

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 200787 Report

This analysis was run 01/18/25 on database version 583.

Pham number 200787 has 13 members, 5 are drafts.

Phages represented in each track:

- Track 1 : BirdsNest 29
- Track 2 : Zenteno07\_30
- Track 3 : Cooper\_24
- Track 4 : KayaCho\_22
- Track 5 : Hosp 020
- Track 6 : Jolie1\_022
- Track 7: 40BC 022
- Track 8 : 39HC 022
- Track 9 : ForDig 23
- Track 10 : Knocker 28
- Track 11: CRB2 29
- Track 12 : PenguinLover67\_28
- Track 13 : Evaa\_24

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

The start number called the most often in the published annotations is 8, it was called in 1 of the 8 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

• 40BC\_022, CRB2\_29, ForDig\_23, Hosp\_020,

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

• 39HC\_022, Jolie1\_022, KayaCho\_22,

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

• BirdsNest\_29, Cooper\_24, Evaa\_24, Knocker\_28, PenguinLover67\_28, Zenteno07\_30,

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

#### Start 3:

- Found in 1 of 13 (7.7%) of genes in pham
- Manual Annotations of this start: 1 of 8

- Called 100.0% of time when present
- Phage (with cluster) where this start called: KayaCho\_22 (B6),

### Start 5:

- Found in 4 of 13 (30.8%) of genes in pham
- No Manual Annotations of this start.
- Called 25.0% of time when present
- Phage (with cluster) where this start called: 39HC\_022 (B6),

### Start 6:

- Found in 1 of 13 (7.7%) of genes in pham
- No Manual Annotations of this start.
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Jolie1\_022 (B6),

#### Start 8:

- Found in 7 of 13 (53.8%) of genes in pham
- Manual Annotations of this start: 1 of 8
- Called 57.1% of time when present
- Phage (with cluster) where this start called: 40BC\_022 (B6), CRB2\_29 (B9), ForDig\_23 (B6), Hosp\_020 (B6),

### Start 10:

- Found in 1 of 13 (7.7%) of genes in pham
- Manual Annotations of this start: 1 of 8
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Knocker\_28 (B9),

#### Start 13:

- Found in 2 of 13 (15.4%) of genes in pham
- Manual Annotations of this start: 1 of 8
- Called 50.0% of time when present
- Phage (with cluster) where this start called: Zenteno07 30 (B13),

### Start 17:

- Found in 2 of 13 (15.4%) of genes in pham
- Manual Annotations of this start: 1 of 8
- Called 50.0% of time when present
- Phage (with cluster) where this start called: BirdsNest\_29 (B13),

#### Start 18:

- Found in 1 of 13 (7.7%) of genes in pham
- Manual Annotations of this start: 1 of 8
- Called 100.0% of time when present
- Phage (with cluster) where this start called: PenguinLover67\_28 (B9),

### Start 21:

- Found in 1 of 13 (7.7%) of genes in pham
- Manual Annotations of this start: 1 of 8
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Cooper\_24 (B4),

#### Start 37:

- Found in 1 of 13 (7.7%) of genes in pham
- Manual Annotations of this start: 1 of 8
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Evaa\_24 (DR),

## **Summary by clusters:**

There are 5 clusters represented in this pham: B4, B6, DR, B13, B9,

Info for manual annotations of cluster B13:

- •Start number 13 was manually annotated 1 time for cluster B13.
- Start number 17 was manually annotated 1 time for cluster B13.

Info for manual annotations of cluster B4:

•Start number 21 was manually annotated 1 time for cluster B4.

Info for manual annotations of cluster B6:

•Start number 3 was manually annotated 1 time for cluster B6.

Info for manual annotations of cluster B9:

- •Start number 8 was manually annotated 1 time for cluster B9.
- •Start number 10 was manually annotated 1 time for cluster B9.
- •Start number 18 was manually annotated 1 time for cluster B9.

Info for manual annotations of cluster DR:

•Start number 37 was manually annotated 1 time for cluster DR.

### Gene Information:

Gene: 39HC 022 Start: 18307, Stop: 19038, Start Num: 5

Candidate Starts for 39HC 022:

(5, 18307), (7, 18340), (Start: 8 @ 18370 has 1 MA's), (9, 18397), (24, 18490), (26, 18496), (29, 18514), (36, 18589), (50, 18709), (58, 18865), (60, 18928), (62, 18943), (64, 18979), (67, 19003),

Gene: 40BC\_022 Start: 18370, Stop: 19038, Start Num: 8

Candidate Starts for 40BC\_022:

(5, 18307), (7, 18340), (Start: 8 @18370 has 1 MA's), (9, 18397), (24, 18490), (26, 18496), (29, 18514), (36, 18589), (50, 18709), (58, 18865), (60, 18928), (62, 18943), (64, 18979), (67, 19003),

Gene: BirdsNest\_29 Start: 19215, Stop: 19856, Start Num: 17

Candidate Starts for BirdsNest 29:

(Start: 13 @19173 has 1 MA's), (Start: 17 @19215 has 1 MA's), (22, 19251), (48, 19473), (54, 19563), (58, 19662), (60, 19725), (62, 19740),

Gene: CRB2\_29 Start: 21997, Stop: 22677, Start Num: 8

Candidate Starts for CRB2 29:

(Start: 8 @21997 has 1 MA's), (19, 22096), (28, 22132), (47, 22300), (60, 22549), (62, 22564), (63, 22591), (64, 22600), (67, 22624), (68, 22627),

Gene: Cooper 24 Start: 18981, Stop: 19541, Start Num: 21

Candidate Starts for Cooper\_24:

(Start: 21 @18981 has 1 MA's), (31, 19050), (40, 19116), (41, 19125), (50, 19212), (51, 19215), (53, 19236), (56, 19356), (57, 19362), (58, 19368), (60, 19431), (62, 19446), (63, 19473), (64, 19482),

Gene: Evaa\_24 Start: 18931, Stop: 19455, Start Num: 37

Candidate Starts for Evaa\_24:

(14, 18775), (20, 18820), (32, 18892), (Start: 37 @18931 has 1 MA's), (39, 18940), (44, 18994), (55, 19171), (56, 19222), (57, 19228), (58, 19234), (59, 19261), (60, 19297), (61, 19300), (62, 19312), (66, 19360), (67, 19372),

Gene: ForDig\_23 Start: 18356, Stop: 19030, Start Num: 8

Candidate Starts for ForDig 23:

(5, 18293), (7, 18326), (Start: 8 @ 18356 has 1 MA's), (9, 18383), (24, 18476), (27, 18488), (29, 18500), (36, 18578), (50, 18698), (60, 18920), (62, 18935), (64, 18971), (67, 18995),

Gene: Hosp\_020 Start: 16572, Stop: 17234, Start Num: 8

Candidate Starts for Hosp\_020:

(5, 16509), (7, 16542), (Start: 8 @ 16572 has 1 MA's), (9, 16599), (24, 16692), (26, 16698), (27, 16704), (29, 16716), (36, 16785), (50, 16905), (58, 17061), (60, 17124), (62, 17139), (64, 17175), (67, 17199),

Gene: Jolie1\_022 Start: 18308, Stop: 19030, Start Num: 6

Candidate Starts for Jolie 1 022:

(6, 18308), (7, 18323), (Start: 8 @18353 has 1 MA's), (9, 18380), (24, 18473), (27, 18485), (29, 18497), (36, 18578), (38, 18587), (50, 18698), (60, 18920), (62, 18935), (64, 18971), (67, 18995),

Gene: KayaCho\_22 Start: 18287, Stop: 19042, Start Num: 3

Candidate Starts for KayaCho\_22:

(Start: 3 @18287 has 1 MA's), (Start: 8 @18365 has 1 MA's), (12, 18419), (23, 18491), (25, 18500), (29, 18521), (36, 18593), (50, 18710), (60, 18932), (62, 18947), (64, 18983), (67, 19007),

Gene: Knocker\_28 Start: 22251, Stop: 22850, Start Num: 10

Candidate Starts for Knocker\_28:

(Start: 10 @22251 has 1 MA's), (11, 22266), (15, 22290), (30, 22371), (33, 22383), (34, 22398), (42, 22443), (47, 22479), (49, 22497), (53, 22539), (56, 22656), (57, 22662), (60, 22731), (62, 22746), (63, 22773), (65, 22791), (67, 22806), (68, 22809), (69, 22818).

Gene: PenguinLover67\_28 Start: 21391, Stop: 22002, Start Num: 18

Candidate Starts for PenguinLover67\_28:

(2, 21145), (4, 21208), (14, 21358), (16, 21370), (Start: 18 @21391 has 1 MA's), (46, 21610), (47, 21625), (52, 21682), (53, 21688), (60, 21883), (62, 21898), (63, 21925), (64, 21934), (68, 21961),

Gene: Zenteno07\_30 Start: 19548, Stop: 20225, Start Num: 13

Candidate Starts for Zenteno07\_30:

(1, 19302), (Start: 13 @19548 has 1 MA's), (Start: 17 @19590 has 1 MA's), (35, 19728), (43, 19800), (45, 19815), (48, 19842), (54, 19932), (58, 20031), (60, 20094), (62, 20109), (67, 20169),