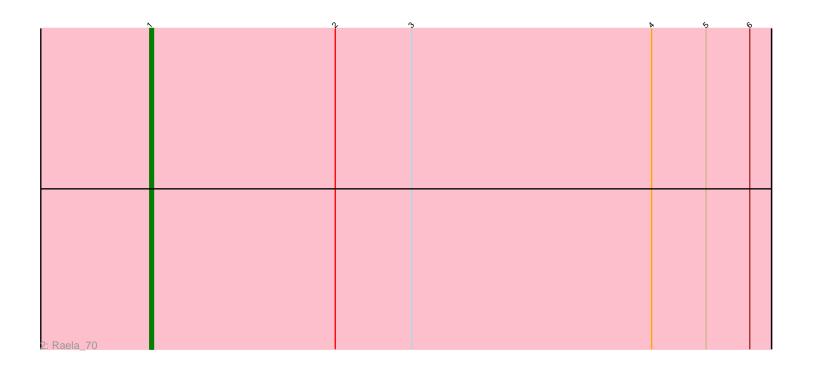
	า	, ,)	b 6	
					-
1: Beelzebub_74 + 6					



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 7199 Report

This analysis was run 04/28/24 on database version 559.

Pham number 7199 has 8 members, 0 are drafts.

Phages represented in each track:

• Track 1 : Beelzebub_74, Huphlepuff_72, Blackbeetle_70, JoieB_71, Poise_70, Clarkson_71, RedRaider77_69

Track 2 : Raela 70

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

The start number called the most often in the published annotations is 1, it was called in 8 of the 8 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

• Beelzebub_74, Blackbeetle_70, Clarkson_71, Huphlepuff_72, JoieB_71, Poise_70, Raela_70, RedRaider77_69,

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 1:

- Found in 8 of 8 (100.0%) of genes in pham
- Manual Annotations of this start: 8 of 8
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Beelzebub_74 (S), Blackbeetle_70 (S), Clarkson_71 (S), Huphlepuff_72 (S), JoieB_71 (S), Poise_70 (S), Raela_70 (S), RedRaider77_69 (S),

Summary by clusters:

There is one cluster represented in this pham: S

Info for manual annotations of cluster S:

•Start number 1 was manually annotated 8 times for cluster S.

Gene Information:

Gene: Beelzebub_74 Start: 40528, Stop: 40698, Start Num: 1

Candidate Starts for Beelzebub 74:

(Start: 1 @ 40528 has 8 MA's), (2, 40579), (3, 40600), (4, 40666), (6, 40693),

Gene: Blackbeetle 70 Start: 39540, Stop: 39710, Start Num: 1

Candidate Starts for Blackbeetle 70:

(Start: 1 @ 39540 has 8 MA's), (2, 39591), (3, 39612), (4, 39678), (6, 39705),

Gene: Clarkson_71 Start: 40256, Stop: 40426, Start Num: 1

Candidate Starts for Clarkson_71:

(Start: 1 @ 40256 has 8 MA's), (2, 40307), (3, 40328), (4, 40394), (6, 40421),

Gene: Huphlepuff_72 Start: 40061, Stop: 40231, Start Num: 1

Candidate Starts for Huphlepuff_72:

(Start: 1 @ 40061 has 8 MA's), (2, 40112), (3, 40133), (4, 40199), (6, 40226),

Gene: JoieB_71 Start: 40494, Stop: 40664, Start Num: 1

Candidate Starts for JoieB_71:

(Start: 1 @ 40494 has 8 MA's), (2, 40545), (3, 40566), (4, 40632), (6, 40659),

Gene: Poise_70 Start: 39540, Stop: 39710, Start Num: 1

Candidate Starts for Poise_70:

(Start: 1 @ 39540 has 8 MA's), (2, 39591), (3, 39612), (4, 39678), (6, 39705),

Gene: Raela_70 Start: 40097, Stop: 40267, Start Num: 1

Candidate Starts for Raela 70:

(Start: 1 @ 40097 has 8 MA's), (2, 40148), (3, 40169), (4, 40235), (5, 40250), (6, 40262),

Gene: RedRaider77 69 Start: 39485, Stop: 39655, Start Num: 1

Candidate Starts for RedRaider77_69:

(Start: 1 @39485 has 8 MA's), (2, 39536), (3, 39557), (4, 39623), (6, 39650),