



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 286270 Report

This analysis was run 03/28/26 on database version 641.

Pham number 286270 has 6 members, 4 are drafts.

Phages represented in each track:

- Track 1 : NikkiHotWheels_39, HotPotato_37, Henrique_38
- Track 2 : PeggyLeg03_38
- Track 3 : Pucara_35
- Track 4 : CocoCinnamon_35

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

The start number called the most often in the published annotations is 7, it was called in 2 of the 2 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- CocoCinnamon_35, Henrique_38, HotPotato_37, NikkiHotWheels_39, PeggyLeg03_38,

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

- Pucara_35,

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

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Summary by start number:

Start 7:

- Found in 6 of 6 (100.0%) of genes in pham
- Manual Annotations of this start: 2 of 2
- Called 83.3% of time when present
- Phage (with cluster) where this start called: CocoCinnamon_35 (FO), Henrique_38 (FA), HotPotato_37 (FA), NikkiHotWheels_39 (FA), PeggyLeg03_38 (FA),

Start 8:

- Found in 3 of 6 (50.0%) of genes in pham
- No Manual Annotations of this start.
- Called 33.3% of time when present
- Phage (with cluster) where this start called: Pucara_35 (FA),

Summary by clusters:

There are 2 clusters represented in this pham: FA, FO,

Info for manual annotations of cluster FA:

- Start number 7 was manually annotated 1 time for cluster FA.

Info for manual annotations of cluster FO:

- Start number 7 was manually annotated 1 time for cluster FO.

Gene Information:

Gene: CocoCinnamon_35 Start: 25963, Stop: 25775, Start Num: 7

Candidate Starts for CocoCinnamon_35:

(4, 26035), (5, 26029), (Start: 7 @25963 has 2 MA's), (8, 25939), (9, 25915), (10, 25813),

Gene: Henrique_38 Start: 27474, Stop: 27286, Start Num: 7

Candidate Starts for Henrique_38:

(2, 27588), (4, 27546), (5, 27540), (6, 27513), (Start: 7 @27474 has 2 MA's),

Gene: HotPotato_37 Start: 28038, Stop: 27850, Start Num: 7

Candidate Starts for HotPotato_37:

(2, 28152), (4, 28110), (5, 28104), (6, 28077), (Start: 7 @28038 has 2 MA's),

Gene: NikkiHotWheels_39 Start: 27467, Stop: 27279, Start Num: 7

Candidate Starts for NikkiHotWheels_39:

(2, 27581), (4, 27539), (5, 27533), (6, 27506), (Start: 7 @27467 has 2 MA's),

Gene: PeggyLeg03_38 Start: 28180, Stop: 27992, Start Num: 7

Candidate Starts for PeggyLeg03_38:

(1, 28318), (3, 28270), (4, 28252), (5, 28246), (Start: 7 @28180 has 2 MA's), (8, 28156),

Gene: Pucara_35 Start: 27761, Stop: 27597, Start Num: 8

Candidate Starts for Pucara_35:

(1, 27923), (3, 27875), (4, 27857), (5, 27851), (Start: 7 @27785 has 2 MA's), (8, 27761),