	٨	ν <u> </u>	×	
HErnants 22 4				
A HErospate 23 a 4				
A HErocotto, 23 a 4				
I HErecotte 23 + 4				
I HExpectite 22 14				
Newports 23 L4				
NETWORKS 22.44				
Li HErmanette 22 L 4				
1 Histophysia 92 + 4				
Liberanotto 23 L 4				
L HEronotto 22 L 4				
1. HEroporto, 22 1.4				
1. HErosopho 22 + 4				
1. HEropoette 22 t 4				
t HErpsoetto 22 i 4				
1. HEropoetto 22 1.4				
1. HErnnotte 22 + 1				
1. HErmontto, 22 + 4				
1. HEronoutto 22 + 1				
1. HEropoetto 22 + 1				
1. HEropoetto 22 + 1				
1. HEropoetto 22 + 1				
1. HEropoetto 22 + 1				
1. HEronootto 22 + 1				
1. HEropoetto 22 i 1				
1. HEropoetto 22 i 1				
1. HEropootto 22 i 1				
t HEronootto 22 - 1				
	1	HFrancette_23 + 1		

Note: Tracks are now grouped by subcluster and scaled. Switching in subcluster is indicated by changes in track color. Track scale is now set by default to display the region 30 bp upstream of start 1 to 30 bp downstream of the last possible start. If this default region is judged to be packed too tightly with annotated starts, the track will be further scaled to only show that region of the ORF with annotated starts. This action will be indicated by adding "Zoomed" to the title. For starts, yellow indicates the location of called starts comprised solely of Glimmer/GeneMark auto-annotations, green indicates the location of called starts with at least 1 manual gene annotation.

Pham 15955 Report

This analysis was run 04/28/24 on database version 559.

Pham number 15955 has 2 members, 0 are drafts.

Phages represented in each track:

Track 1 : HFrancette_23, Ignacio_23

Summary of Final Annotations (See graph section above for start numbers):

The start number called the most often in the published annotations is 3, it was called in 2 of the 2 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

• HFrancette_23, Ignacio_23,

Genes that have the "Most Annotated" start but do not call it:

•

Genes that do not have the "Most Annotated" start:

•

Summary by start number:

Start 3:

- Found in 2 of 2 (100.0%) of genes in pham
- Manual Annotations of this start: 2 of 2
- Called 100.0% of time when present
- Phage (with cluster) where this start called: HFrancette_23 (BP), Ignacio_23 (BP),

Summary by clusters:

There is one cluster represented in this pham: BP

Info for manual annotations of cluster BP:

•Start number 3 was manually annotated 2 times for cluster BP.

Gene Information:

Gene: HFrancette_23 Start: 18663, Stop: 18959, Start Num: 3

Candidate Starts for HFrancette_23:

(1, 18600), (2, 18603), (Start: 3 @18663 has 2 MA's), (4, 18819), (5, 18903),

Gene: Ignacio_23 Start: 18663, Stop: 18959, Start Num: 3

Candidate Starts for Ignacio_23:

(1, 18600), (2, 18603), (Start: 3 @18663 has 2 MA's), (4, 18819), (5, 18903),