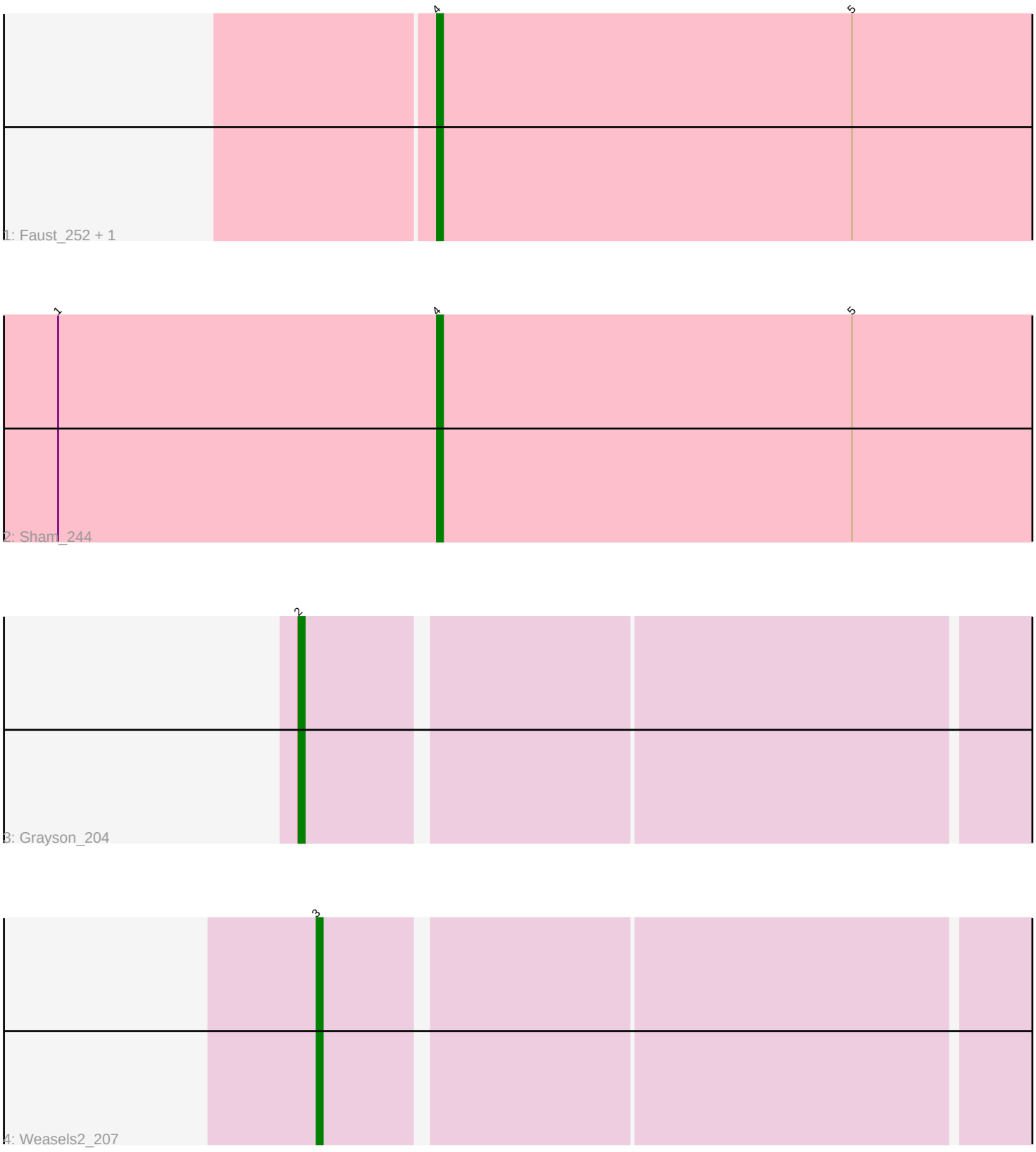
Pham 348453



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 348453 Report

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

Pham number 348453 has 5 members, 1 are drafts.

Phages represented in each track:
• Track 1 : Faust_252, SeresaTree_256
• Track 2 : Sham_244
• Track 3 : Grayson_204
• Track 4 : Weasels2_207

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

The start number called the most often in the published annotations is 4, it was called in 2 of the 4 non-draft genes in the pham.

Genes that call this "Most Annotated" start:
• Faust_252, SeresaTree_256, Sham_244,

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

Genes that do not have the "Most Annotated" start:
• Grayson_204, Weasels2_207,

Summary by start number:

Start 2:
• Found in 1 of 5 (20.0%) of genes in pham
• Manual Annotations of this start: 1 of 4
• Called 100.0% of time when present
• Phage (with cluster) where this start called: Grayson_204 (CB),

Start 3:
• Found in 1 of 5 (20.0%) of genes in pham
• Manual Annotations of this start: 1 of 4
• Called 100.0% of time when present
• Phage (with cluster) where this start called: Weasels2_207 (CB),

Start 4:

• Found in 3 of 5 (60.0%) of genes in pham
• Manual Annotations of this start: 2 of 4
• Called 100.0% of time when present
• Phage (with cluster) where this start called: Faust_252 (BK1), SeresaTree_256 (BK1), Sham_244 (BK1),

Summary by clusters:

There are 2 clusters represented in this pham: CB, BK1,

Info for manual annotations of cluster BK1:
•Start number 4 was manually annotated 2 times for cluster BK1.

Info for manual annotations of cluster CB:
•Start number 2 was manually annotated 1 time for cluster CB.
•Start number 3 was manually annotated 1 time for cluster CB.

Gene Information:

Gene: Faust_252 Start: 121119, Stop: 121268, Start Num: 4
Candidate Starts for Faust_252:
(Start: 4 @121119 has 2 MA's), (5, 121188),

Gene: Grayson_204 Start: 104125, Stop: 104274, Start Num: 2
Candidate Starts for Grayson_204:
(Start: 2 @104125 has 1 MA's),

Gene: SeresaTree_256 Start: 121344, Stop: 121493, Start Num: 4
Candidate Starts for SeresaTree_256:
(Start: 4 @121344 has 2 MA's), (5, 121413),

Gene: Sham_244 Start: 121606, Stop: 121755, Start Num: 4
Candidate Starts for Sham_244:
(1, 121543), (Start: 4 @121606 has 2 MA's), (5, 121675),

Gene: Weasels2_207 Start: 107383, Stop: 107529, Start Num: 3
Candidate Starts for Weasels2_207:
(Start: 3 @107383 has 1 MA's),