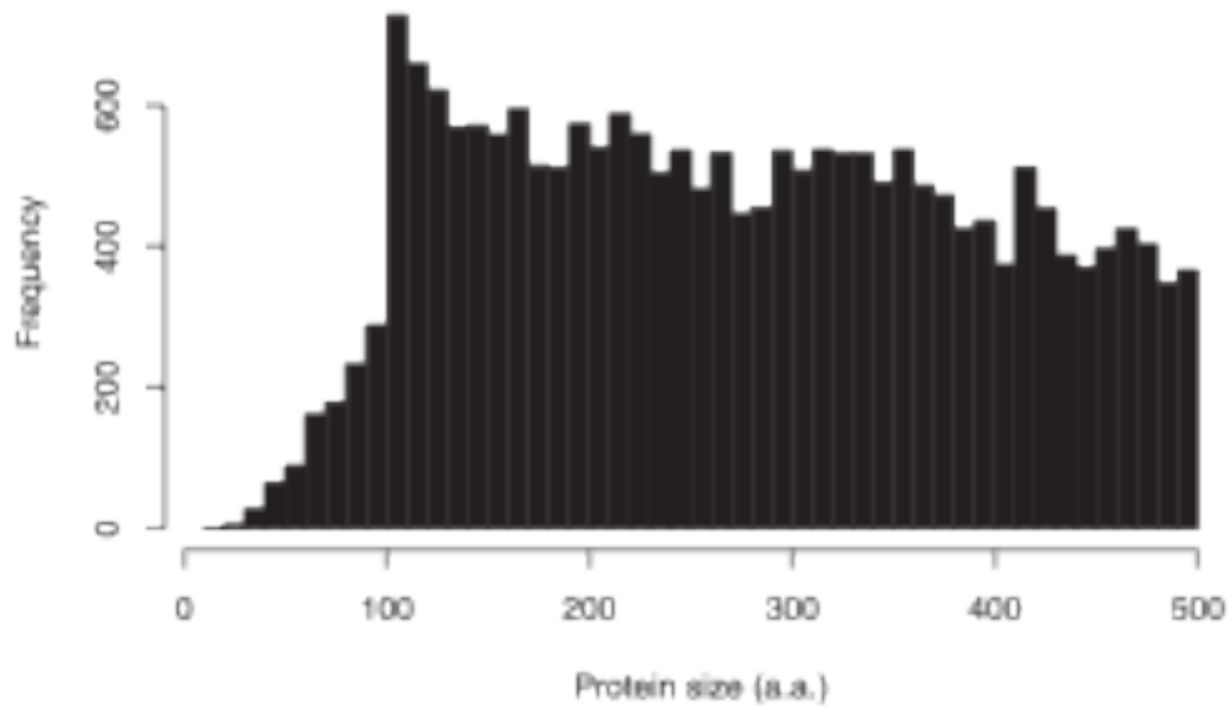


# Proteogenomics with Galaxy

Ira Cooke: LSCC / La Trobe



# Protein Sizes in Uniprot



# Lots of Different Data



Ion Trap

120 Samples



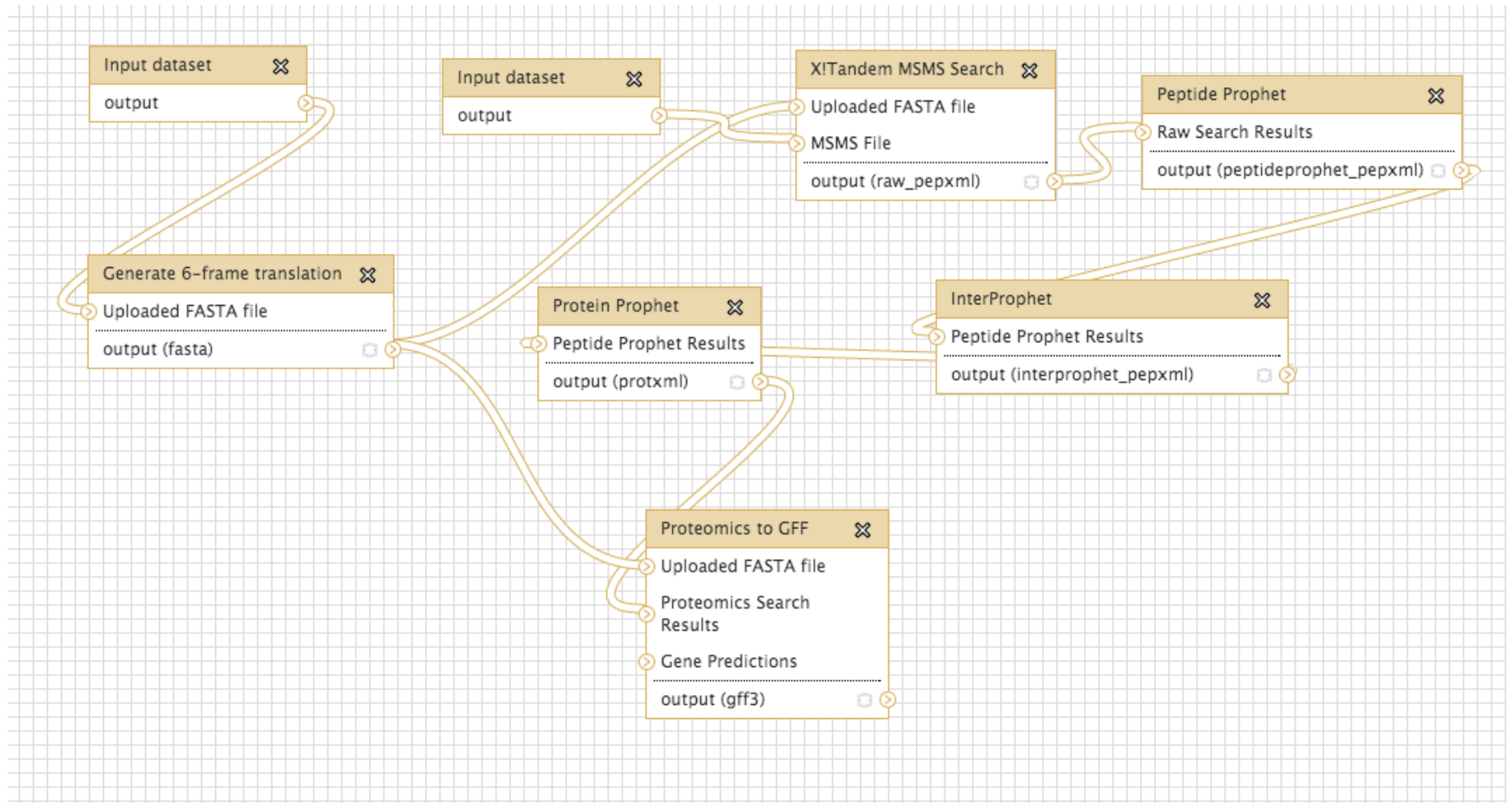
Q-TOF

4 Samples



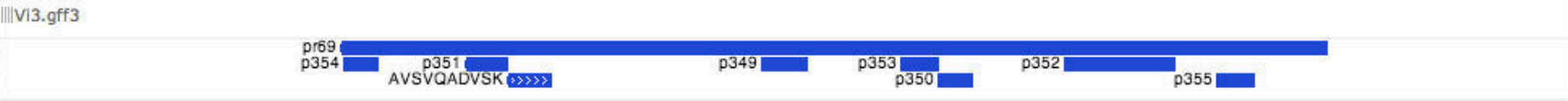
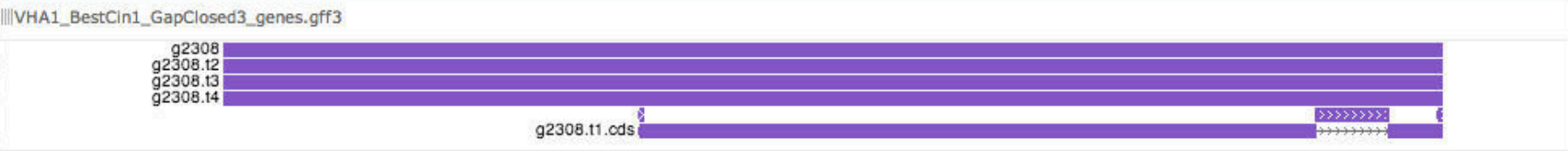
MALDI-TOF

2 Plates



Vi PG2 (i-cooke:vinaeqaulis\_v2) scaffold\_301 274,723 - 275,842

275,000



Vi PG2 (i-cooke:vinaeqaulis\_v2)

scaffold\_515

7,319 - 12,119



8,000 9,000 10,000 11,000 12,000

|||s\_7\_accepted\_hits.bam

10221 757 976 1064 4 12 49

|||VHA1\_BestCin1\_GapClosed3\_genes.gff3



|||VI3.gff3

pr143  
p716  
APAFSLSTAAAAAR  
p717

|||augustus.hints.GeneWithUTR.withNCBI\_Fungi.gff

