Pham 107295


2
5
6
11
12
14
15
17
20
1: Che9c_28
3
4
5
6
7
9
11
13
14
17
18
19
2: Nazo_29 + 1
1
3
4
5
6
8
11
12
13
14
15
16
17
3: Phayonce_29
1
3
4
5
6
8
10
11
12
13
14
15
16
17
4: ThulaThula_29

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 107295 Report

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

Pham number 107295 has 5 members, 0 are drafts.

Phages represented in each track:
• Track 1 : Che9c_28
• Track 2 : Nazo_29, BigNuz_29
• Track 3 : Phayonce_29
• Track 4 : ThulaThula_29

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

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

Genes that call this "Most Annotated" start:
• BigNuz_29, Che9c_28, Nazo_29, Phayonce_29, ThulaThula_29,

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

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

Summary by start number:

Start 5:
• Found in 5 of 5 (100.0%) of genes in pham
• Manual Annotations of this start: 5 of 5
• Called 100.0% of time when present
• Phage (with cluster) where this start called: BigNuz_29 (P4), Che9c_28 (I2), Nazo_29 (P4), Phayonce_29 (P5), ThulaThula_29 (P5),

Summary by clusters:

There are 3 clusters represented in this pham: I2, P4, P5,

Info for manual annotations of cluster I2:
•Start number 5 was manually annotated 1 time for cluster I2.

Info for manual annotations of cluster P4:
•Start number 5 was manually annotated 2 times for cluster P4.

Info for manual annotations of cluster P5:
•Start number 5 was manually annotated 2 times for cluster P5.

Gene Information:

Gene: BigNuz_29 Start: 26343, Stop: 26825, Start Num: 5
Candidate Starts for BigNuz_29:
(3, 26247), (4, 26265), (Start: 5 @26343 has 5 MA's), (6, 26373), (7, 26379), (9, 26451), (11, 26511), (13, 26529), (14, 26634), (17, 26763), (18, 26787), (19, 26790),

Gene: Che9c_28 Start: 25654, Stop: 26139, Start Num: 5
Candidate Starts for Che9c_28:
(2, 25567), (Start: 5 @25654 has 5 MA's), (6, 25684), (11, 25822), (12, 25828), (14, 25945), (15, 26011), (17, 26074), (20, 26128),

Gene: Nazo_29 Start: 26346, Stop: 26828, Start Num: 5
Candidate Starts for Nazo_29:
(3, 26250), (4, 26268), (Start: 5 @26346 has 5 MA's), (6, 26376), (7, 26382), (9, 26454), (11, 26514), (13, 26532), (14, 26637), (17, 26766), (18, 26790), (19, 26793),

Gene: Phayonce_29 Start: 26266, Stop: 26739, Start Num: 5
Candidate Starts for Phayonce_29:
(1, 25966), (3, 26170), (4, 26188), (Start: 5 @26266 has 5 MA's), (6, 26296), (8, 26365), (11, 26434), (12, 26440), (13, 26452), (14, 26557), (15, 26623), (16, 26647), (17, 26686),

Gene: ThulaThula_29 Start: 26272, Stop: 26745, Start Num: 5
Candidate Starts for ThulaThula_29:
(1, 25972), (3, 26176), (4, 26194), (Start: 5 @26272 has 5 MA's), (6, 26302), (8, 26371), (10, 26389), (11, 26440), (12, 26446), (13, 26458), (14, 26563), (15, 26629), (16, 26653), (17, 26692),