Lab 3 GSI

Name of studen	t *					
Todd Faulkenberry						
Readability of re	port (5 p	oints) *				
	1	2	3	4	5	
Narrative unclear and/or difficult to read and/or there is not much detail in the explanations			0			Narrative very clear and/or easy to read and lots of detail is given
Grammar of repo	ort (5 po	ints) *				
	1	2	3	4	5	
Incorrect of written grammar pervasive	0	0	\bigcirc	•	0	Excellent written grammar
Level of written or runtime *	detail on	comparis	on of R	and C++ i	mpleme	ntation and
	0	1		2	3	
Did not write about a comparison of the R and C++ implementation				0	0	Wrote a detailed comparison between the R and C++ implementations

R/C++ (3 points) *						
	0	1	2	3		
Incorrect implementations	0	•	0	\bigcirc	Seems correct to me	
Comments on in	nplementa	tion of para	allelization o	the simil	arity measure	
I am not sure exactly			-		em to be computing	
things correctly base	ed on your be	n Hur ligures,	also there is no	C++ code		
Efficiency and pr	acticality o	of R and C+	+ (3 points)	*		
a p.	0	1	2	3		
Inefficient (e.g. repeated computations, unnecessarily saved objects, etc.)					Very efficient and practical	
Suggestions for improving efficiency of R and/or C++ code *						
Looks good						
Does the author satisfy the following code readability requirements? (3 points)						
✓ Always spacing before and after variable assignment and addition symbols (" = ", " + ") and after commas						
No line of code exceeds 80 characters						
✓ Consistent variable naming for all variables (words always separated by one of "_" or "."						

Correctly coded the parallelization of k-means and pairwise similarity in

Clarity of variable	names (2 pe	oints) *		
	0	1	2	
Variable names are unclear and meaningless (e,g, 'df', 'x', 'data2)'				Variable names are helpful and unambiguous
Quality of code co	omments (2	points) *		
	0	1	2	
There are almost no comments				The comments explain clearly what is being done and why
Suggestion for im	proving read	dability of R code	*	
Overall, it looks good!				
Did the student preport (note: repo		·		their results AND
	0	1	2	
Incomplete code or no .Rnw/.Rmd file provided				Everything was provided
Clarity of folder st	tructure (2 p	oints) *		
	0	1	2	
Many excess files not relevant to report	0		0	The purpose of each file is clear and there are no excess files in the lab2 folder

Comments on folder structure

It's okay to have a data folder that has your results but remember not to upload the lingBinary.Rdata in that folder

Correctly produce	e Ben-Hur	type figures	s (3 points)	*	
	0	1	2	3	
Did not provide a figure like Ben- Hur	0	•	0	0	Figures look correct
Discuss one (or r	nore thing	ງs) that you	liked about	the author	r's figure *
I like that you kept ev	erything on t	he same scale	e, it makes sor	e easy compa	arison
Discuss one (or r	nore thing	gs) that coul	ld be impro	ved the au	thor's figure *
Again, I am not quite 3 should be the large getting larger for bigg	st (even with	_	· · · · · · · · · · · · · · · · · · ·	-	•
Also, I would choose	wider bin's f	or your histogr	am		
Justification of c	onclusion 0	s drawn fro 1	m Ben-Hur ⁻ 2	figures (3	ooints) *
Did not write about any conclusions drawn from the figures	0				Clearly outlines interpretation of the figures and drew reasonable conclusions (e.g. found k =3, or some other value, is the best

and provides reasons why)

Comments on the conclusions and interpretations of the Ben-Hur type figures *

It is unclear whether you are saying k = 3 stick out in a good or bad way here? In your plots it seems like it is bad because the similarity values are all very low and there seem to be three distinct clusters of similarity scores potential implying the points have three different ways of being clustered by k means

Things the author did well *

Your discussion of the Ben Hur algorithm was good

Things the author could improve on *

Sorry I couldn't help with the SCF thing, but if you ever can't figure something out (like the C++ code) please come ask. I'll do my best to help you figure.

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