

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 225172 Report

This analysis was run 03/28/25 on database version 593.

Pham number 225172 has 8 members, 3 are drafts.

Phages represented in each track:

- Track 1 : Coriander_54, Doggs_52
- Track 2 : GEazy_61, HannahD_58
- Track 3 : BackstagePass_62
- Track 4 : Schnabeltier_59
- Track 5 : Eyes_54
- Track 6 : TaronosaurasRx_63

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 4 of the 5 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

• BackstagePass_62, Eyes_54, GEazy_61, HannahD_58, Schnabeltier_59, TaronosaurasRx 63,

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:

Coriander_54, Doggs_52,

Summary by start number:

Start 4:

- Found in 2 of 8 (25.0%) of genes in pham
- Manual Annotations of this start: 1 of 5
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Coriander_54 (DB), Doggs_52 (DB),

Start 5:

- Found in 6 of 8 (75.0%) of genes in pham
- Manual Annotations of this start: 4 of 5
- Called 100.0% of time when present

 Phage (with cluster) where this start called: BackstagePass_62 (DB), Eyes_54 (DB), GEazy_61 (DB), HannahD_58 (DB), Schnabeltier_59 (DB), TaronosaurasRx_63 (DB),

Summary by clusters:

There is one cluster represented in this pham: DB

Info for manual annotations of cluster DB:

- •Start number 4 was manually annotated 1 time for cluster DB.
- •Start number 5 was manually annotated 4 times for cluster DB.

Gene Information:

Gene: BackstagePass 62 Start: 39231, Stop: 39605, Start Num: 5

Candidate Starts for BackstagePass 62:

(Start: 5 @ 39231 has 4 MA's), (6, 39261), (11, 39477), (12, 39507), (15, 39573),

Gene: Coriander 54 Start: 38252, Stop: 38635, Start Num: 4

Candidate Starts for Coriander 54:

(Start: 4 @38252 has 1 MA's), (8, 38462), (9, 38483), (11, 38507), (12, 38537), (13, 38549), (15, 38603),

Gene: Doggs_52 Start: 38571, Stop: 38954, Start Num: 4

Candidate Starts for Doggs 52:

(Start: 4 @38571 has 1 MA's), (8, 38781), (9, 38802), (11, 38826), (12, 38856), (13, 38868), (15, 38922),

Gene: Eyes_54 Start: 39592, Stop: 39966, Start Num: 5

Candidate Starts for Eyes 54:

(3, 39448), (Start: 5 @39592 has 4 MA's), (6, 39622), (7, 39700), (8, 39793), (11, 39838), (12, 39868), (15, 39934),

Gene: GEazy_61 Start: 39774, Stop: 40148, Start Num: 5

Candidate Starts for GEazy_61:

(1, 39486), (2, 39513), (Start: 5 @39774 has 4 MA's), (6, 39804), (7, 39882), (8, 39975), (11, 40020), (12, 40050), (15, 40116),

Gene: HannahD 58 Start: 39141, Stop: 39515, Start Num: 5

Candidate Starts for HannahD 58:

(1, 38853), (2, 38880), (Start: 5 @39141 has 4 MA's), (6, 39171), (7, 39249), (8, 39342), (11, 39387), (12, 39417), (15, 39483),

Gene: Schnabeltier_59 Start: 40175, Stop: 40549, Start Num: 5

Candidate Starts for Schnabeltier 59:

(Start: 5 @ 40175 has 4 MA's), (6, 40205), (7, 40283), (8, 40376), (11, 40421), (12, 40451), (15, 40517),

Gene: TaronosaurasRx 63 Start: 40175, Stop: 40549, Start Num: 5

Candidate Starts for TaronosaurasRx 63:

(3, 40028), (Start: 5 @40175 has 4 MA's), (6, 40205), (8, 40376), (10, 40409), (12, 40451), (13, 40463), (14, 40472),