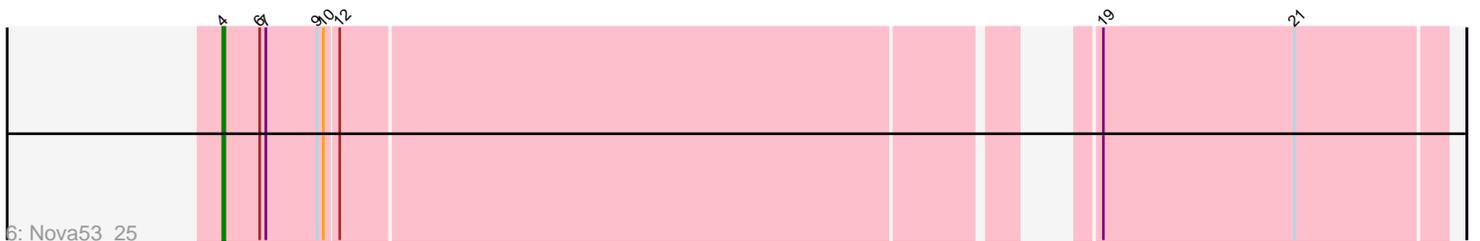
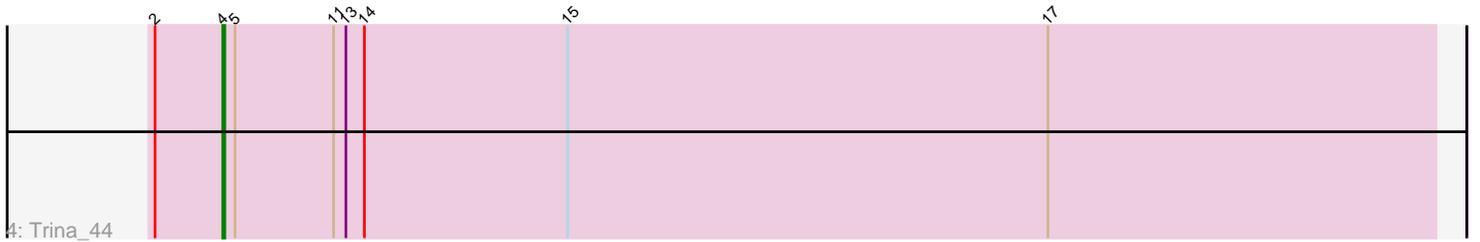


Pham 286194



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 286194 Report

This analysis was run 03/28/26 on database version 641.

Pham number 286194 has 8 members, 0 are drafts.

Phages represented in each track:

- Track 1 : Peregrin_44, Grayson_46
- Track 2 : Weasels2_46
- Track 3 : NiceHouse_39
- Track 4 : Trina_44
- Track 5 : Dorin_28, Francesca_27
- Track 6 : Nova53_25

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

The start number called the most often in the published annotations is 4, it was called in 5 of the 8 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Dorin_28, Francesca_27, NiceHouse_39, Nova53_25, Trina_44,

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

-

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

- Grayson_46, Peregrin_44, Weasels2_46,

Summary by start number:

Start 3:

- Found in 3 of 8 (37.5%) of genes in pham
- Manual Annotations of this start: 3 of 8
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Grayson_46 (CB), Peregrin_44 (CB), Weasels2_46 (CB),

Start 4:

- Found in 5 of 8 (62.5%) of genes in pham
- Manual Annotations of this start: 5 of 8
- Called 100.0% of time when present

- Phage (with cluster) where this start called: Dorin_28 (CG), Francesca_27 (CG), NiceHouse_39 (CE), Nova53_25 (CG), Trina_44 (CE),

Summary by clusters:

There are 3 clusters represented in this pham: CB, CG, CE,

Info for manual annotations of cluster CB:

- Start number 3 was manually annotated 3 times for cluster CB.

Info for manual annotations of cluster CE:

- Start number 4 was manually annotated 2 times for cluster CE.

Info for manual annotations of cluster CG:

- Start number 4 was manually annotated 3 times for cluster CG.

Gene Information:

Gene: Dorin_28 Start: 12313, Stop: 11747, Start Num: 4

Candidate Starts for Dorin_28:

(1, 12403), (Start: 4 @12313 has 5 MA's), (6, 12295), (7, 12292), (9, 12268), (10, 12265), (16, 12001),

Gene: Francesca_27 Start: 12435, Stop: 11869, Start Num: 4

Candidate Starts for Francesca_27:

(1, 12525), (Start: 4 @12435 has 5 MA's), (6, 12417), (7, 12414), (9, 12390), (10, 12387), (16, 12123),

Gene: Grayson_46 Start: 16195, Stop: 15629, Start Num: 3

Candidate Starts for Grayson_46:

(Start: 3 @16195 has 3 MA's), (20, 15757), (22, 15697), (23, 15637),

Gene: NiceHouse_39 Start: 16995, Stop: 16417, Start Num: 4

Candidate Starts for NiceHouse_39:

(Start: 4 @16995 has 5 MA's), (8, 16971), (15, 16827), (18, 16593),

Gene: Nova53_25 Start: 12099, Stop: 11551, Start Num: 4

Candidate Starts for Nova53_25:

(Start: 4 @12099 has 5 MA's), (6, 12081), (7, 12078), (9, 12054), (10, 12051), (12, 12045), (19, 11715), (21, 11622),

Gene: Peregrin_44 Start: 15553, Stop: 14987, Start Num: 3

Candidate Starts for Peregrin_44:

(Start: 3 @15553 has 3 MA's), (20, 15115), (22, 15055), (23, 14995),

Gene: Trina_44 Start: 17405, Stop: 16815, Start Num: 4

Candidate Starts for Trina_44:

(2, 17438), (Start: 4 @17405 has 5 MA's), (5, 17399), (11, 17351), (13, 17345), (14, 17336), (15, 17237), (17, 17003),

Gene: Weasels2_46 Start: 16496, Stop: 15930, Start Num: 3

Candidate Starts for Weasels2_46:

(Start: 3 @16496 has 3 MA's), (20, 16058), (23, 15938),

