

Magic Pie (mgcpy)

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Date: 9/12/2018

Sprint 1: Create mgcpy package

- Task 1: Implement dHSIC, HHG, Pearson, Spearman, and Mic into package (Sambit)
- Task 2: Implement MGC into the package (Satish)
- Task 3: Implement MDMR and FastMGC into package (Sandhya)
- Task 4: Implement MCORR, DCORR, and Mantel into package (Bear)
- Task 5: Implement Random Forest independence tests into package (Richard)
- Task 6: Implement 2- sample tests into package (Ananya)

Task 1 (Sambit): Implement dHSIC, HHG, Pearson, Spearman, and Mic into package

- Translated Pearson's correlation data (RVCorr.m function in fastMGC) into python and sent pull request to development branch (pending approval)
- Tested code for the function (works for both lower dimensional and high dimensional tests)

Function is able to calculate correlations between distance matrices.

Matrix A

0	23	56	90	5	63	49
23	0	80	15	95	4	43
56	80	0	94	27	41	90
90	15	94	0	95	95	89
5	95	27	95	0	35	37

1. Compare A, A: Correlation = 1.0

Function is able to calculate correlations between distance matrices.

Matrix A

Matrix B

0	23	56	90	5	63	49	0	18	5	3	6	75	72	
23	0	80	15	95	4	43	18	0	75	46	50	6	41	
56	80	0	94	27	41	90	5	75	0	35	31	44	92	
90	15	94	0	95	95	89	3	46	35	0	87	62	55	
5	95	27	95	0	35	37	6	50	31	87	0	57	55	

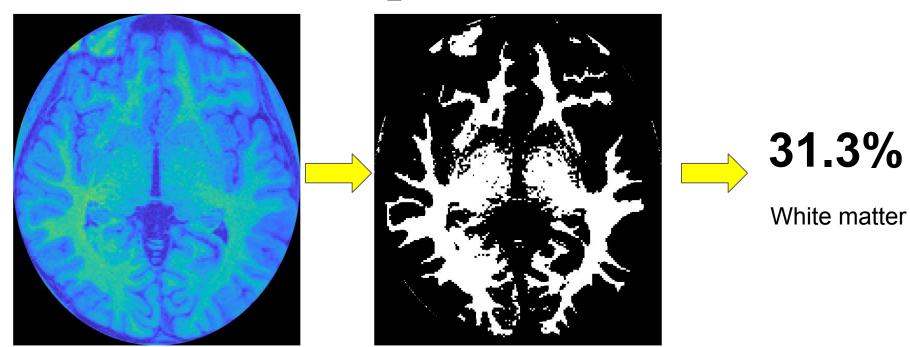
0	18	5	3	6	75	72
18	0	75	46	50	6	41
5	75	0	35	31	44	92
3	46	35	0	87	62	55
6	50	31	87	0	57	55

2. Compare A, B: **Correlation =** 0.54479

Visualization of the week

Cropped Section

sub-NDARAA536PTU_T1w.nii File



Filtered White Matter

Stuff to do this week

- More robust testing on higher dimensional data sets for RVCorr
- Begin working on dHSIC function (little bit more involved)
- Finalize PR with RVCorr function and merge with development branch
- Make RVCorr compliant with documentation guidelines

Task 2 (Satish): Implement MGC into the package

Last Week Accomplishments:

- Added the <u>Apache 2.0</u> LICENSE file (and the batch to <u>README.md</u>)
- Added <u>pre-commit</u> git hooks to ensure PEP8 style guide is followed
 - Added a PEP8 badge as well
- Added draft version of the contribution guidelines (<u>CONTRIBUTING.md</u>)
- Setup code coverage (<u>coveralls</u>) and added a badge
- Implemented the DistRanks (in R MGC) method with unit test in mgcpy

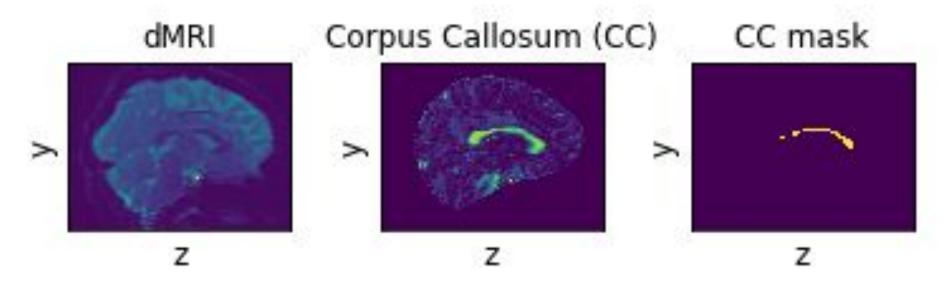
All the changes above have been reviewed & merged to the *development* branch.

```
Sorts the entries within each column in ascending order
> DistRanks <- function(dis) {</pre>
                                                             For ties, the "minimum" ranking is used, e.g. if there are
       n=nrow(dis)
                                                             repeating distance entries, The order is like 1,2,2,3,3,4,...
       disRank=matrix(0,n,n)
                                                             :param distance_matrix: a symmetric distance matrix.
       for (i in (1:n)){
                                                             :return: column-wise ranked matrice of ``distance_matrix``
            v=dis[,i]
            tmp=rank(v,ties.method="min")
                                                             n_rows = distance_matrix.shape[0]
                                                             ranked distance matrix = np.zeros(distance matrix.shape)
            tu=unique(tmp)
                                                             for i in range(n_rows):
            if (length(tu)!=max(tmp)){
                                                                 column = distance_matrix[:, i]
                tu=sort(tu)
                                                                 ranked column = np.array(scipy.stats.rankdata(column, "min"))
                 for (j in (1:length(tu))){
                                                                 sorted_unique_ranked_column = sorted(list(set(ranked_column)))
                      kk=which(tmp==tu[j])
                                                                 if (len(ranked_column) != len(sorted_unique_ranked_column)):
                                                                    for j, rank in enumerate(sorted_unique_ranked_column):
                     tmp[kk]=j
                                                                        ranked column[ranked column == rank] = j + 1
                                                                 ranked distance matrix[:, i] = ranked column
                                                             return ranked_distance_matrix
            disRank[,i]=tmp
                                                         def test rank distance matrix():
                                                             a = np.array([[1, 4, 6],
       return(disRank)
                                                                             [2, 5, 7],
                                                                             [1, 4, 6]])
                                                             ranked_a = np.array([[1, 1, 1],
                                                                                    [2, 2, 2],
                                                                                    [1, 1, 1]])
                                                             assert np.array equal(rank distance matrix(a), ranked a)
```

def rank_distance_matrix(distance_matrix):

Visualization of the week

Extracting the Corpus Callosum of Subject-NDARBN100LCD (using *dipy*)



Stuff to do this week

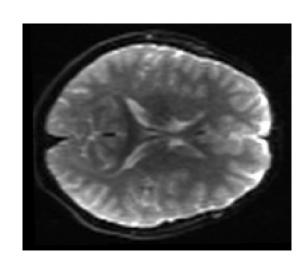
- Study the pseudo code in the MGC paper and the R code, and list down all the functions and its dependencies required to port to Python
- Create stubbed versions of all the above functions and add proper docstrings to define the inputs and outputs
- Implement one of the above function with tests in Python and raise a PR

Task 3 (Sandhya): Implement MDMR and FastMGC into package

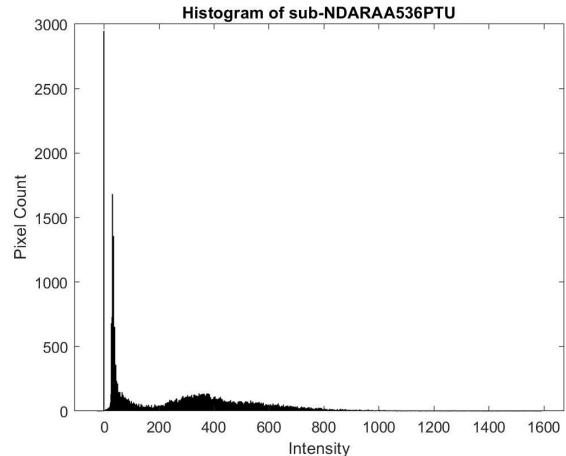
Last Week's Accomplishments:

- Attempted to run cython version of MDMR
 - Hard on windows: visual studios build issue, vcvarsall.bat
 - o In general more accessible if not in cython
- Began to create a python version of MDMR (previously cython)
 - All base functions running in python
 - Mdmr function runs, but unclear input "columns"
- Attempted to contact author- no response yet
- Read up on MDMR from R edition's documentation

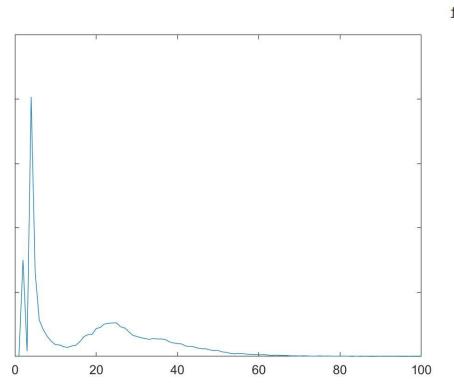
Visualization of the Week



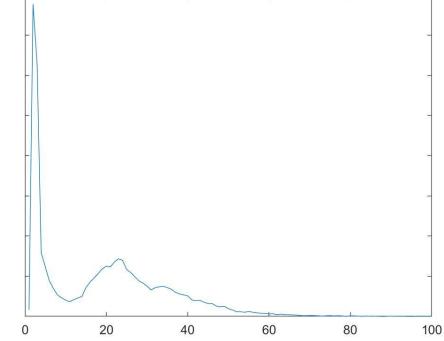
- sub-NDARAA536PTU
- Registered DWI



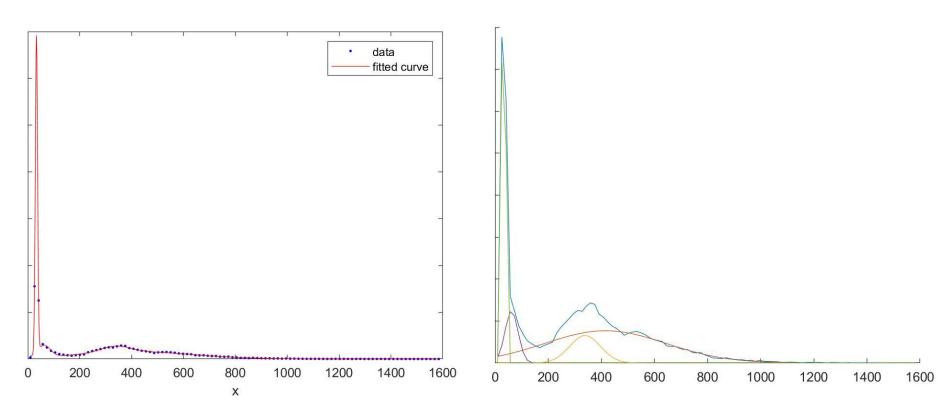
Background Strip:



```
load('a.mat')
a = double(a(a>0));
[p,center] = hist(double(a(a>min(a(:)))),100);
p = p/sum(p);
figure; plot(p);
fitobject = fit(center',p','gauss4');
figure; plot(fitobject, center', p');
```



Gaussian Mixture Model



Next week:

- Convert remaining MDMR code to python
- Contact author of MDMR code for help
- If successful, run MDMR on spiral data, linear data
- If successful, convert MDMR code to our coding structure/documentation
 - o Continue learning how git works and what our coding structure is from my teammates/google

Task 4 (Bear): Implement MCORR, DCORR, and Mantel into package

Last Week's Accomplishments:

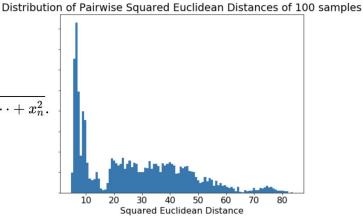
- Implemented DCORR. Results matched matlab code on simulated data
 - Specifics of simulated data: all 20 types of dependence, X, Y are 100 examples and 1 dimensional
 - DCORR test statistics matched original results with less than 1e-5 absolute error
- Initiated connectome preprocessing pipeline
 - n Edgelists → n d by d Adjacency matrices → one n by p data matrix (p is the dimension of each example)
 - Most naive approach: flatten the matrices!
 - 48 x 48 adjacency matrix → 2304-dimensional vector

Pairwise distances between connectome graphs

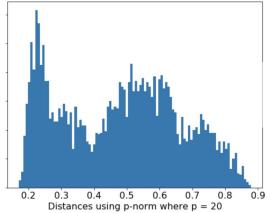
• 100 "task-rest_bold_JHU_res-2x2x2_measure-correlation.edgelist" examples

Distribution of Pairwise Euclidean Distances of 100 samples

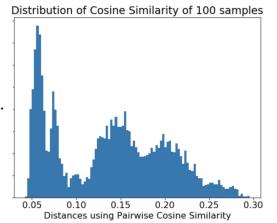
 $\|oldsymbol{x}\|_2 := \sqrt{x_1^2 + \dots + x_n^2}.$



Euclidean Distance
Distribution of Pairwise 20-norm Distances of 100 samples



 $\left\|\mathbf{x}
ight\|_{p}:=\left(\sum_{i=1}^{n}\left|x_{i}
ight|^{p}
ight)^{1/p}$



 $1-rac{u\cdot v}{||u||_2||v||_2}$

Next week

- Merge DCORR, MCORR, Mantel modules (with unit tests) into mgcpy
 - DoD: Closed PR
- Potential new tasks
 - Port simulation code into mgcpy
 - DoD: Scatter plots of different dependency
 - Build connectome-to-correlation-test data pipeline
 - DoD: Module which converts edgelist files into formats usable for mgcpy functions

Task 5 (Richard): Random Forest Based Independence Test

Last Week's Accomplishments:

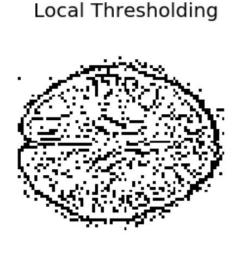
- Setup travis-ci (merged into dev)
- Setup sphinx for documentation (PR ready)

Visualization of the week:

fMRI of Subject NDARMR277TT7 Watching Despicable Me

Original





Next week

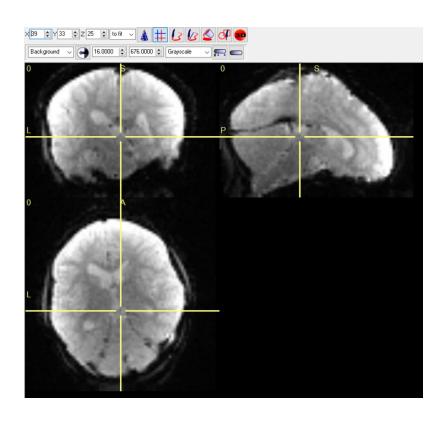
- Detailed guide on how to generate documentation and overall github workflow
 - DoD: docs in sphinx
- Pseudocode on using random forest to estimate conditional entropy
 - DoD: LaTeX write up
- Reach: Implement algorithm using scikit learn random forest embedding code
 - DoD: Jupyter Notebook showing results on easy simulated dataset

Task 6 (Ananya): Implement 2-sample tests into package

Last Week's Accomplishments:

 Read and took notes on "Equivalence of Distance and Kernel Methods" to learn more about ENERGY and MMD

Visualization of the Week

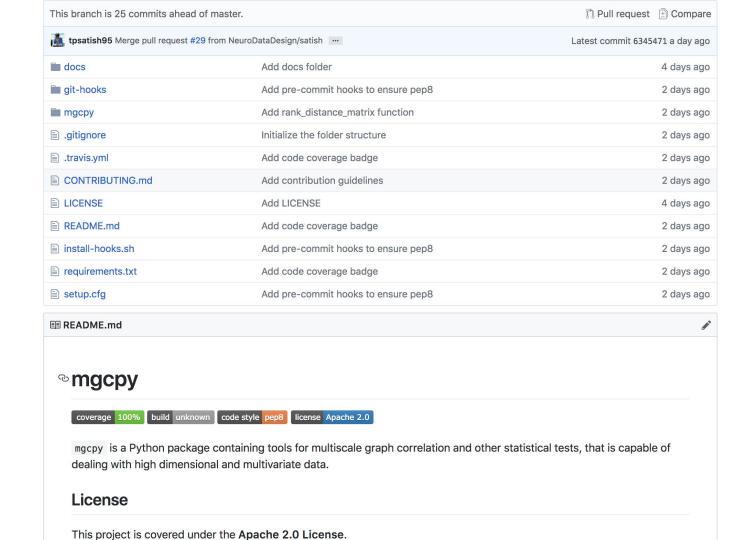


Resting state fMRI of subject NDARMN450PEH

TP = 12

Next week

- Run ENERGY
 - o DoD: Output of ENERGY
- Run MMD
 - o DoD: Output of MMD



Collective Team Task:

Last week: Completed first draft of AWS proposal

This week: Visualization of data, Better second draft of AWS proposal, Make PR on something in repo