

BIO247 Exam 1

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Part 1 - Written

Problem 1

Being robust is being able to change any variables but the experiment is still able to work. Robustness is not definitive and can be narrow or wide depending upon how transferable the experiment guidelines are. Sharing code and commenting on it to make it readable for humans is also part of being robust.

Being reproducible means that the experiment has clearly described protocol that can be easily followed and conducted by others. In order to be reproducible, one should share their script used or at least supply a benchmark to reference as well. Clearly stating how things were done and why they were makes an experiment reproducible.

Problem 2

This problem has been drawn on the last page of the exam.

Problem 3

This conservation study is not reproducible because it does not give a clear procedure and is vague in its description. Also, the experiment itself is not sound due to the fact that they're asking a hypothetical question on how someone would behave in a situation, but their response could be far from the truth had the participants actually encountered the snake. As for robustness, one could argue that this is not robust because the age group of participants is not wide and is in fact limited to only undergraduate students. This experimnt then is very limited in its reach.

Problem 4

This bioinformatics study is robust and reproducible. It is robust and reproducible becasue it uses data accessed from open-source technologies in R programming languages which means that data for this can be easily (for the most part) accessed by others who wish to do this with different variables. Going off of that, the variables can easily be changed to something else and we could get viable results. They included where they got all of their data from and what programming they used for steps. This study includes well laid out steps that could be followed by others and apply under different variables.

Part 2 - Practical

Problem 5

This is a coding diagram beneath the chunk because I am unsure of the code to conduct this.

```
interact <- rep("interactions",25)
```

Use If/Else Loop to say if interactions are positive/negative and then print the suffix

Find the positive and negatives

Vectorize the suffixes that are negative

Vectorize the suffixes that are positive

Use the paste function

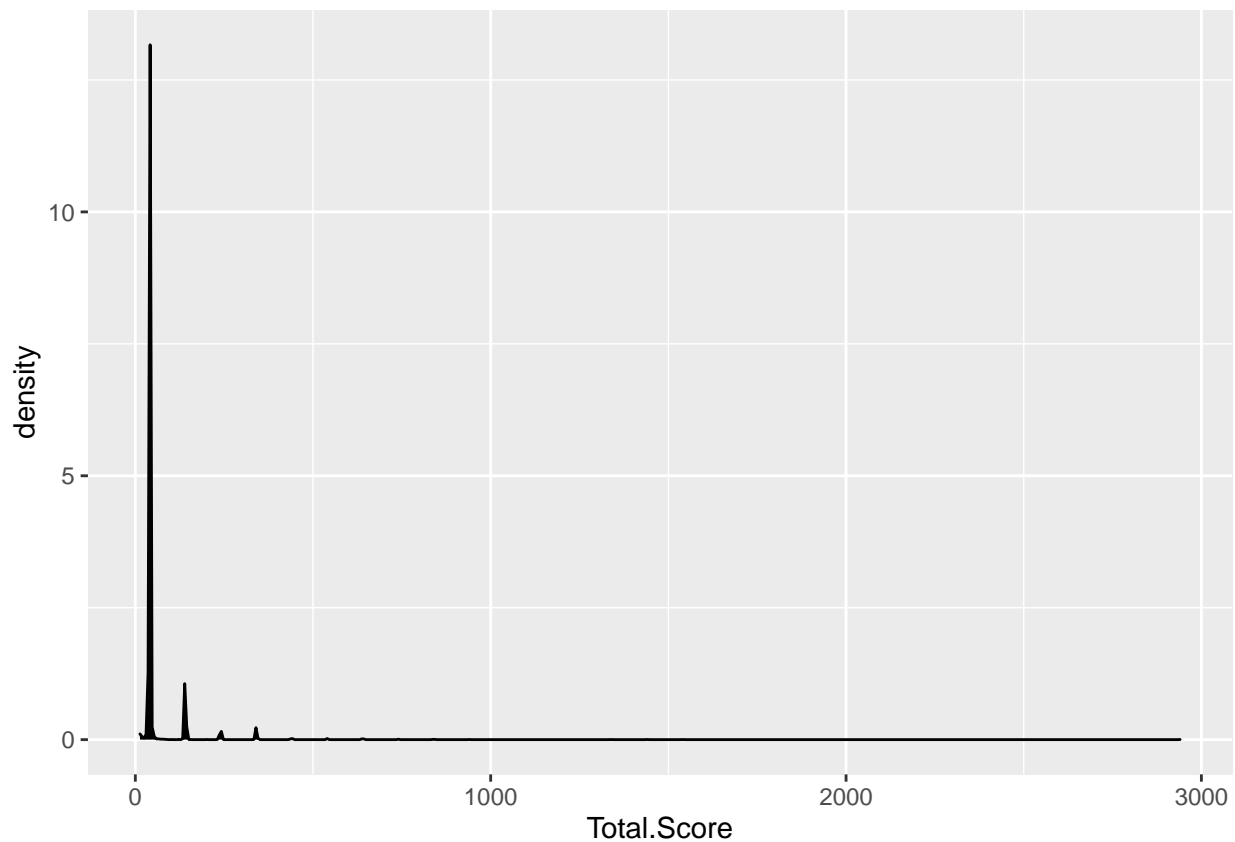
Problem 6

This is importing the dataset and installing the ggplot package in order to create a density plot.

```
GBM_data <- read.csv("/cloud/project/GBM_data.csv")
install.packages("ggplot2")
```

```
## Installing package into '/cloud/lib/x86_64-pc-linux-gnu-library/4.2'
## (as 'lib' is unspecified)
```

```
library(ggplot2)
ggplot(GBM_data)+geom_density(aes(x=Total.Score),fill="black")
```



Problem 7

subsetting the rows to the top 100 interactions only.

```
GBM_data[1:100,]
```

##	X	Element.Name	Element.Type	Database	PosReg.Name
## 1	2184	egfr	protein	p00533	pm

## 2	2188	egfr	protein	p00533	sd
## 3	5045	rb			p38
## 4	4989	ras	family	pf00071	
## 5	4994	ras	family	pf00071	
## 6	201	akt	family	akt	
## 7	2174	egfr	protein	p00533	ano1
## 8	3930	mrn	protein	o60934	
## 9	3661	map	protein	p12872	
## 10	2293	erk	family	erk	
## 11	203	akt	family	akt	
## 12	211	akt	family	akt	atra
## 13	177	akt	family	akt	
## 14	242	akt	family	akt	p110
## 15	204	akt	family	akt	
## 16	4996	ras	family	pf00071	
## 17	2094	e2f1	protein	q01094	
## 18	4285	p53	protein	p04637	
## 19	4291	p53	protein	p04637	
## 20	144	akt			
## 21	2307	erk	family	erk	insulin
## 22	4625	pi3k	protein	pi3k	insulin
## 23	4200	p21	protein	p38936	
## 24	2312	erk	family	erk	pmek
## 25	4288	p53	protein	p04637	
## 26	1107	caspase-9	protein	p55211	cytochrome c
## 27	6032	vegf	protein	p15692	vgpcr
## 28	3663	map	protein	p12872	integrin
## 29	3714	mdm2	protein	q00987	
## 30	4306	p53	protein	p04637	etv1
## 31	4634	pi3k	protein	pi3k	rtks
## 32	2346	es	simple-chemical	6995653	
## 33	4322	p53	protein	p04637	tert
## 34	5005	ras	family	pf00071	grb10
## 35	1142	cd44	protein	p16070	
## 36	3723	mdm2	protein	q00987	daxx
## 37	3664	map	protein	p12872	integrin-
## 38	3715	mdm2	protein	q00987	
## 39	4992	ras	family	pf00071	
## 40	4275	p53	protein	p04637	
## 41	3980	mtor	protein	p42345	vgpcr
## 42	212	akt	family	akt	bm11
## 43	4635	pi3k	protein	pi3k	runx2
## 44	2169	egfr	protein	p00533	
## 45	219	akt	family	akt	doxorubicin
## 46	186	akt	family	akt	
## 47	196	akt	family	akt	
## 48	195	akt	family	akt	
## 49	4609	pi3k	protein	pi3k	atra
## 50	4919	pten	protein	p60484	pi3k inhibitors
## 51	3972	mtor	protein	p42345	
## 52	146	akt			
## 53	150	akt			
## 54	151	akt			
## 55	157	akt			bdnf

## 56	161	akt			igf-i	
## 57	163	akt			il-3	
## 58	181	akt	family	akt		
## 59	185	akt	family	akt		
## 60	225	akt	family	akt	growth factor receptor	
## 61	187	akt	family	akt		
## 62	4626	pi3k	protein	pi3k	integrin	
## 63	4632	pi3k	protein	pi3k	rac1	
## 64	224	akt	family	akt	fbs	
## 65	253	akt	family	akt	resveratrol	
## 66	228	akt	family	akt	hdl	
## 67	4641	pi3k	protein	pi3k	tsc	
## 68	4223	p27	protein	p40305		
## 69	5056	rb	protein	p06400		
## 70	5059	rb	protein	p06400		
## 71	6022	vegf	protein	p15692		
## 72	4905	pten	protein	p60484		
## 73	4902	pten	protein	p60484		
## 74	5120	rb1	protein	p06400		
## 75	4409	parp1	protein	p09874		
## 76	5333	shc	protein	p29353	t antigen	
## 77	2708	fkhr	protein	q12778		
## 78	2756	foxo1a	protein	q12778		
## 79	5168	rheb	protein	q15382	insulin	
## 80	5261	s6k1	protein	p23443		
## 81	5185	ros	simple-chemical	chebi:26523		
## 82	5940	tuberin	protein	p49815		
## 83	5914	tsc2	protein	p49815		
## 84	5194	ros	simple-chemical	chebi:26523	selenite	
## 85	4224	p27	protein	p40305		
## 86	5054	rb	protein	p06400		
## 87	1047	ca	simple-chemical	271	snare	
## 88	4237	p27	protein	p40305	ubiquitin	
## 89	234	akt	family	akt	insulin	
## 90	2655	fas	protein	p25445		
## 91	251	akt	family	akt	rapamycin	
## 92	255	akt	family	akt	runx2	
## 93	256	akt	family	akt	sr-bi	
## 94	5035	rb				
## 95	3029	hif-1	protein	hif1	pi3k inhibitor	
## 96	4235	p27	protein	p40305	scf	
## 97	5048	rb			tigar	
## 98	4233	p27	protein	p40305	proteasome	
## 99	5041	rb			cyclin	
## 100	4229	p27	protein	p40305	e3	
##		PosReg.Type	PosReg.ID	X.1	NegReg.Name	NegReg.Type
## 1		simple-chemical	1052	NA		
## 2		protein	p46060	NA		
## 3		family	p38	NA		
## 4				NA	grb10	protein
## 5				NA	nf1	protein
## 6				NA	pten	protein
## 7		protein	q5xxa6	NA		
## 8				NA	atrx	protein

## 9			NA	pten	protein
## 10			NA	atorvastatin	simple-chemical
## 11			NA	resveratrol	simple-chemical
## 12	simple-chemical	44795	NA		
## 13			NA	atorvastatin	simple-chemical
## 14	family	pi3k_p110	NA		
## 15			NA	selenite	simple-chemical
## 16			NA	pten	protein
## 17			NA	kdm2a	protein
## 18			NA	mdm2	protein
## 19			NA	slc12a5	protein
## 20			NA	ly294002	simple-chemical
## 21	protein	p01308	NA		
## 22	protein	p01308	NA		
## 23			NA	slc12a5	protein
## 24	protein	uaz07614	NA		
## 25			NA	pike-a	protein
## 26	protein	p99999	NA		
## 27	protein	q98146	NA		
## 28	protein	p08648	NA		
## 29			NA	akt	family
## 30	protein	p50549	NA		
## 31	protein	uaz00058	NA		
## 32			NA	pml	protein
## 33	protein	o14746	NA		
## 34	protein	q13322	NA		
## 35			NA	rb	protein
## 36	protein	q9uer7	NA		
## 37	protein	p08648	NA		
## 38			NA	arf	protein
## 39			NA	neurofibromin	protein
## 40			NA	akt	family
## 41	protein	q98146	NA		
## 42	protein	p35226	NA		
## 43	protein	q13950	NA		
## 44			NA	cas	protein
## 45	simple-chemical	40469134	NA		
## 46			NA	kp-372-2	protein
## 47			NA	p110	family
## 48			NA	nqo2	protein
## 49	simple-chemical	44795	NA		
## 50	simple-chemical	uaz00013	NA		
## 51			NA	rapamycin	simple-chemical
## 52			NA	p110	family
## 53			NA	pten	protein
## 54			NA	rapamycin	
## 55	protein	p23560	NA		
## 56	protein	p05019	NA		
## 57	protein	p08700	NA		
## 58			NA	egfr inhibitors	simple-chemical
## 59			NA	kp-372-1	protein
## 60	protein	ua-bp-ggp-00052	NA		
## 61			NA	kp372-1	protein
## 62	protein	p08648	NA		

## 63	protein	p63000	NA		
## 64	protein	q9hah7	NA		
## 65	simple-chemical	445154	NA		
## 66	protein	hdl	NA		
## 67	family	tsc	NA		
## 68			NA	abl	protein
## 69			NA	e2f3	protein
## 70			NA	hdm2	protein
## 71			NA	ly294002	simple-chemical
## 72			NA	foxo1a	protein
## 73			NA	bmi1	protein
## 74			NA	cdk4	protein
## 75			NA	arsenite	simple-chemical
## 76	simple-chemical	6857385	NA		
## 77			NA	insulin	protein
## 78			NA	pi3k	protein
## 79	protein	p01308	NA		
## 80			NA	tuberin	protein
## 81			NA	idh1	protein
## 82			NA	akt	family
## 83			NA	akt	family
## 84	simple-chemical	1090	NA		
## 85			NA	bcr	protein
## 86			NA	e2f1	protein
## 87	family	pf05739	NA		
## 88	protein	p0cg47	NA		
## 89	protein	p01308	NA		
## 90			NA	pi3k inhibitors	simple-chemical
## 91	simple-chemical	5284616	NA		
## 92	protein	q13950	NA		
## 93	protein	q8wtv0	NA		
## 94			NA	pp1c	family
## 95	simple-chemical	uaz00019	NA		
## 96	protein	p21583	NA		
## 97	protein	q9nq88	NA		
## 98	protein	proteasome	NA		
## 99	protein	p12004	NA		
## 100	simple-chemical	5756	NA		
##	NegReg.ID X.2 Connection.Type				
## 1		NA	i		
## 2		NA	i		
## 3		NA	i		
## 4	q13322	NA	i		
## 5	p21359	NA	i		
## 6	p60484	NA	i		
## 7		NA	i		
## 8	p46100	NA	i		
## 9	p60484	NA	i		
## 10	60823	NA	i		
## 11	445154	NA	i		
## 12		NA	i		
## 13	60823	NA	i		
## 14		NA	i		
## 15	1090	NA	i		

## 16	p60484	NA	i
## 17	q9y2k7	NA	i
## 18	q00987	NA	i
## 19	q9h2x9	NA	i
## 20	3973	NA	i
## 21		NA	i
## 22		NA	i
## 23	q9h2x9	NA	i
## 24		NA	i
## 25	uaz09795	NA	i
## 26		NA	i
## 27		NA	i
## 28		NA	i
## 29	akt	NA	i
## 30		NA	i
## 31		NA	i
## 32	p29590	NA	i
## 33		NA	i
## 34		NA	i
## 35	p06400	NA	i
## 36		NA	i
## 37		NA	i
## 38	q8n726	NA	i
## 39	p21359	NA	i
## 40	akt	NA	i
## 41		NA	i
## 42		NA	i
## 43		NA	i
## 44	o60716	NA	i
## 45		NA	i
## 46	uaz07773	NA	i
## 47	pi3k_p110	NA	i
## 48	p16083	NA	i
## 49		NA	i
## 50		NA	i
## 51	5284616	NA	i
## 52	pi3k_p110	NA	i
## 53	p60484	NA	i
## 54	5284616	NA	i
## 55		NA	i
## 56		NA	i
## 57		NA	i
## 58	uaz00082	NA	i
## 59	uaz00147	NA	i
## 60		NA	i
## 61	uaz00147	NA	i
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## 63		NA	i
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## 66		NA	i
## 67		NA	i
## 68	p00519	NA	i
## 69	o00716	NA	i

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##

1 ['activation', 'activation', 'activation', 'activation', 'activation', 'activation', 'activation', 'activation'
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```
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##
## 1  ['pm activation of egfr is achieved by activation of endocytosis deficient mutant egfr101011/aa :
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##	Evidence.Score	Match.Score	Kind.Score	Epistemic.Value	Total.Score
## 1	29	100	40	1	2940
## 2	15	100	40	1	1540
## 3	14	100	40	1	1440
## 4	13	100	40	1	1340
## 5	9	100	40	1	940
## 6	8	100	40	1	840
## 7	8	100	40	1	840
## 8	8	100	40	1	840
## 9	7	100	40	1	740
## 10	7	100	40	1	740
## 11	7	100	40	1	740
## 12	6	100	40	1	640
## 13	6	100	40	1	640
## 14	6	100	40	1	640
## 15	6	100	40	1	640
## 16	6	100	40	1	640
## 17	6	100	40	1	640
## 18	6	100	40	1	640
## 19	6	100	40	1	640
## 20	6	100	40	1	640
## 21	6	100	40	1	640
## 22	5	100	40	1	540
## 23	5	100	40	1	540
## 24	5	100	40	1	540
## 25	5	100	40	1	540
## 26	5	100	40	1	540
## 27	5	100	40	1	540
## 28	5	100	40	1	540
## 29	5	100	40	1	540
## 30	4	100	40	1	440
## 31	4	100	40	1	440
## 32	4	100	40	1	440
## 33	4	100	40	1	440
## 34	4	100	40	1	440
## 35	4	100	40	1	440
## 36	4	100	40	1	440
## 37	4	100	40	1	440
## 38	4	100	40	1	440
## 39	4	100	40	1	440
## 40	4	100	40	1	440
## 41	4	100	40	1	440
## 42	3	100	40	1	340
## 43	3	100	40	1	340
## 44	3	100	40	1	340
## 45	3	100	40	1	340
## 46	3	100	40	1	340
## 47	3	100	40	1	340
## 48	3	100	40	1	340
## 49	3	100	40	1	340
## 50	3	100	40	1	340

## 51	3	100	40	1	340
## 52	3	100	40	1	340
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## 57	3	100	40	1	340
## 58	3	100	40	1	340
## 59	3	100	40	1	340
## 60	3	100	40	1	340
## 61	3	100	40	1	340
## 62	3	100	40	1	340
## 63	3	100	40	1	340
## 64	3	100	40	1	340
## 65	3	100	40	1	340
## 66	3	100	40	1	340
## 67	3	100	40	1	340
## 68	3	100	40	1	340
## 69	3	100	40	1	340
## 70	3	100	40	1	340
## 71	3	100	40	1	340
## 72	3	100	40	1	340
## 73	3	100	40	1	340
## 74	3	100	40	1	340
## 75	3	100	40	1	340
## 76	3	100	40	1	340
## 77	3	100	40	1	340
## 78	3	100	40	1	340
## 79	3	100	40	1	340
## 80	3	100	40	1	340
## 81	3	100	40	1	340
## 82	3	100	40	1	340
## 83	3	100	40	1	340
## 84	3	100	40	1	340
## 85	3	100	40	1	340
## 86	3	100	40	1	340
## 87	3	100	40	1	340
## 88	3	100	40	1	340
## 89	3	100	40	1	340
## 90	3	100	40	1	340
## 91	3	100	40	1	340
## 92	3	100	40	1	340
## 93	3	100	40	1	340
## 94	3	100	40	1	340
## 95	3	100	40	1	340
## 96	3	100	40	1	340
## 97	3	100	40	1	340
## 98	3	100	40	1	340
## 99	3	100	40	1	340
## 100	3	100	40	1	340

I chose to do the top 100 interactions because the density plot doesn't give me a clear indication of the number at which it changes, so instead I chose to focus on the top 100 interactions which would be the smaller humps on the density graph. By focusing on these as my threshold, I am narrowing down the data I

am looking at and instead observing the top scored interactions.

Problem 8

This is subbing out the special characters and then unlisting the IDs so that the unique ones can be counted in the dataset. Part of this solution was sourced from the RMD created for Lab4/ Lab4 Code.

```
GBM_data$Paper.ID <-gsub("\\[", "", GBM_data$Paper.ID)
GBM_data$Paper.ID <-gsub("\\`", "", GBM_data$Paper.ID)
GBM_data$Paper.ID <-gsub("\\]", "", GBM_data$Paper.ID)
GBM_data$Paper.ID <-gsub(" ", "", GBM_data$Paper.ID)
unique(unlist(GBM_data$Paper.ID))
```

```
## [1] "'pmc3440385', 'pmc3440385', 'pmc3440385', 'pmc3440385', 'pmc3440385', 'pmc3440385', 'pmc3440385', 'p
## [2] "'pmc3440385', 'pmc3440385', 'pmc3440385', 'pmc3440385', 'pmc3440385', 'pmc3440385', 'pmc3440385', 'p
## [3] "'pmc3012146', 'pmc3012146', 'pmc3012146', 'pmc3012146', 'pmc3012146', 'pmc3012146', 'pmc3012146', 'p
## [4] "'pmc4441450', 'pmc4441450', 'pmc4441450', 'pmc4441450', 'pmc4441450', 'pmc4441450', 'pmc4441450', 'p
## [5] "'pmc4402595', 'pmc3474823', 'pmc3492274', 'pmc3492274', 'pmc3492274', 'pmc3492274', 'pmc3492274', 'p
## [6] "'pmc2777864', 'pmc2652403', 'pmc2652403', 'pmc4560817', 'pmc2652403', 'pmc2652403', 'pmc2652403', 'p
## [7] "'pmc4496210', 'pmc4496210', 'pmc4496210', 'pmc4496210', 'pmc4496210', 'pmc4496210', 'pmc4496210', 'p
## [8] "'pmc4501375', 'pmc4501375', 'pmc4501375', 'pmc4501375', 'pmc4501375', 'pmc4501375', 'pmc4501375', 'p
## [9] "'pmc2133067', 'pmc2133067', 'pmc2133067', 'pmc2133067', 'pmc2133067', 'pmc2133067', 'pmc2133067', 'p
## [10] "'pmc3789171', 'pmc3789171', 'pmc3789171', 'pmc3789171', 'pmc3789171', 'pmc3789171', 'pmc3789171', 'p
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