BIOINF 545 / BIOSTAT 646 Group Project Written Report - 2020 Due: April 21 at 11:59pm

Remember: Everyone in the group must perform some of the analysis.

Structure of project report:

- Title. Authors
- Abstract (<=300 words)
- Introduction (1-2 pages)
- Methods (3-4 pages)
- Results (3-4 pages, cannot cite data not shown)
- Discussion (1-2 pages)
- Acknowledgements
- Project contributions from each person
- References
- Length & format (8-12 pages for Introduction, Methods, Results, and Discussion)
 - Title page, abstract, figures, tables, acknowledgments and contributions and references do not count toward the page total
 - 1½ spaced
 - Size 11 font, Arial
 - Regular 1" page margins
- Statements of existing knowledge that are not general knowledge should be cited
- If someone outside of class helped with the project please include them in the Acknowledgements
- Figures & Tables
 - May have up to 6 Tables and/or Figures
 - All Figures must have a Figure Legend; Tables can have titles and footnotes if necessary
 - Each figure or table should be placed in the report close to the place where it's described in the text.

Guidelines for writing each section

- Overall writing and presentation
 - Writing is grammatically correct and as clear as possible
 - Text is concise, not repetitive
- Title page
 - Title describes the work
 - List all authors and affiliations
- Abstract (<=300 words):
 - State question of interest/hypothesis
 - Include number of samples/ replicates and type of sample
 - Include type of data (eg, ChIP-seq, RNA-seq, ATAC-seq)
 - Describe the main method(s) used for data analysis
 - Describe the main results
 - Include a summary statement of what was learned

- Introduction:
 - Describe current state of knowledge on the question
 - Give motivation for question of interest
 - Include adequate references
 - State question(s) of interest at the end of the Introduction
- Methods:
- Give enough details that the analysis could be repeated
- Reference papers describing the methods used
- Describe the source of data and give the appropriate link or accession numbers if publicly available
- Results:
- Describe characteristics of the samples, if known (include a table if appropriate) and QC stats of the data
- Provide descriptive information about the experiment
- Describe the results of each analysis
- If sequencing, provide read depth, range of number of reads or Mb or Gb sequenced per sample
- Discussion:
 - Don't simply summarize the results
 - Put results in context of other work
 - Provide further interpretation of the results
 - Explain what you would do differently if you started over
 - Suggest future work (work that you would do if you had more time)
 - Describe any limitations of the study
- Tables
- Columns and rows are clearly labeled
- Easy to read
- Only has data that is relevant to the point being made
- Table Legend has table name
- Table legend contains all footnotes and any needed explanation of column or row headers
- Figures
- All plot axes have word labels (not the variable names used in the analysis)
- Are easy to interpret
- Symbols, plots and text are all large enough to read as printed
- Legend describes the figure content so that can understand what is on the plot without reference to the text
- Axis for bar plots should start at zero
- Be careful to not over-plot, which obscures the density of the date, if using a scatter plot. To prevent this, use an alpha value so points are semi-transparent, which allows one to see the level of overlap between points.
- Acknowledgements
 - Acknowledge students outside of your group or people outside of class who have helped with the project or who have provided data (if not publicly available)
- Project contributions

- Briefly describe each person's specific contribution to the project in terms of:
 - Analysis performed
 - Writing, report preparation
- References
 - Consistent style that includes the titles