\_exact)
   ▶ Recovery
       maxnull \leftarrow zeros(1, numPermutations)
 6:
       subset \leftarrow zeros(1, samplingRate * numVoxels)
       for i \leftarrow 1, numPermutations do
 8:
          subset \leftarrow randsample(1:numVoxels, samplingRate*numVoxels)
          p \leftarrow PermTest(Data(:, subset), nGroup1, 1)
10:
          w \leftarrow EstimateCoeffMat(U(:,subU),p)
          s \leftarrow EstimateNoise(U(:, subU), p)
12:
          p \leftarrow U * w + s
          maxnull(1, i) \leftarrow max(p)
14:
       end for
       return maxnull
                                                                      ▶ Maximum Null Distribution
16:
   end procedure
```

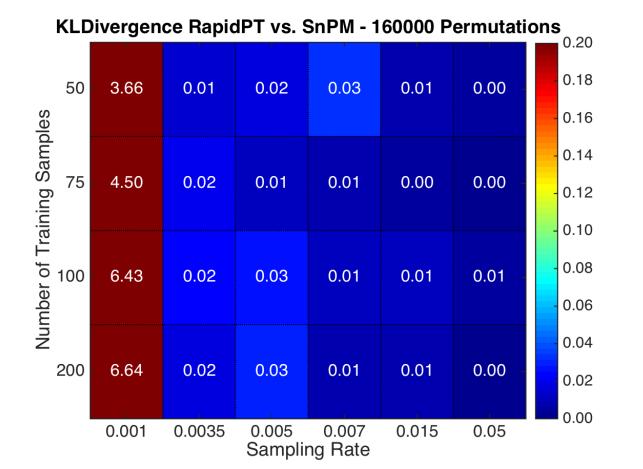
### **Evaluations Setup**

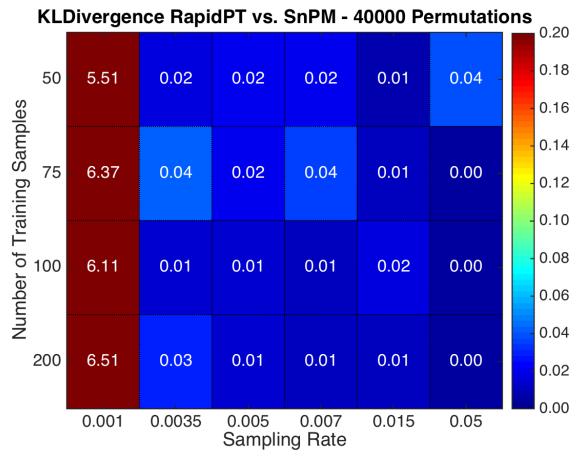
- Data: T1 MRIs from ADNI2 are used.
  - 601 subjects (259 AD and 342 CN)
  - SPM preprocessing is applied.
  - GM images with approx. 500,000+ voxels are extracted.
  - Multiple combinations of dataset sizes
- Experiments: Can we recover the Maxnull distribution?
  - Stability of hyperparameters Sub-sampling rates, and training samples
  - Computational Speedups (RapidPT vs. SnPM, RapidPT vs. NaivePT)

### Recovered MaxNull Distribution Accuracy

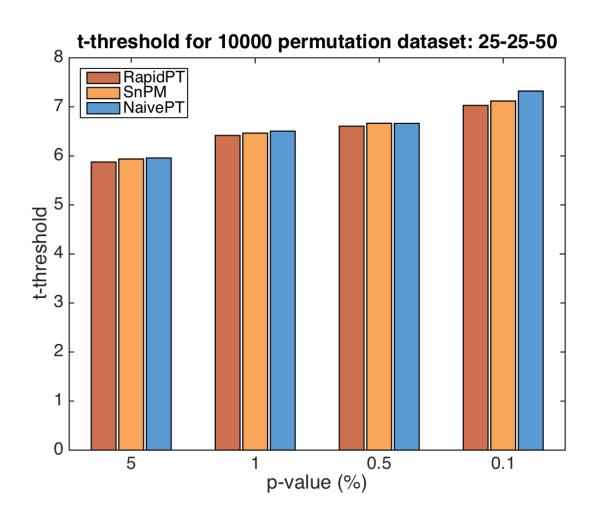
KLDivergence: Measure of the difference between two distributions

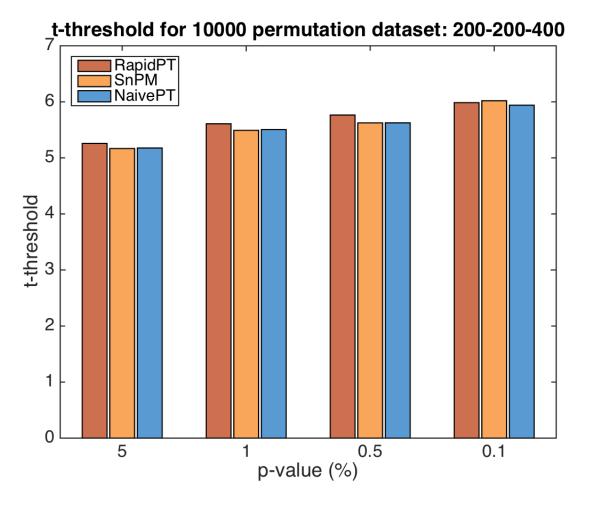
Dataset: 100 subject, 50 AD 50 CN



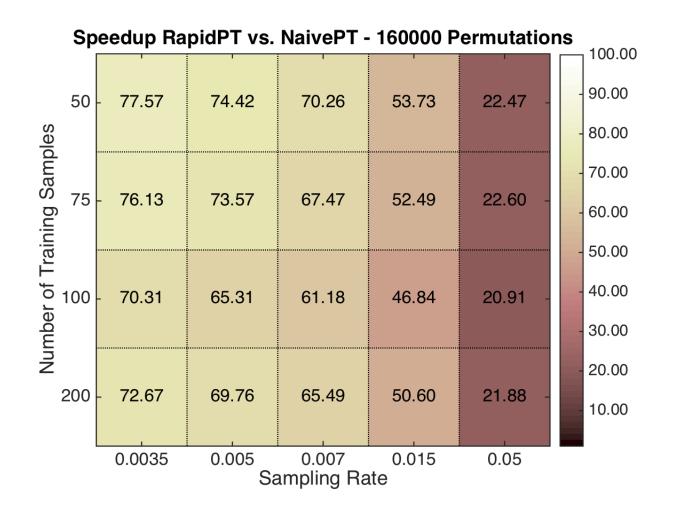


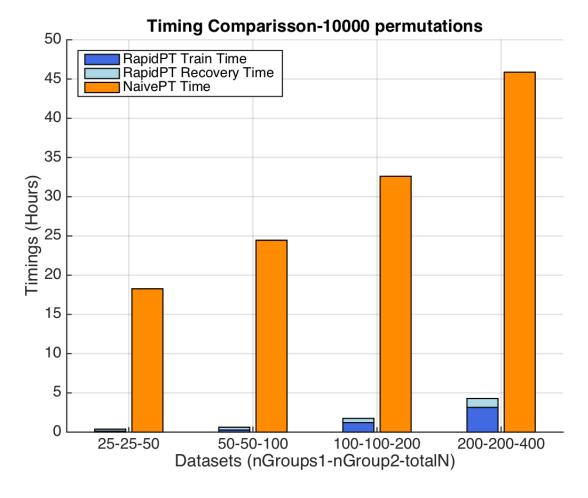
### Calculated T-Threshold for a Given P-Value





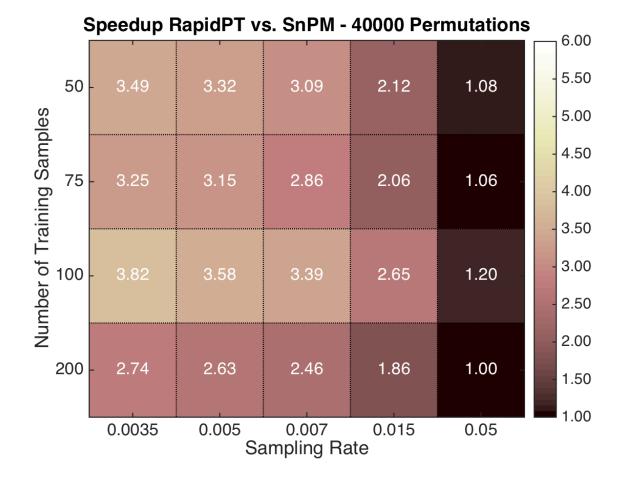
### Speedup RapidPT vs. NaivePT

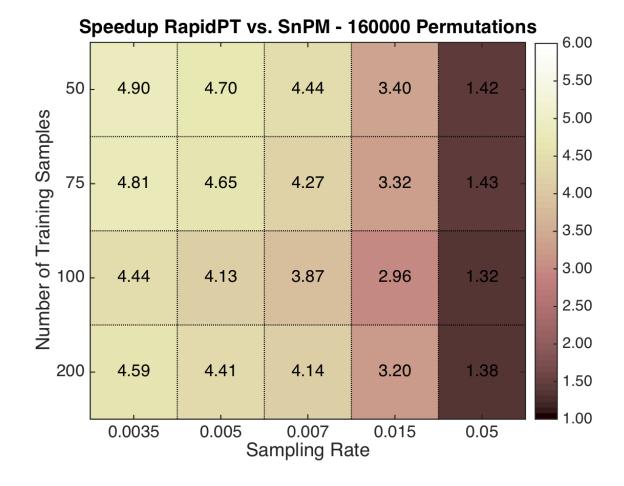




### Speedup RapidPT vs. SnPM

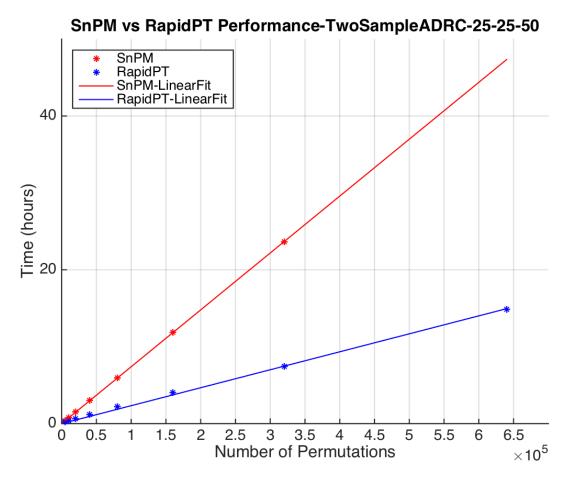
Dataset used: 100 subjects, 50 AD 50 CN

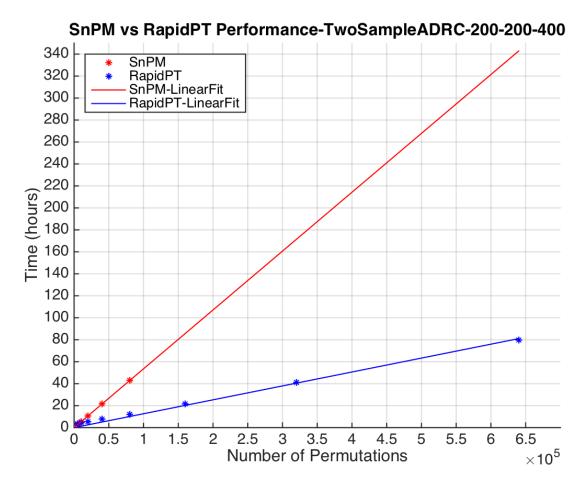




# Scaling analysis RapidPT vs. SnPM

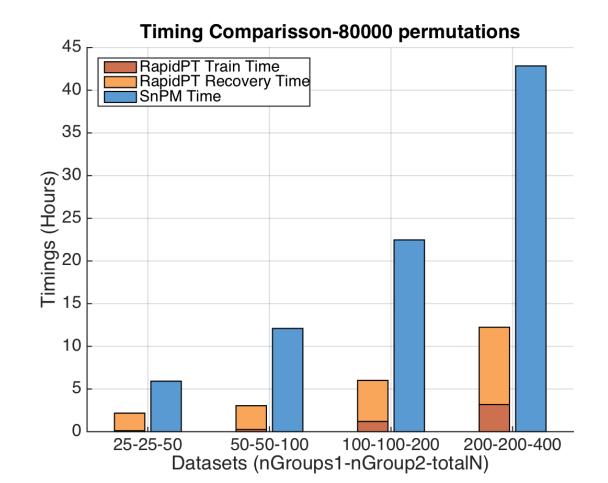
Putting the speedups into context...

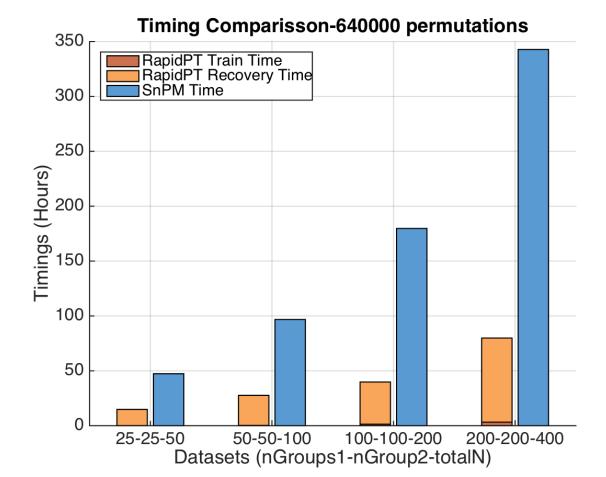




# Timings RapidPT vs. SnPM

Putting the speedups into context...





### Discussion

- RapidPT achieves state of the art performance
  - It is able to recover the Maximum null distribution to a high degree of accuracy. Good control for FWER.
  - Fast A few hours vs. days, and a day or two vs. a week or two...
  - Scalable As the number of permutations increases and the dataset size increases the speedups against SnPM and NaivePT improve...
  - Easy to use????

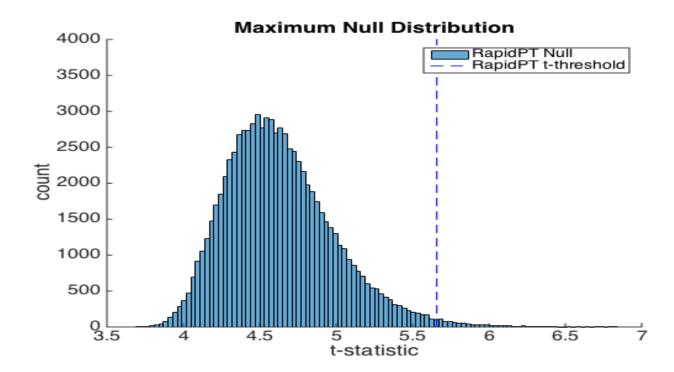
### RapidPT - Usage

Do pre-processing of .nii images for each subject and construct an NxV data matrix.

```
% Addpath RapidPT Repository Path (current working dir in this case)
RapidPTLibraryPath = '.';
addpath(RapidPTLibraryPath);
% Load input data and input labels
dataPath = '~/PermTest/data/ADRC/TwoSample/ADRC 50 25 25.mat';
% dataPath = '~/PermTest/data/face/Data face.mat';
load(dataPath):
% N subjects, V voxels (or statistics)
[N,V] = size(Data);
numPermutations = 40000;
nGroup1 = 25; % You should what is the size of one of your groups prior.
% Set write to 1 if you want the matrices used to recover the permutation matrix.
% Setting this to 1 will make outputs a very large variable, but may be
% useful in certain cases.
write = 0;
[outputs, timings] = TwoSampleRapidPT(Data, numPermutations, nGroup1, write, RapidPTLibraryPath);
save(strcat('outputs/outputs_TwoSampleFace_',num2str(numPermutations),'.mat|'),'outputs');
save(strcat('timings/timings_TwoSampleFace_',num2str(numPermutations),'.mat|'),'timings');
```

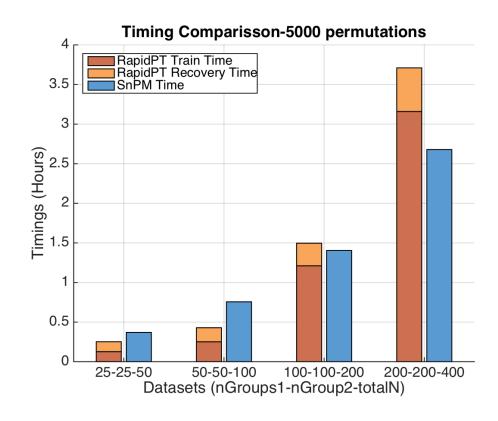
### RapidPT – Postprocess Example

```
% Get the outputs struct you obtained from RapidPT
load('~/PermTest/outputs/TwoSample_ADRC_200_200_400/rapidpt/outputs_80000_0
alpha = 0.01; % Significance level of 1 percent
tThresh_RapidPT = prctile(outputs.maxT, 100 - (100*alpha));
% Get the data
load('~/PermTest/data/ADRC/TwoSample/ADRC_400_200_200.mat');
[h,p,ci,stats]=ttest2(Data(1:200,:),Data(201:400,:),0.05,'both','unequal');
SampleMaxT = max(stats.tstat);
```



# Warnings for RapidPT and Permutation Testing

- Small datasets and few permutations
  - A couple minutes vs one minute
  - Large uncertainty due to the smallest possible p-value being large.
    - 10 subjects  $\rightarrow$  10 choose 5 = 252
    - Smallest possible p-val = 1/252.



### Acknoledgements and Website

- Vikas Singh
- Vamsi Ithapu
- ADNI and ADRC

- Repository and project website:
  - https://github.com/felipegb94/RapidPT (Repository)
  - http://felipegb94.github.io/RapidPT/ (Website)

### More Motivation: Implementation

- Implementation can easily be made inneficient
  - Inneficient data accesses
  - Not take advantage of MATLAB's optimized features

Algorithm 1 Traditional Two-Sample Permutation Testing. In this implementation ttest2 refers to the two-sample t-test operation described in equation 1.

```
1: procedure TWOSAMPLENAIVEPT(\mathbf{X}, r, N_1, N_2)
2: for i \leftarrow 1, r do
3: shuffleRows(\mathbf{X})
4: \mathbf{X}_1 \leftarrow \mathbf{X}(1:N_1,:)
5: \mathbf{X}_2 \leftarrow \mathbf{X}(N_1+1:end,:)
6: \mathbf{t} \leftarrow ttest2(\mathbf{X}_1, \mathbf{X}_2)
7: \mathbf{T}(i,:) \leftarrow \mathbf{t}
8: end for \triangleright Mean and Variance Calculation
9: end procedure
```

