Pham 7217

1: Katyusha_43 + 5

2: Niagara_43

3: Hollow_44
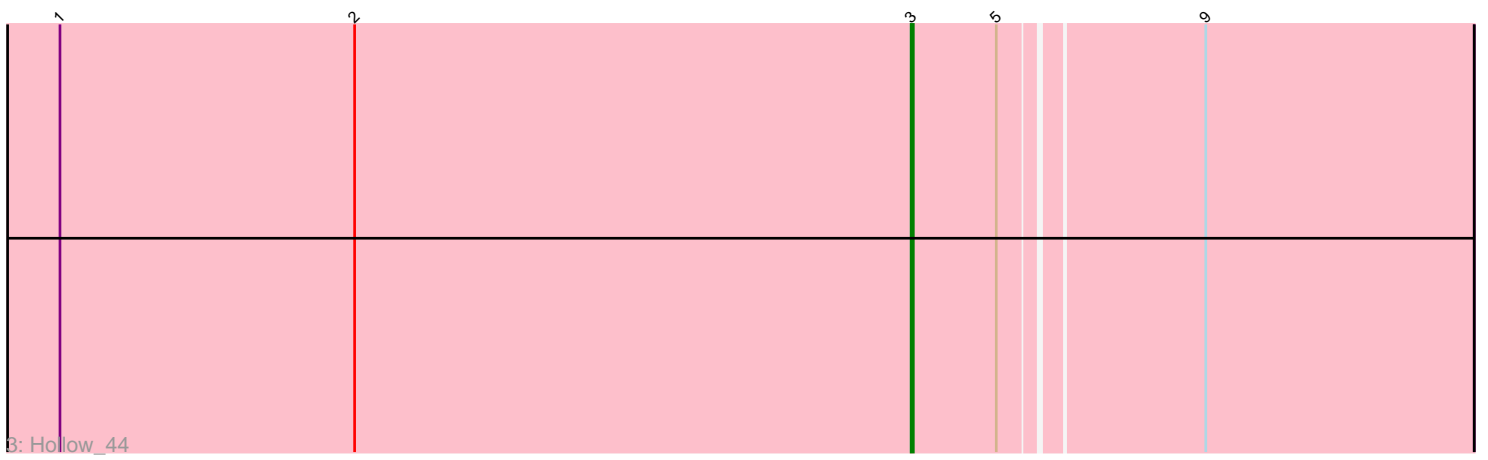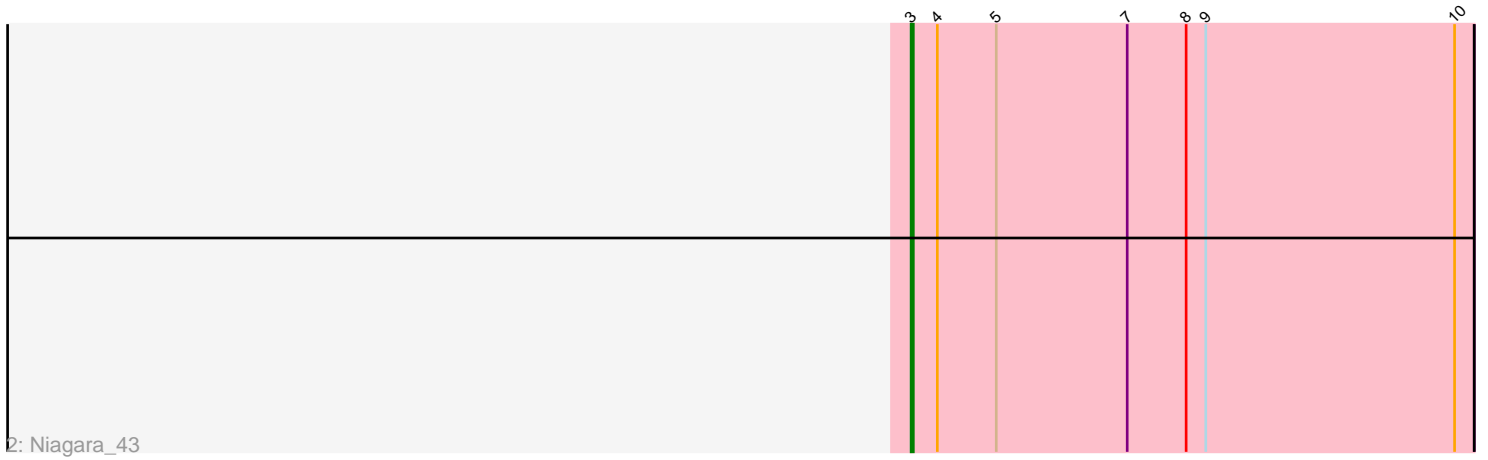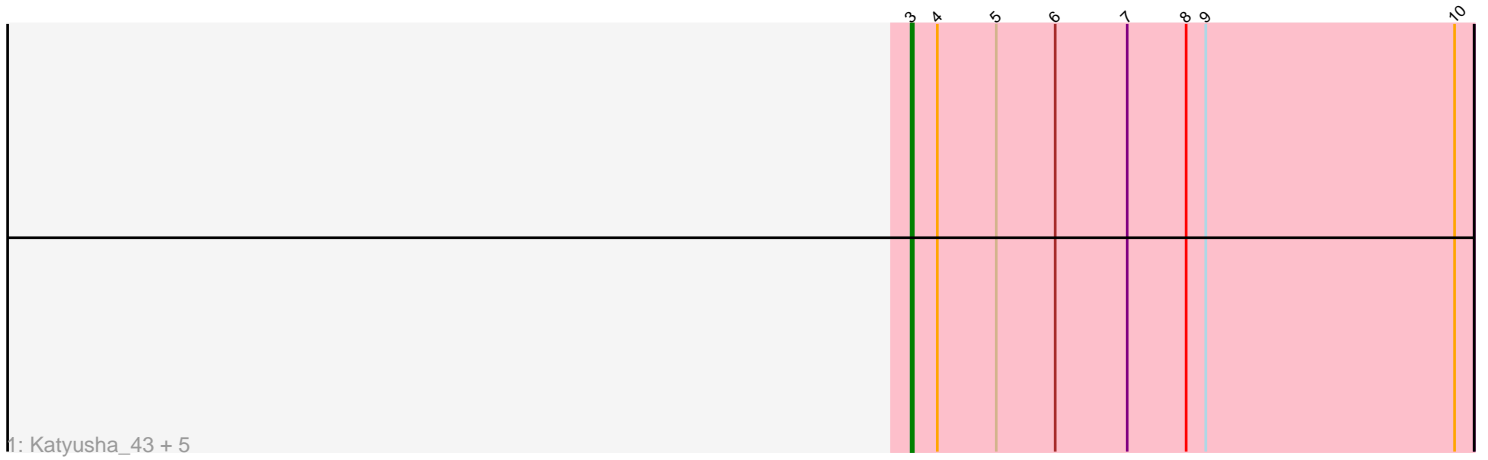
Note: Tracks are now grouped by subcluster and scaled. Switching in subcluster is indicated by changes in track color. Track scale is now set by default to display the region 30 bp upstream of start 1 to 30 bp downstream of the last possible start. If this default region is judged to be packed too tightly 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 7217 Report

This analysis was run 07/09/24 on database version 566.

Pham number 7217 has 8 members, 1 are drafts.

Phages represented in each track:
• Track 1 : Katyusha_43, Teatealatte_44, Tredge_44, Benczkowski14_43, Teech_44, Kvothe_43
• Track 2 : Niagara_43
• Track 3 : Hollow_44

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

The start number called the most often in the published annotations is 3, it was called in 7 of the 7 non-draft genes in the pham.

Genes that call this "Most Annotated" start:
• Benczkowski14_43, Hollow_44, Katyusha_43, Kvothe_43, Niagara_43, Teatealatte_44, Teech_44, Tredge_44,

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 3:
• Found in 8 of 8 (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: Benczkowski14_43 (CS4), Hollow_44 (CS4), Katyusha_43 (CS4), Kvothe_43 (CS4), Niagara_43 (CS4), Teatealatte_44 (CS4), Teech_44 (CS4), Tredge_44 (CS4),

Summary by clusters:

There is one cluster represented in this pham: CS4

Info for manual annotations of cluster CS4:
•Start number 3 was manually annotated 7 times for cluster CS4.

Gene Information:

Gene: Benczkowski14_43 Start: 45905, Stop: 45648, Start Num: 3
Candidate Starts for Benczkowski14_43:
(Start: 3 @45905 has 7 MA's), (4, 45893), (5, 45866), (6, 45839), (7, 45806), (8, 45779), (9, 45770),
(10, 45656),

Gene: Hollow_44 Start: 46242, Stop: 45994, Start Num: 3
Candidate Starts for Hollow_44:
(1, 46632), (2, 46497), (Start: 3 @46242 has 7 MA's), (5, 46203), (9, 46116),

Gene: Katyusha_43 Start: 45905, Stop: 45648, Start Num: 3
Candidate Starts for Katyusha_43:
(Start: 3 @45905 has 7 MA's), (4, 45893), (5, 45866), (6, 45839), (7, 45806), (8, 45779), (9, 45770),
(10, 45656),

Gene: Kvothe_43 Start: 45804, Stop: 45547, Start Num: 3
Candidate Starts for Kvothe_43:
(Start: 3 @45804 has 7 MA's), (4, 45792), (5, 45765), (6, 45738), (7, 45705), (8, 45678), (9, 45669),
(10, 45555),

Gene: Niagara_43 Start: 45883, Stop: 45626, Start Num: 3
Candidate Starts for Niagara_43:
(Start: 3 @45883 has 7 MA's), (4, 45871), (5, 45844), (7, 45784), (8, 45757), (9, 45748), (10, 45634),

Gene: Teatealatte_44 Start: 45807, Stop: 45550, Start Num: 3
Candidate Starts for Teatealatte_44:
(Start: 3 @45807 has 7 MA's), (4, 45795), (5, 45768), (6, 45741), (7, 45708), (8, 45681), (9, 45672),
(10, 45558),

Gene: Teech_44 Start: 45608, Stop: 45351, Start Num: 3
Candidate Starts for Teech_44:
(Start: 3 @45608 has 7 MA's), (4, 45596), (5, 45569), (6, 45542), (7, 45509), (8, 45482), (9, 45473),
(10, 45359),

Gene: Tredge_44 Start: 45807, Stop: 45550, Start Num: 3
Candidate Starts for Tredge_44:
(Start: 3 @45807 has 7 MA's), (4, 45795), (5, 45768), (6, 45741), (7, 45708), (8, 45681), (9, 45672),
(10, 45558),