

Progress Report

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Last Week

- found some tools to do phylogenetic HGT inference, not tested yet
- Scraped all CRISPR annotaion info from CRISPRone database
- scraped NCBI data to match CRISPRone data to CRISPRdb to RefSeq files (one large dataframe now)

This Week

- Email HiDe author about error, other HGT inference tool he wrote
- Actually test the HGT programs identified
- Start testing P/A matrix scripts, tree building on genera from Athena's thesis

Next Week

- Have P/A matrix, gene,species trees for athena genomes
- Have one working inference program, at least for testing purposes