

Basic Analysis: We have

an assembly of an archaeal organism

gene annotation

TF binding sites

Which genes have most overlapping TFBs?

<http://cloud1.galaxyproject.org/>

<http://cloud2.galaxyproject.org/>

<http://cloud3.galaxyproject.org/>

(~ <http://usegalaxy.org/galaxy101>)

Genes & TFBs: A General Plan

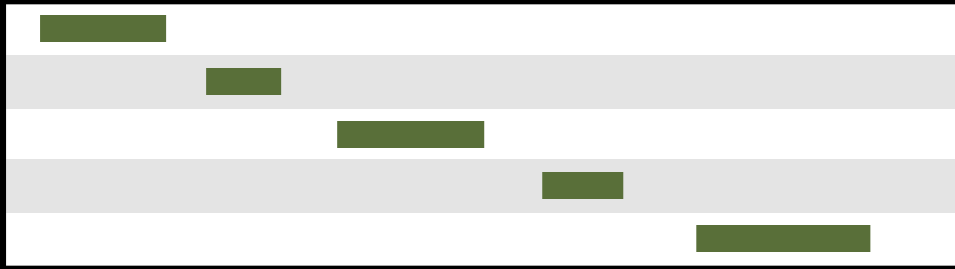
- Get some data
 - Sequence, genes/exons, TFBs
- Mess with it
 - Identify which genes/exons have TFBs
 - Count TFBs per exon
 - Visualize, save, download, ... exons with most TFBs

<http://cloud1.galaxyproject.org/>

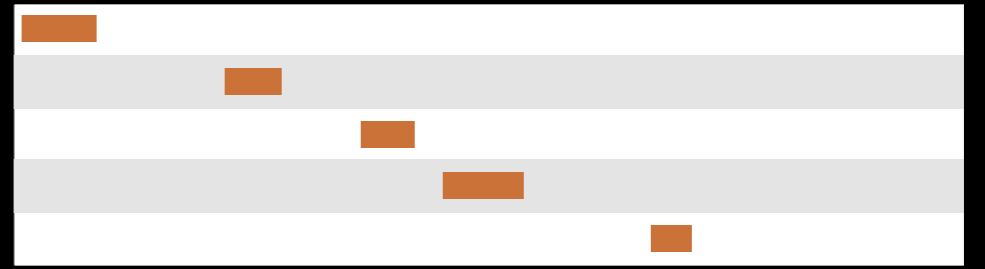
<http://cloud2.galaxyproject.org/>

<http://cloud3.galaxyproject.org/>

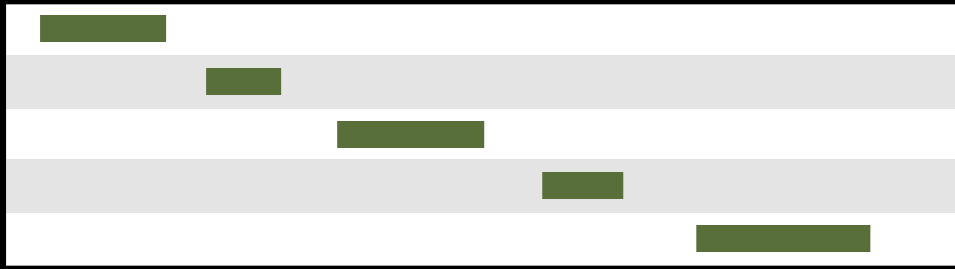
(~ <http://usegalaxy.org/galaxy101>)



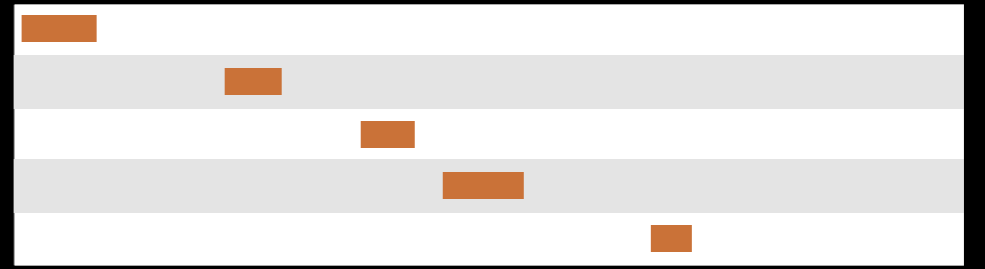
Exons



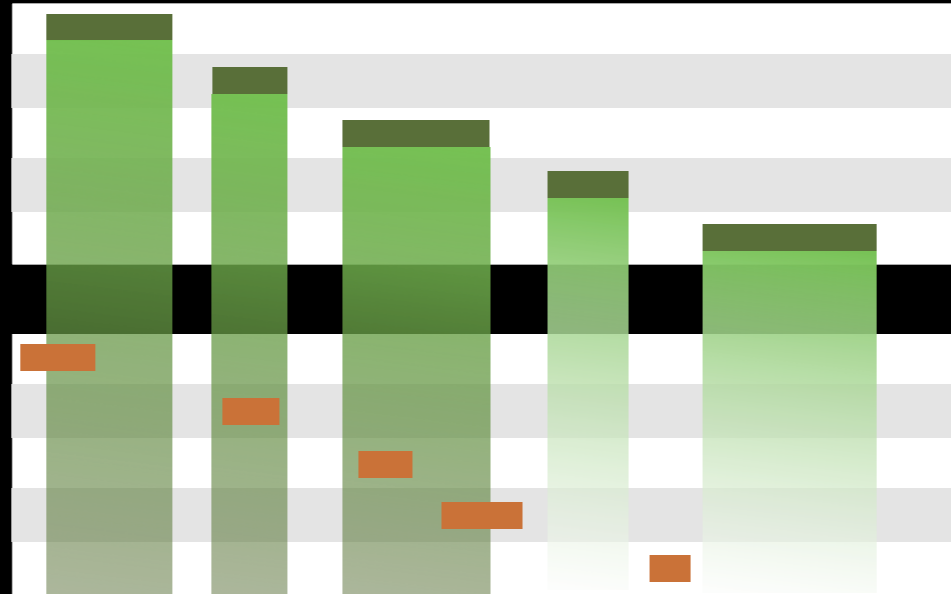
TFBs



Exons



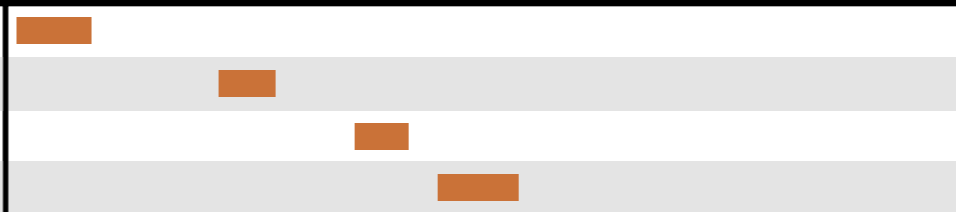
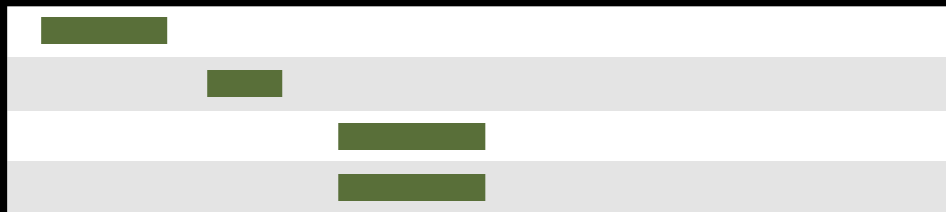
TFBs

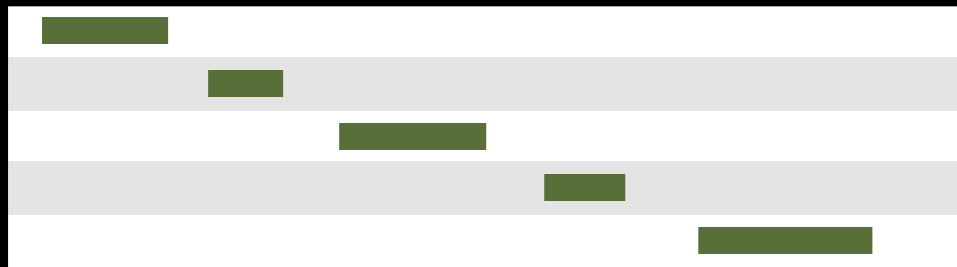


Exons

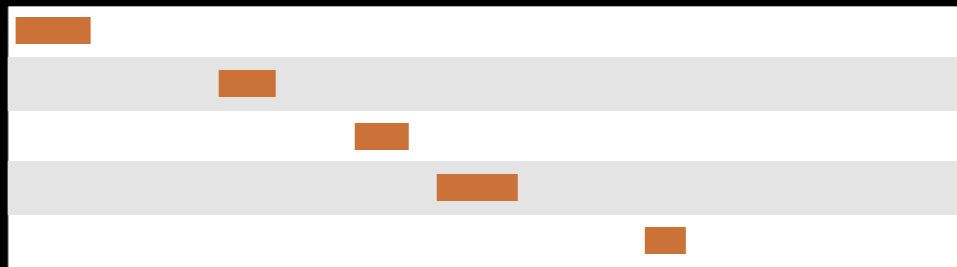
TFBs

Overlap pairings

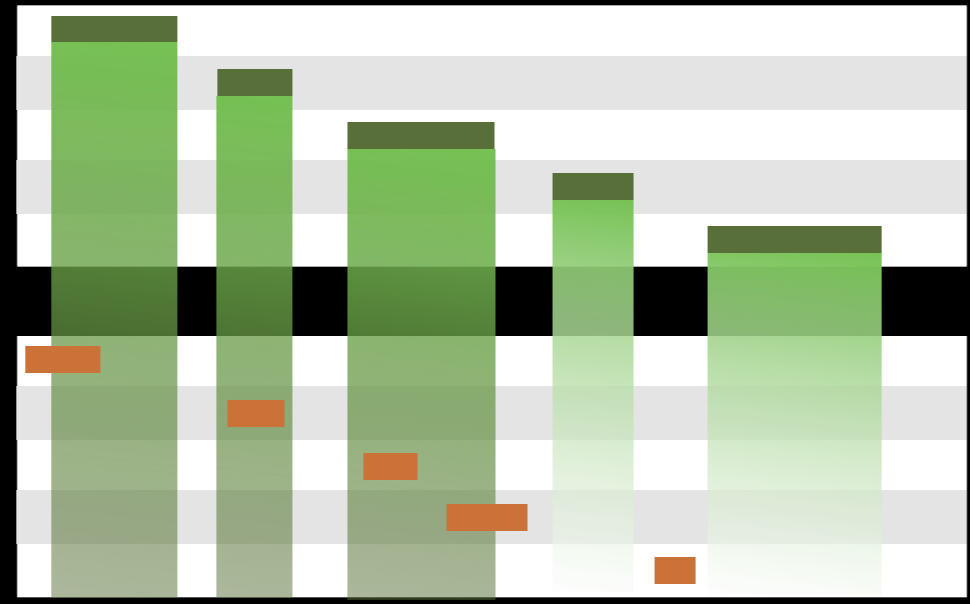




Exons



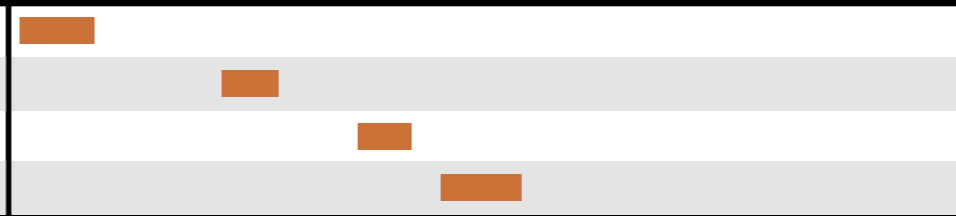
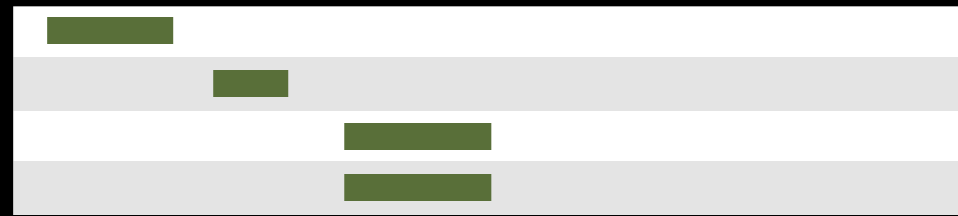
TFBS



Exons

TFBS

Overlap pairings

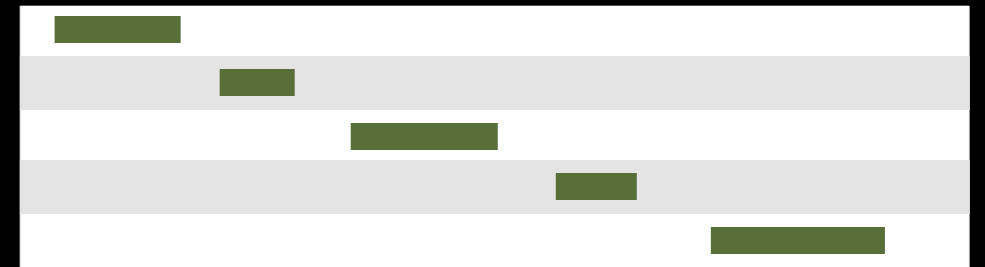


	1
	1
	2

Exon overlap counts

	1
	1
	2

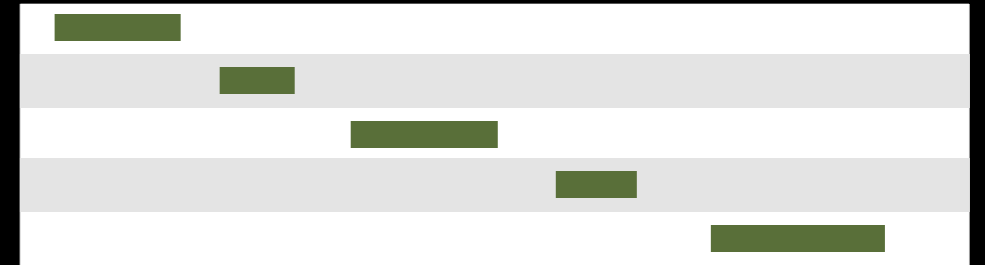
Exon overlap counts




Exons

	1
	1
	2

Exon overlap counts



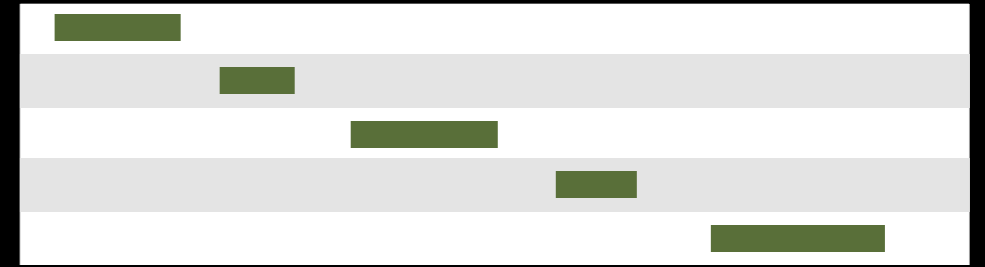
Exons

	1		0
	1		0
	2		0




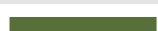
Join on exon name

	1
	1
	2

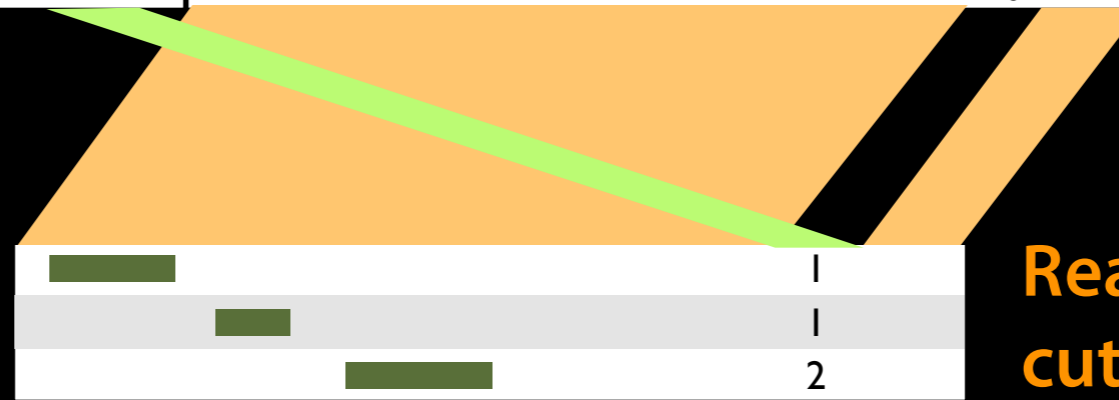
Exon overlap counts



Exons

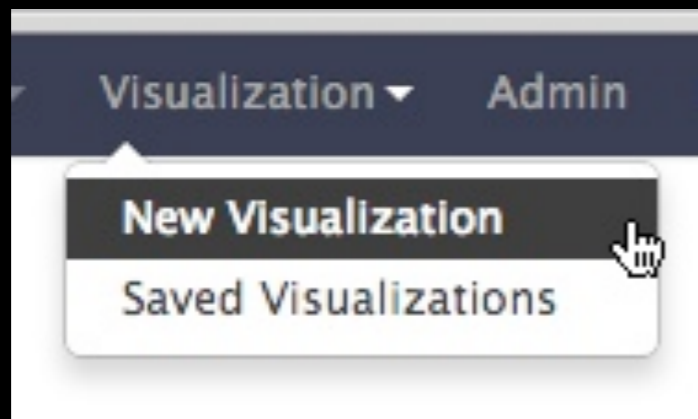
	1		0
	1		0
	2		0

Join on exon name



Rearrange columns w/
cut

Visualize results



or

