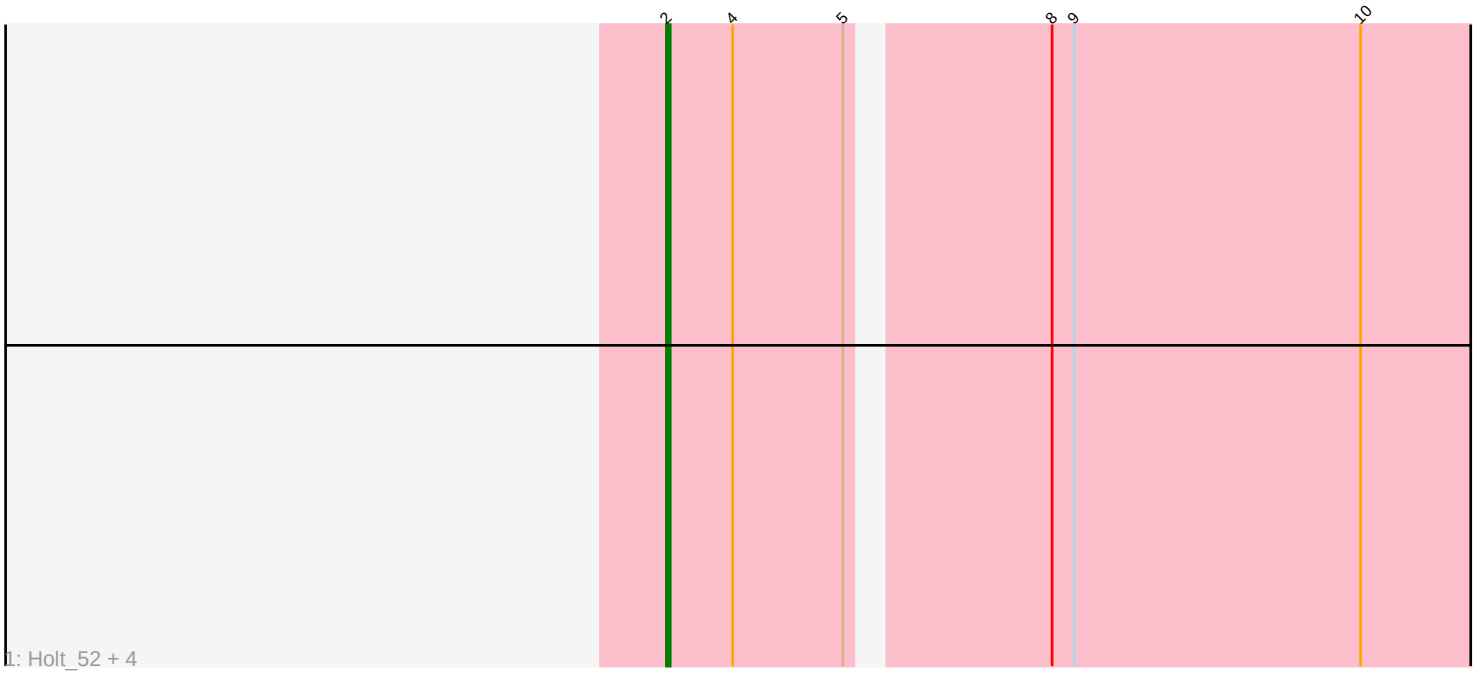
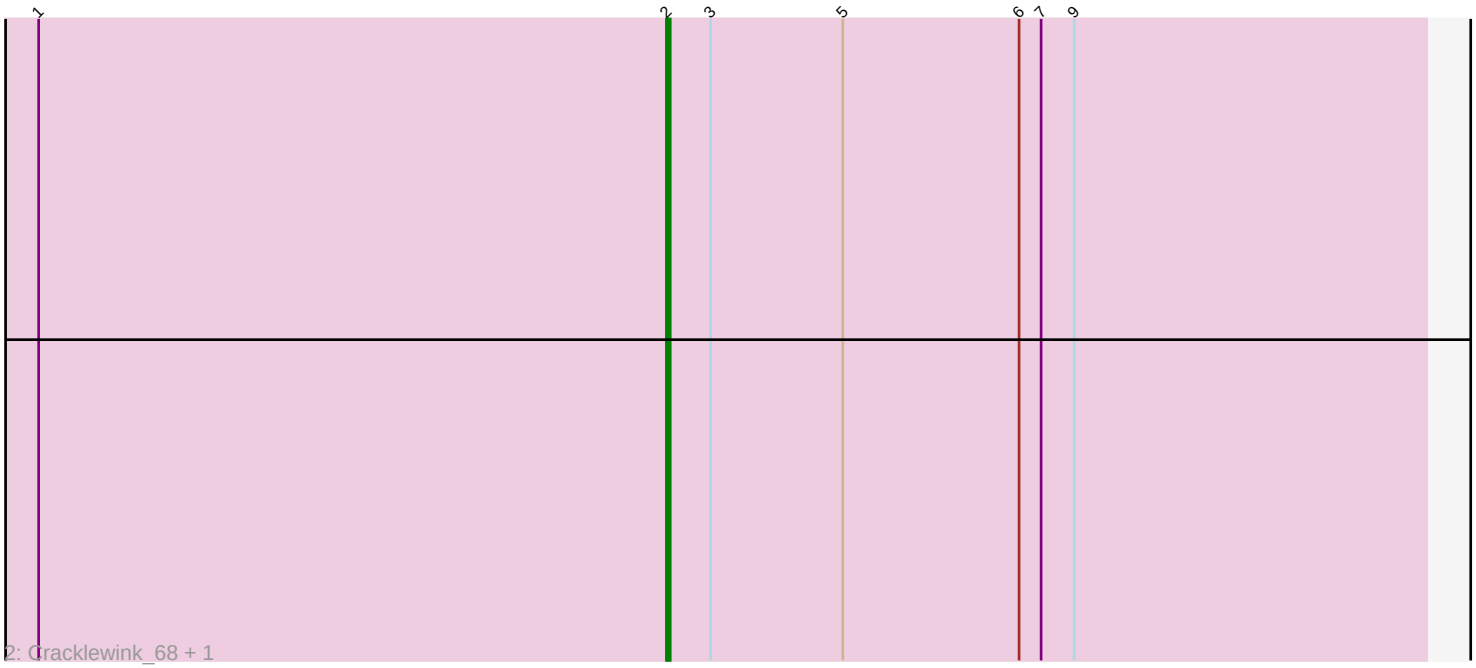
Pham 344757





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 344757 Report

This analysis was run 11/14/26 on database version 672.

Pham number 344757 has 7 members, 0 are drafts.

Phages represented in each track:
• Track 1 : Holt_52, DoctorDiddles_47, ChotaBhai_47, Pat3_45, Amao_47
• Track 2 : Cracklewink_68, Bipper_69

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

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

Genes that call this "Most Annotated" start:
• Amao_47, Bipper_69, ChotaBhai_47, Cracklewink_68, DoctorDiddles_47, Holt_52, Pat3_45,

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 2:
• Found in 7 of 7 (100.0%) of genes in pham
• Manual Annotations of this start: 7 of 7
• Called 100.0% of time when present
• Phage (with cluster) where this start called: Amao_47 (E), Bipper_69 (Y), ChotaBhai_47 (E), Cracklewink_68 (Y), DoctorDiddles_47 (E), Holt_52 (E), Pat3_45 (E),

Summary by clusters:

There are 2 clusters represented in this pham: Y, E,

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

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

Gene Information:

Gene: Amao_47 Start: 35798, Stop: 35586, Start Num: 2
Candidate Starts for Amao_47:
(Start: 2 @35798 has 7 MA's), (4, 35780), (5, 35750), (8, 35702), (9, 35696), (10, 35618),

Gene: Bipper_69 Start: 48431, Stop: 48637, Start Num: 2
Candidate Starts for Bipper_69:
(1, 48260), (Start: 2 @48431 has 7 MA's), (3, 48443), (5, 48479), (6, 48527), (7, 48533), (9, 48542),

Gene: ChotaBhai_47 Start: 35729, Stop: 35517, Start Num: 2
Candidate Starts for ChotaBhai_47:
(Start: 2 @35729 has 7 MA's), (4, 35711), (5, 35681), (8, 35633), (9, 35627), (10, 35549),

Gene: Cracklewink_68 Start: 47465, Stop: 47671, Start Num: 2
Candidate Starts for Cracklewink_68:
(1, 47294), (Start: 2 @47465 has 7 MA's), (3, 47477), (5, 47513), (6, 47561), (7, 47567), (9, 47576),

Gene: DoctorDiddles_47 Start: 35765, Stop: 35553, Start Num: 2
Candidate Starts for DoctorDiddles_47:
(Start: 2 @35765 has 7 MA's), (4, 35747), (5, 35717), (8, 35669), (9, 35663), (10, 35585),

Gene: Holt_52 Start: 36600, Stop: 36388, Start Num: 2
Candidate Starts for Holt_52:
(Start: 2 @36600 has 7 MA's), (4, 36582), (5, 36552), (8, 36504), (9, 36498), (10, 36420),

Gene: Pat3_45 Start: 34930, Stop: 34718, Start Num: 2
Candidate Starts for Pat3_45:
(Start: 2 @34930 has 7 MA's), (4, 34912), (5, 34882), (8, 34834), (9, 34828), (10, 34750),