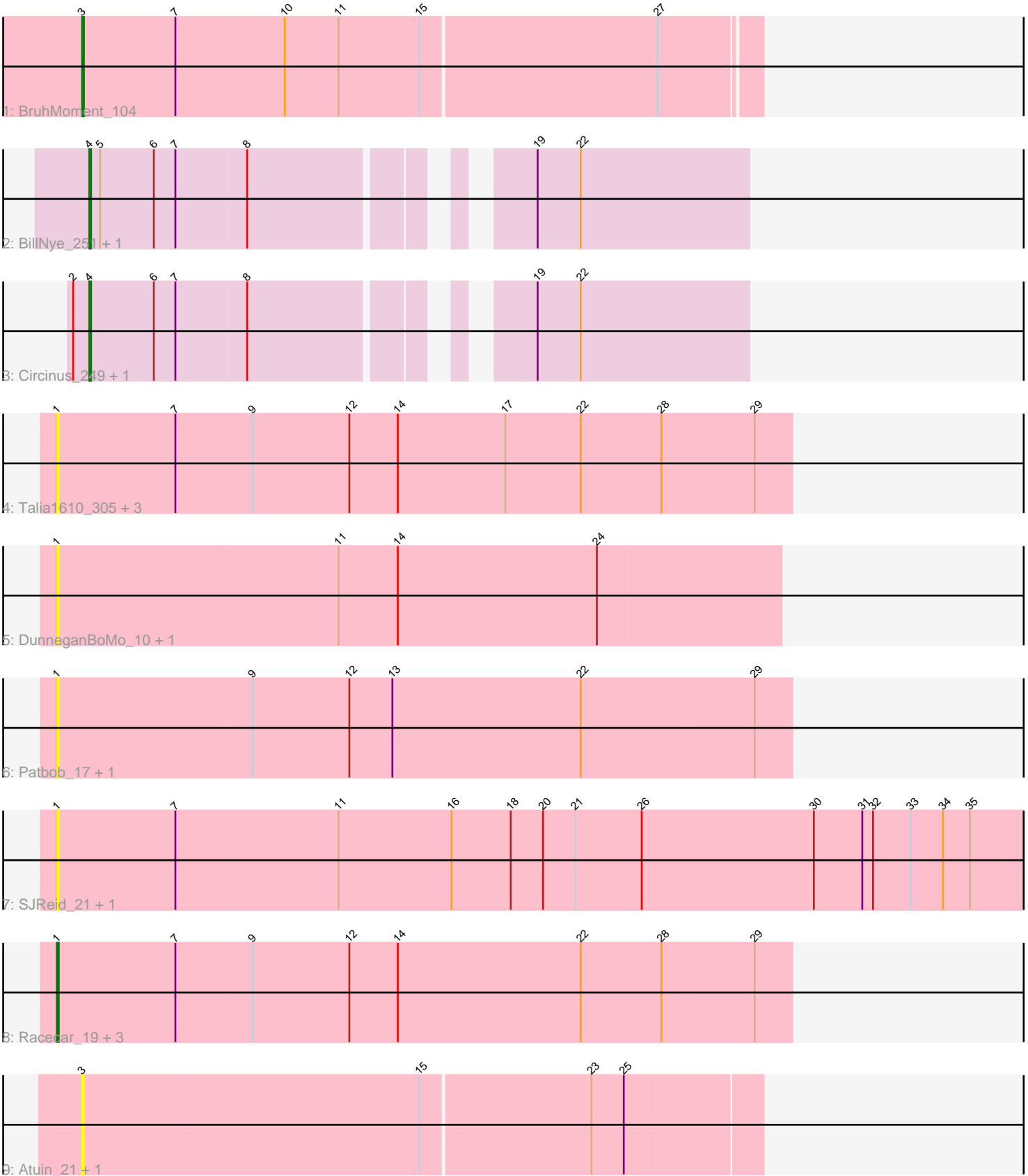


Pham 153508



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 153508 Report

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

Pham number 153508 has 21 members, 14 are drafts.

Phages represented in each track:

- Track 1 : BruhMoment\_104
- Track 2 : BillNye\_251, BillNye\_5
- Track 3 : Circinus\_249, Circinus\_4
- Track 4 : Talia1610\_305, Talia1610\_18, Mimi\_20, Mimi\_310
- Track 5 : DunneganBoMo\_10, DunneganBoMo\_313
- Track 6 : Patbob\_17, Patbob\_307
- Track 7 : SJReid\_21, SJReid\_332
- Track 8 : Racecar\_19, Bloom\_307, Racecar\_308, Bloom\_20
- Track 9 : Atuin\_21, Atuin\_328

### ***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 4 of the 7 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- BillNye\_251, BillNye\_5, Circinus\_249, Circinus\_4,

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

- 

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

- Atuin\_21, Atuin\_328, Bloom\_20, Bloom\_307, BruhMoment\_104, DunneganBoMo\_10, DunneganBoMo\_313, Mimi\_20, Mimi\_310, Patbob\_17, Patbob\_307, Racecar\_19, Racecar\_308, SJReid\_21, SJReid\_332, Talia1610\_18, Talia1610\_305,

### **Summary by start number:**

Start 1:

- Found in 14 of 21 ( 66.7% ) of genes in pham
- Manual Annotations of this start: 2 of 7
- Called 100.0% of time when present

- Phage (with cluster) where this start called: Bloom\_20 (FC), Bloom\_307 (FC), DunneganBoMo\_10 (FC), DunneganBoMo\_313 (FC), Mimi\_20 (FC), Mimi\_310 (FC), Patbob\_17 (FC), Patbob\_307 (FC), Racecar\_19 (FC), Racecar\_308 (FC), SJReid\_21 (FC), SJReid\_332 (FC), Talia1610\_18 (FC), Talia1610\_305 (FC),

Start 3:

- Found in 3 of 21 ( 14.3% ) of genes in pham
- Manual Annotations of this start: 1 of 7
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Atuin\_21 (FC), Atuin\_328 (FC), BruhMoment\_104 (AP3),

Start 4:

- Found in 4 of 21 ( 19.0% ) of genes in pham
- Manual Annotations of this start: 4 of 7
- Called 100.0% of time when present
- Phage (with cluster) where this start called: BillNye\_251 (BK2), BillNye\_5 (BK2), Circinus\_249 (BK2), Circinus\_4 (BK2),

### **Summary by clusters:**

There are 3 clusters represented in this pham: AP3, FC, BK2,

Info for manual annotations of cluster AP3:

- Start number 3 was manually annotated 1 time for cluster AP3.

Info for manual annotations of cluster BK2:

- Start number 4 was manually annotated 4 times for cluster BK2.

Info for manual annotations of cluster FC:

- Start number 1 was manually annotated 2 times for cluster FC.

### **Gene Information:**

Gene: Atuin\_21 Start: 9989, Stop: 10360, Start Num: 3

Candidate Starts for Atuin\_21:

(Start: 3 @9989 has 1 MA's), (15, 10175), (23, 10268), (25, 10286),

Gene: Atuin\_328 Start: 186877, Stop: 187248, Start Num: 3

Candidate Starts for Atuin\_328:

(Start: 3 @186877 has 1 MA's), (15, 187063), (23, 187156), (25, 187174),

Gene: BillNye\_251 Start: 126516, Stop: 126845, Start Num: 4

Candidate Starts for BillNye\_251:

(Start: 4 @126516 has 4 MA's), (5, 126522), (6, 126552), (7, 126564), (8, 126603), (19, 126729), (22, 126753),

Gene: BillNye\_5 Start: 1694, Stop: 2023, Start Num: 4

Candidate Starts for BillNye\_5:

(Start: 4 @1694 has 4 MA's), (5, 1700), (6, 1730), (7, 1742), (8, 1781), (19, 1907), (22, 1931),

Gene: Bloom\_307 Start: 182892, Stop: 183299, Start Num: 1

Candidate Starts for Bloom\_307:

(Start: 1 @182892 has 2 MA's), (7, 182958), (9, 183000), (12, 183054), (14, 183081), (22, 183183), (28, 183228), (29, 183279),

Gene: Bloom\_20 Start: 9417, Stop: 9824, Start Num: 1

Candidate Starts for Bloom\_20:

(Start: 1 @9417 has 2 MA's), (7, 9483), (9, 9525), (12, 9579), (14, 9606), (22, 9708), (28, 9753), (29, 9804),

Gene: BruhMoment\_104 Start: 63229, Stop: 62861, Start Num: 3

Candidate Starts for BruhMoment\_104:

(Start: 3 @63229 has 1 MA's), (7, 63178), (10, 63118), (11, 63088), (15, 63043), (27, 62914),

Gene: Circinus\_249 Start: 125815, Stop: 126144, Start Num: 4

Candidate Starts for Circinus\_249:

(2, 125806), (Start: 4 @125815 has 4 MA's), (6, 125851), (7, 125863), (8, 125902), (19, 126028), (22, 126052),

Gene: Circinus\_4 Start: 1488, Stop: 1817, Start Num: 4

Candidate Starts for Circinus\_4:

(2, 1479), (Start: 4 @1488 has 4 MA's), (6, 1524), (7, 1536), (8, 1575), (19, 1701), (22, 1725),

Gene: DunneganBoMo\_10 Start: 4570, Stop: 4971, Start Num: 1

Candidate Starts for DunneganBoMo\_10:

(Start: 1 @4570 has 2 MA's), (11, 4726), (14, 4759), (24, 4870),

Gene: DunneganBoMo\_313 Start: 183982, Stop: 184383, Start Num: 1

Candidate Starts for DunneganBoMo\_313:

(Start: 1 @183982 has 2 MA's), (11, 184138), (14, 184171), (24, 184282),

Gene: Mimi\_20 Start: 8854, Stop: 9261, Start Num: 1

Candidate Starts for Mimi\_20:

(Start: 1 @8854 has 2 MA's), (7, 8920), (9, 8962), (12, 9016), (14, 9043), (17, 9103), (22, 9145), (28, 9190), (29, 9241),

Gene: Mimi\_310 Start: 181514, Stop: 181921, Start Num: 1

Candidate Starts for Mimi\_310:

(Start: 1 @181514 has 2 MA's), (7, 181580), (9, 181622), (12, 181676), (14, 181703), (17, 181763), (22, 181805), (28, 181850), (29, 181901),

Gene: Patbob\_17 Start: 8860, Stop: 9267, Start Num: 1

Candidate Starts for Patbob\_17:

(Start: 1 @8860 has 2 MA's), (9, 8968), (12, 9022), (13, 9046), (22, 9151), (29, 9247),

Gene: Patbob\_307 Start: 184319, Stop: 184726, Start Num: 1

Candidate Starts for Patbob\_307:

(Start: 1 @184319 has 2 MA's), (9, 184427), (12, 184481), (13, 184505), (22, 184610), (29, 184706),

Gene: Racecar\_19 Start: 9417, Stop: 9824, Start Num: 1

Candidate Starts for Racecar\_19:

(Start: 1 @9417 has 2 MA's), (7, 9483), (9, 9525), (12, 9579), (14, 9606), (22, 9708), (28, 9753), (29, 9804),

Gene: Racecar\_308 Start: 183126, Stop: 183533, Start Num: 1

Candidate Starts for Racecar\_308:

(Start: 1 @183126 has 2 MA's), (7, 183192), (9, 183234), (12, 183288), (14, 183315), (22, 183417), (28, 183462), (29, 183513),

Gene: SJReid\_21 Start: 9248, Stop: 9805, Start Num: 1

Candidate Starts for SJReid\_21:

(Start: 1 @9248 has 2 MA's), (7, 9314), (11, 9404), (16, 9467), (18, 9500), (20, 9518), (21, 9536), (26, 9572), (30, 9668), (31, 9695), (32, 9701), (33, 9722), (34, 9740), (35, 9755),

Gene: SJReid\_332 Start: 182087, Stop: 182644, Start Num: 1

Candidate Starts for SJReid\_332:

(Start: 1 @182087 has 2 MA's), (7, 182153), (11, 182243), (16, 182306), (18, 182339), (20, 182357), (21, 182375), (26, 182411), (30, 182507), (31, 182534), (32, 182540), (33, 182561), (34, 182579), (35, 182594),

Gene: Talia1610\_305 Start: 183328, Stop: 183735, Start Num: 1

Candidate Starts for Talia1610\_305:

(Start: 1 @183328 has 2 MA's), (7, 183394), (9, 183436), (12, 183490), (14, 183517), (17, 183577), (22, 183619), (28, 183664), (29, 183715),

Gene: Talia1610\_18 Start: 8856, Stop: 9263, Start Num: 1

Candidate Starts for Talia1610\_18:

(Start: 1 @8856 has 2 MA's), (7, 8922), (9, 8964), (12, 9018), (14, 9045), (17, 9105), (22, 9147), (28, 9192), (29, 9243),