

Genomics Final Project Bibliography

Project Idea: I am interested in performing some sort of statistical analysis on my gene (SCN5A) and its orthologs to see how it has genetically drifted over time. My current idea is to use the statistical programming language R to perform a multi-dimensional scaling (MDS) of SCN5A with a large number of orthologs and create a 2-dimensional scatterplot; from that data I could then see and write about what inferences come about from it in terms of evolution. My hypothesis is that since sodium voltage gated channels in heart tissue are critical in survival, that the overall genetic drift between orthologs is low. The level of drift could be quantitatively described using tdN/dS ratios amongst other statistical tools.

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