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[illegible]

Input file2
example

Input file 1
example

Output file example

Input file 2
example

Visualization example

BLAST

InterProScan

Amino acid sequences

E-value cut-off 10^{-5}

BLASTp output format 6

Amino acid sequences

Tsv format output

Pfam domains etc.

Output file example

HSDFinder

Within 10 amino acids
>90% pairwise identities
8-column spreadsheet

Visualization example

Heatmap

At least two species
KEGG pathway KO num
6-column spreadshee

