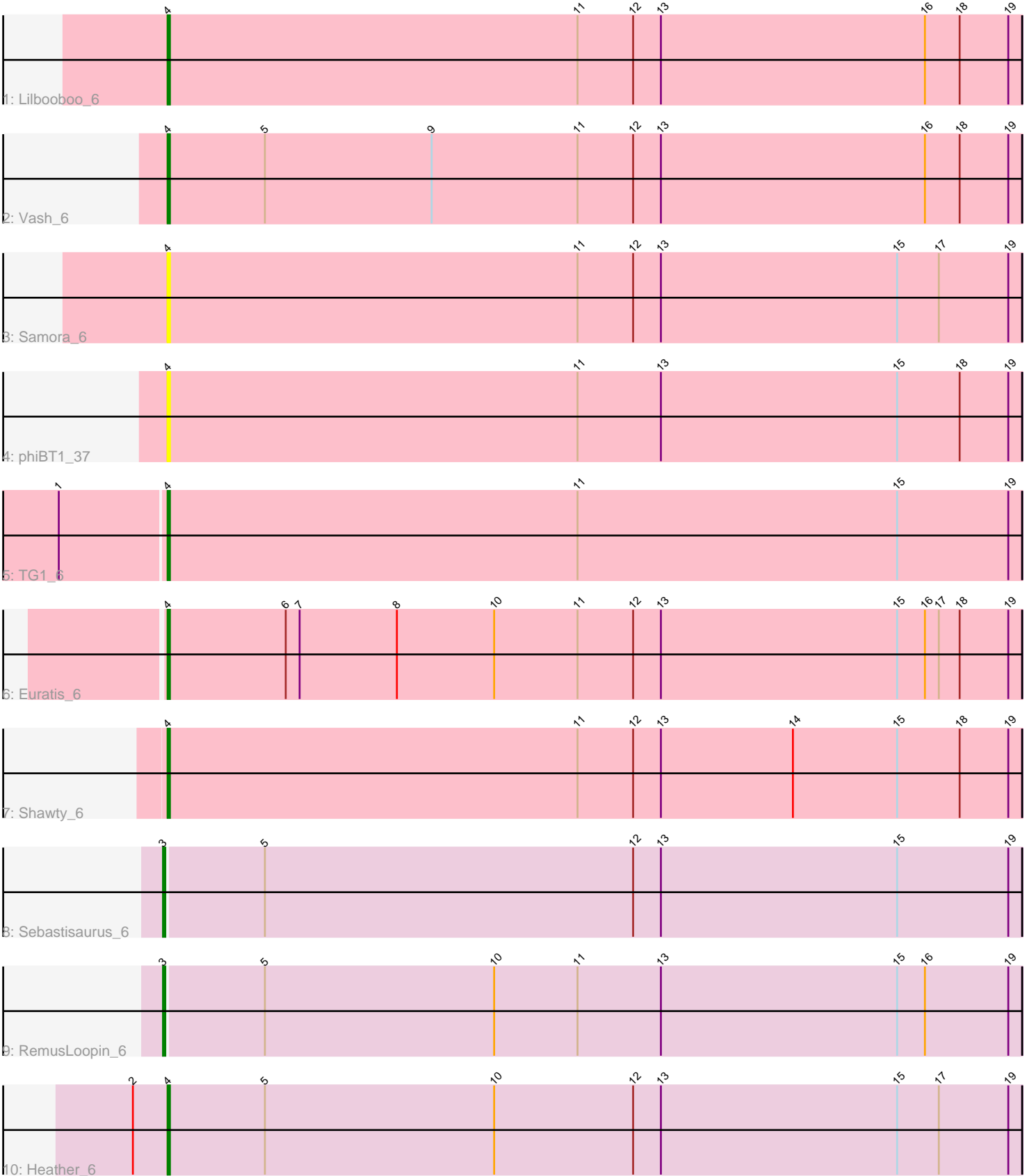


Pham 5518



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 5518 Report

This analysis was run 04/05/24 on database version 557.

WARNING: Pham size does not match number of genes in report. Either unphamerated genes have been added (by you) or starterator has removed genes due to invalid start codon.

Pham number 5518 has 10 members, 2 are drafts.

Phages represented in each track:

- Track 1 : Lilbooboo_6
- Track 2 : Vash_6
- Track 3 : Samora_6
- Track 4 : phiBT1_37
- Track 5 : TG1_6
- Track 6 : Euratis_6
- Track 7 : Shawty_6
- Track 8 : Sebastisaurus_6
- Track 9 : RemusLoopin_6
- Track 10 : Heather_6

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 6 of the 8 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Euratis_6, Heather_6, Lilbooboo_6, Samora_6, Shawty_6, TG1_6, Vash_6, phiBT1_37,

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

-

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

- RemusLoopin_6, Sebastisaurus_6,

Summary by start number:

Start 3:

- Found in 2 of 10 (20.0%) of genes in pham

- Manual Annotations of this start: 2 of 8
- Called 100.0% of time when present
- Phage (with cluster) where this start called: RemusLoopin_6 (BB2), Sebastisaurus_6 (BB2),

Start 4:

- Found in 8 of 10 (80.0%) of genes in pham
- Manual Annotations of this start: 6 of 8
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Euratis_6 (BB1), Heather_6 (BB2), Lilbooboo_6 (BB1), Samora_6 (BB1), Shawty_6 (BB1), TG1_6 (BB1), Vash_6 (BB1), phiBT1_37 (BB1),

Summary by clusters:

There are 2 clusters represented in this pham: BB2, BB1,

Info for manual annotations of cluster BB1:

- Start number 4 was manually annotated 5 times for cluster BB1.

Info for manual annotations of cluster BB2:

- Start number 3 was manually annotated 2 times for cluster BB2.
- Start number 4 was manually annotated 1 time for cluster BB2.

Gene Information:

Gene: Euratis_6 Start: 5442, Stop: 5810, Start Num: 4

Candidate Starts for Euratis_6:

(Start: 4 @5442 has 6 MA's), (6, 5493), (7, 5499), (8, 5541), (10, 5583), (11, 5619), (12, 5643), (13, 5655), (15, 5757), (16, 5769), (17, 5775), (18, 5784), (19, 5805),

Gene: Heather_6 Start: 5469, Stop: 5837, Start Num: 4

Candidate Starts for Heather_6:

(2, 5454), (Start: 4 @5469 has 6 MA's), (5, 5511), (10, 5610), (12, 5670), (13, 5682), (15, 5784), (17, 5802), (19, 5832),

Gene: Lilbooboo_6 Start: 5456, Stop: 5824, Start Num: 4

Candidate Starts for Lilbooboo_6:

(Start: 4 @5456 has 6 MA's), (11, 5633), (12, 5657), (13, 5669), (16, 5783), (18, 5798), (19, 5819),

Gene: RemusLoopin_6 Start: 5452, Stop: 5820, Start Num: 3

Candidate Starts for RemusLoopin_6:

(Start: 3 @5452 has 2 MA's), (5, 5494), (10, 5593), (11, 5629), (13, 5665), (15, 5767), (16, 5779), (19, 5815),

Gene: Samora_6 Start: 5486, Stop: 5854, Start Num: 4

Candidate Starts for Samora_6:

(Start: 4 @5486 has 6 MA's), (11, 5663), (12, 5687), (13, 5699), (15, 5801), (17, 5819), (19, 5849),

Gene: Sebastisaurus_6 Start: 5474, Stop: 5842, Start Num: 3

Candidate Starts for Sebastisaurus_6:

(Start: 3 @5474 has 2 MA's), (5, 5516), (12, 5675), (13, 5687), (15, 5789), (19, 5837),

Gene: Shawty_6 Start: 5474, Stop: 5842, Start Num: 4

Candidate Starts for Shawty_6:

(Start: 4 @5474 has 6 MA's), (11, 5651), (12, 5675), (13, 5687), (14, 5744), (15, 5789), (18, 5816), (19, 5837),

Gene: TG1_6 Start: 5464, Stop: 5832, Start Num: 4

Candidate Starts for TG1_6:

(1, 5419), (Start: 4 @5464 has 6 MA's), (11, 5641), (15, 5779), (19, 5827),

Gene: Vash_6 Start: 5464, Stop: 5832, Start Num: 4

Candidate Starts for Vash_6:

(Start: 4 @5464 has 6 MA's), (5, 5506), (9, 5578), (11, 5641), (12, 5665), (13, 5677), (16, 5791), (18, 5806), (19, 5827),

Gene: phiBT1_37 Start: 5485, Stop: 5853, Start Num: 4

Candidate Starts for phiBT1_37:

(Start: 4 @5485 has 6 MA's), (11, 5662), (13, 5698), (15, 5800), (18, 5827), (19, 5848),