

BI296: Linux Programming For Bioinformatics  
Course Projects: Suggestions  
2018 Spring

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In this project, you need to write a package/module to take advantage of at least one of the following packages or modules by grouping based on 3-4 persons per-group.

1. matplotlib (<https://matplotlib.org>)
2. scipy (<https://www.scipy.org>)
3. numpy (<http://www.numpy.org>)
4. pandas (<https://pandas.pydata.org>)
5. scikit-learn (<http://scikit-learn.org>)
6. statsmodels (<https://www.statsmodels.org>)
7. pysam (<https://pysam.readthedocs.io>)
8. khmer (<https://khmer.readthedocs.io>)
9. biopython (<http://biopython.org>)
10. scikit-bio (<http://scikit-bio.org>)
11. AfterQC (<https://github.com/OpenGene/AfterQC>)