

# Documentation\_Groupproject\_BSL

*December 8, 2017*

**Project title: Obesity Induced Changes in Gene Expression**

**Group partners: Shane Davitt, Bhavana Palakurthi, Lindsay Serene**

**Scripting Language: R**

**Teaching Assistant: Ann Raiho**

***Tasks due prior to class, 29 November 2017:***

1. Use BLAST to identify the genes encoded by the 6 differentially expressed transcripts listed in uniquetranscripts.fasta (*Lindsay*)
2. Search the NCBI protein database (Nucleotide collection (nr/nt) Database) for amino acid sequences corresponding to these 6 transcripts (*Lindsay*)
3. Translate the 4 provided files of “RNAseq data”. Write a R script to translate nucleotide sequences into amino acid sequences (*Pseudocode by Shane and Bhavana*)

***Tasks in class, 29 November 2017:***

4. Translate pseudo code into R code. (*All three*)
5. Translating using grep, substring, and while loop. (*Lindsay*)
6. Translating using regex and translate function. (*Bhavana*)

***Tasks due by the end of class, 1 December 2017:***

7. Translate pseudo code into R code. (*Lindsay and Bhavana*)
8. Build a Hidden Markov Model for each of the 6 transcript proteins and search the 4 translated “RNAseq files”. (*All the three*)

***Tasks due prior to class, 6 December 2017:***

9. Documentation of the project plan. (*Bhavana*)
10. Translate pseudo code into R code. (*Lindsay and Bhavana*)

***Tasks in class, 6 December 2017:***

11. Translate pseudo code into R code. (*Lindsay and Bhavana*)
12. Compare and discuss the code for translating RNAseq data and choose the best and efficient code. (*All the three*)
13. Peer review of the code. (*All the three*)

***Tasks due prior to class, 8 December 2017:***

14. Translate pseudo code into R code. (*Lindsay and Bhavana*)
15. Further exploration 1: Examine how discontinuous megablast and blastn qualitatively change your table of BLAST hits. (*Shane*)
16. Further exploration 2: Explore the effects of phylogenetic relatedness of amino acid sequences on the performance of the HMM mode. (*Lindsay and Bhavana*)
17. Make changes as per discussion and peer review on the final code. (*All the three*)
18. Comment collectively on the final code for all the sub-problems. (*All the three*)
19. Final check to make sure all the files are uploaded to github. (*All the three*)

***Tasks due before deadline, noon of 8, December 2017:***

19. Submission of the url for the github repo for the group project emailed to TA. (*All the three*)