

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 194391 Report

This analysis was run 11/02/24 on database version 579.

Pham number 194391 has 17 members, 3 are drafts.

Phages represented in each track:

- Track 1 : Katyusha_35, Benczkowski14_35, Teatealatte_35, Teech_35, Niagara_35, Tredge_35
- Track 2 : Kvothe_35
- Track 3: Hollow_36
- Track 4: Demosthenes 35
- Track 5 : ASerpRocky_35
- Track 6: Philon9 37
- Track 7 : Wocket 24
- Track 8 : LonelyBoi 24
- Track 9 : Sidious 23
- Track 10 : Camerico 40
- Track 11 : GMA3 28
- Track 12 : Jumbo 37

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

The start number called the most often in the published annotations is 10, it was called in 9 of the 14 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

• ASerpRocky_35, Benczkowski14_35, Demosthenes_35, Hollow_36, Katyusha_35, Kvothe_35, Niagara_35, Philon9_37, Teatealatte_35, Teech_35, Tredge_35,

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

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

Camerico_40, GMA3_28, Jumbo_37, LonelyBoi_24, Sidious_23, Wocket_24,

Summary by start number:

Start 2:

- Found in 1 of 17 (5.9%) of genes in pham
- Manual Annotations of this start: 1 of 14

- Called 100.0% of time when present
- Phage (with cluster) where this start called: Camerico_40 (DF),

Start 4:

- Found in 1 of 17 (5.9%) of genes in pham
- No Manual Annotations of this start.
- Called 100.0% of time when present
- Phage (with cluster) where this start called: GMA3_28 (DF2),

Start 5:

- Found in 1 of 17 (5.9%) of genes in pham
- Manual Annotations of this start: 1 of 14
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Jumbo_37 (DF3),

Start 7:

- Found in 1 of 17 (5.9%) of genes in pham
- Manual Annotations of this start: 1 of 14
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Wocket_24 (CV),

Start 9:

- Found in 2 of 17 (11.8%) of genes in pham
- Manual Annotations of this start: 2 of 14
- Called 100.0% of time when present
- Phage (with cluster) where this start called: LonelyBoi_24 (CY), Sidious_23 (CZ7),

Start 10:

- Found in 11 of 17 (64.7%) of genes in pham
- Manual Annotations of this start: 9 of 14
- Called 100.0% of time when present
- Phage (with cluster) where this start called: ASerpRocky_35 (CS4),
 Benczkowski14_35 (CS4), Demosthenes_35 (CS4), Hollow_36 (CS4), Katyusha_35 (CS4), Kvothe_35 (CS4), Niagara_35 (CS4), Philon9_37 (CS4), Teatealatte_35 (CS4), Teech_35 (CS4), Tredge_35 (CS4),

Summary by clusters:

There are 7 clusters represented in this pham: DF, CZ7, DF3, DF2, CY, CS4, CV,

Info for manual annotations of cluster CS4:

•Start number 10 was manually annotated 9 times for cluster CS4.

Info for manual annotations of cluster CV:

•Start number 7 was manually annotated 1 time for cluster CV.

Info for manual annotations of cluster CY:

•Start number 9 was manually annotated 1 time for cluster CY.

Info for manual annotations of cluster CZ7:

•Start number 9 was manually annotated 1 time for cluster CZ7.

Info for manual annotations of cluster DF:

•Start number 2 was manually annotated 1 time for cluster DF.

Info for manual annotations of cluster DF3:

•Start number 5 was manually annotated 1 time for cluster DF3.

Gene Information:

Gene: ASerpRocky 35 Start: 41389, Stop: 41841, Start Num: 10

Candidate Starts for ASerpRocky 35:

(3, 41335), (Start: 10 @41389 has 9 MA's), (20, 41494), (26, 41557), (31, 41611), (33, 41623), (38, 41647), (48, 41743), (51, 41770), (52, 41779),

Gene: Benczkowski14 35 Start: 41422, Stop: 41874, Start Num: 10

Candidate Starts for Benczkowski14 35:

(3, 41368), (Start: 10 @41422 has 9 MA's), (20, 41527), (31, 41644), (33, 41656), (38, 41680), (44, 41728), (48, 41776), (51, 41803), (52, 41812),

Gene: Camerico 40 Start: 43256, Stop: 43774, Start Num: 2

Candidate Starts for Camerico 40:

(Start: 2 @43256 has 1 MA's), (8, 43295), (16, 43376), (18, 43382), (22, 43421), (33, 43538), (36, 43550), (37, 43559), (45, 43616),

Gene: Demosthenes_35 Start: 41362, Stop: 41814, Start Num: 10

Candidate Starts for Demosthenes 35:

(3, 41308), (Start: 10 @41362 has 9 MA's), (20, 41467), (26, 41530), (31, 41584), (33, 41596), (37, 41617), (38, 41620), (48, 41716), (51, 41743), (52, 41752),

Gene: GMA3_28 Start: 36944, Stop: 37429, Start Num: 4

Candidate Starts for GMA3 28:

(1, 36902), (4, 36944), (15, 37061), (16, 37064), (19, 37088), (23, 37124), (26, 37160), (33, 37226), (53, 37391),

Gene: Hollow_36 Start: 41768, Stop: 42220, Start Num: 10

Candidate Starts for Hollow_36:

(3, 41714), (Start: 10 @41768 has 9 MA's), (20, 41873), (31, 41990), (33, 42002), (38, 42026), (48, 42122), (51, 42149), (52, 42158),

Gene: Jumbo 37 Start: 42278, Stop: 42769, Start Num: 5

Candidate Starts for Jumbo 37:

(Start: 5 @ 42278 has 1 MA's), (6, 42296), (11, 42338), (16, 42398), (25, 42491), (28, 42530), (33, 42560), (34, 42563), (35, 42566), (40, 42593), (41, 42599), (47, 42665), (53, 42728),

Gene: Katyusha_35 Start: 41422, Stop: 41874, Start Num: 10

Candidate Starts for Katyusha_35:

(3, 41368), (Start: 10 @41422 has 9 MA's), (20, 41527), (31, 41644), (33, 41656), (38, 41680), (44, 41728), (48, 41776), (51, 41803), (52, 41812),

Gene: Kvothe 35 Start: 41321, Stop: 41773, Start Num: 10

Candidate Starts for Kvothe 35:

(3, 41267), (Start: 10 @41321 has 9 MA's), (20, 41426), (31, 41543), (33, 41555), (38, 41579), (44, 41627), (48, 41675), (51, 41702), (52, 41711),

Gene: LonelyBoi_24 Start: 21255, Stop: 21728, Start Num: 9

Candidate Starts for LonelyBoi_24:

(Start: 9 @21255 has 2 MA's), (13, 21315), (14, 21321), (16, 21336), (21, 21378), (28, 21465), (29, 21471), (32, 21489), (33, 21495), (39, 21522), (43, 21543), (49, 21636), (51, 21666),

Gene: Niagara_35 Start: 41379, Stop: 41831, Start Num: 10

Candidate Starts for Niagara 35:

(3, 41325), (Start: 10 @41379 has 9 MA's), (20, 41484), (31, 41601), (33, 41613), (38, 41637), (44, 41685), (48, 41733), (51, 41760), (52, 41769),

Gene: Philon9 37 Start: 42589, Stop: 43029, Start Num: 10

Candidate Starts for Philon9_37:

(3, 42535), (Start: 10 @42589 has 9 MA's), (17, 42661), (20, 42691), (30, 42805), (31, 42808), (33, 42820), (38, 42844), (42, 42859), (46, 42901), (48, 42940), (50, 42961),

Gene: Sidious 23 Start: 20906, Stop: 21379, Start Num: 9

Candidate Starts for Sidious_23:

(Start: 9 @20906 has 2 MA's), (13, 20966), (14, 20972), (16, 20987), (21, 21029), (24, 21050), (28, 21116), (32, 21140), (33, 21146), (43, 21194), (44, 21218), (51, 21317),

Gene: Teatealatte_35 Start: 41324, Stop: 41776, Start Num: 10

Candidate Starts for Teatealatte 35:

(3, 41270), (Start: 10 @41324 has 9 MA's), (20, 41429), (31, 41546), (33, 41558), (38, 41582), (44, 41630), (48, 41678), (51, 41705), (52, 41714),

Gene: Teech_35 Start: 41125, Stop: 41577, Start Num: 10

Candidate Starts for Teech_35:

(3, 41071), (Start: 10 @41125 has 9 MA's), (20, 41230), (31, 41347), (33, 41359), (38, 41383), (44, 41431), (48, 41479), (51, 41506), (52, 41515),

Gene: Tredge 35 Start: 41324, Stop: 41776, Start Num: 10

Candidate Starts for Tredge 35:

(3, 41270), (Start: 10 @41324 has 9 MA's), (20, 41429), (31, 41546), (33, 41558), (38, 41582), (44, 41630), (48, 41678), (51, 41705), (52, 41714),

Gene: Wocket_24 Start: 20570, Stop: 21052, Start Num: 7

Candidate Starts for Wocket_24:

(Start: 7 @ 20570 has 1 MA's), (12, 20609), (14, 20642), (16, 20657), (27, 20786), (31, 20807), (33, 20819), (37, 20840), (43, 20867),