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# Laboratory Pipelines

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# QIIME

## 1 QIIME2 Microbiome

Cite vHMM [Earth Virome Project](#).

```
makeblastdb -in Combined_VF.faa -dbtype prot -out Combined_VF -title "Combined_VF"
```

```
blastp -db /Users/stronglab2/blastdb/Combined_VF/Combined_VF -out results.txt  
-outfmt 6 -query phage_proteins.faa
```

---

```
## Hello World  
print(x)
```

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# Genome

## 1 Genomic Comparisons

Use Cicos

# HMM

hello world