Data Analytics - Assignment IV

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Effect of Smoking

Part 1

To identify genes which respond differently to smoke in men vs. women (Smoking Status X Gender model vs. Smoking Status + Gender null model) We computed A and A' using :

$$h = AB + Error$$

h: gene expression

A: Alternative Hypothesis

A': Null Hypothesis

B: mean vector

For Null Model A':

For Alternative Model A:

$$\begin{bmatrix} h_1 \\ h_2 \\ \vdots \\ \vdots \\ h_{48} \end{bmatrix} = \begin{bmatrix} 1 & 0 & 0 & 0 \\ 1 & 0 & 0 & 0 \\ \vdots & & & & \\ \vdots & & & & \\ 0 & 0 & 0 & 1 \end{bmatrix} \begin{bmatrix} male_{nonsmoker} \\ male_{smoker} \\ female_{nonsmoker} \\ female_{smoker} \end{bmatrix} + \begin{bmatrix} \epsilon_1 \\ \epsilon_2 \\ \vdots \\ \epsilon_{48} \end{bmatrix}$$

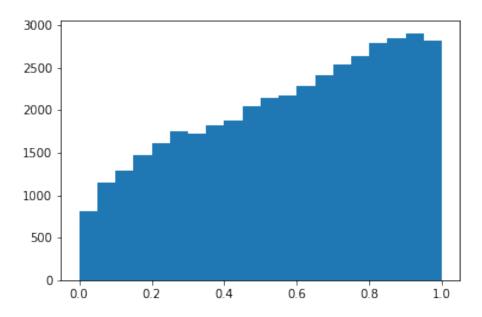
I had computed F-Statistic by using formula:

F-statistic
$$\hat{f} = \frac{\vec{\hat{h}}^T (A(A^TA)^{\dagger}A^T - A'(A'^TA')^{\dagger}A'^T)\vec{\hat{h}}}{\vec{\hat{h}}^T (I - (A(A^TA)^{\dagger}A^T)\vec{\hat{h}}} * \frac{n - rank(A)}{rank(A) - rank(A')}$$

After this i had computed p-value by using scipy library (refer code part)

Part 2

To draw the histogram of p-values



Part 3
Refer code

Part 4

Refer code

Part 5

Refer genes-symbol-list.txt file

Part 6

Genes Intersection with Xenobiotic metabolism :: ['SULT1A1', 'AOC2', 'CYP2S1', 'AADAC', 'HNF4A', 'AS3MT']

Genes Intersection with Free Radical Response:: None

Genes Intersection with DNA Repair :: [PNKP]

Genes Intersection with Natural Killer Cell Cytotoxicity :: ['IFNG', 'KLRC2', 'PTPN6', 'HLA-C', 'PRF1', 'HLA-E', 'HLA-G']

Part 7

Intersection Count:

Xenobiotic metabolism :: 6 Free Radical Response :: 0

DNA Repair :: 1

Natural Killer Cell Cytotoxicity:: 7

Groups are as follows:

Female Smokers up genes ::

['HLA-C', 'CYP2S1', 'PNKP', 'HLA-E', 'SULT1A1', 'AOC2', 'HLA-G', 'HNF4A', 'PTPN6']

Female Smokers Down genes ::

['KLRC2', 'IFNG', 'AADAC', 'PRF1', 'AS3MT']

Male Smokers up genes::

['KLRC2', 'IFNG', 'AADAC', 'HLA-G', 'PRF1', 'AS3MT', 'HNF4A', 'HLA-E']

Male Smokers down genes::

['HLA-C', 'CYP2S1', 'PNKP', 'HLA-E', 'SULT1A1', 'AOC2', 'HLA-G', 'HNF4A', 'PTPN6']

Here is the detailed table of how i had splitted in 4 Groups :

		Xenobiotic metabolism				
Probe-Name	Gene-Symbol	Male Non Smoker	Male Smoker	Female Non Smoker	Female Smoker	
A_24_P10751	HNF4A	3.180579583	4.063394983	3.663447817	3.933650767	
A_23_P434212	SULT1A1	11.3759045	10.8529605	10.68061133	10.89170938	
A_23_P4133	AOC2	5.1779153	4.923473625	4.634595858	5.342885492	
A_23_P101374	CYP2S1	7.079202892	6.751945342	6.362079875	6.9155088	
A_23_P80570	AADAC	0.631956892	2.254574383	1.525998573	1.273477925	
A_32_P169688	HNF4A	0.805925633	0.526938294	0.352009458	2.103507458	
A_23_P12643	AS3MT	2.470355533	3.834752258	3.629293979	2.58933331	
A_23_P28761	HNF4A	0.7819002	0.915570567	0	0.569839333	
			Going down from Nor			
			Going up from Non Sn			
			Going up from Non Sn	noker to Smoker		

			DNA Repair		
Probe-Name	Gene-Symbol	Male Non Smoker	Male Smoker	Female Non Smoker	Female Smoker
A_23_P164883	PNKP	8.52977525	8.27714125	8.020962417	8.2652624
			Going down from Nor		
			Going up from Non Sr		

		Natural Killer Ce	ell Cytotoxicity		
Probe-Name	Gene-Symbol	Male Non Smoker	Male Smoker	Female Non Smoker	Female Smoker
A_23_P151294	IFNG	5.915514025	6.710654067	6.606077317	5.854469467
A_23_P22232	KLRC2	9.717394458	10.67619267	10.30146854	9.54930685
A_23_P113716	HLA-C	18.47654708	18.442839	18.48375175	18.57580192
A_24_P936272	HLA-C	15.18613767	14.98331017	14.90501925	15.30503171
A_23_P162486	PTPN6	12.29216738	11.98547275	11.93286204	12.23317408
A_23_P70539	HLA-C	16.35341483	16.180303	16.02259142	16.37197258
A_23_P95917	HLA-C	17.98420892	17.97244417	17.96767792	18.20373358
A_23_P1473	PRF1	10.366631	10.68289221	10.01158479	9.422278417
A_24_P298409	HLA-C	17.93432892	17.87332817	17.88560383	18.0351685
A_24_P326082	HLA-E	17.694454	17.70754658	17.5425235	17.80270308
A_23_P30848	HLA-E	16.18347967	15.99989933	15.82391817	16.11673117
A_24_P311926	HLA-G	18.459436	18.45459983	18.42325108	18.55368683
A_23_P300112	HLA-G	10.004409	10.22921967	10.07251369	10.39354538
A_23_P361614	HLA-G	7.697568258	7.559727542	7.030861325	7.784258217
A_32_P460973	HLA-E	17.21206008	17.18141983	17.09641017	17.33577933
			Going down from Non		
			Going up from Non Sn		