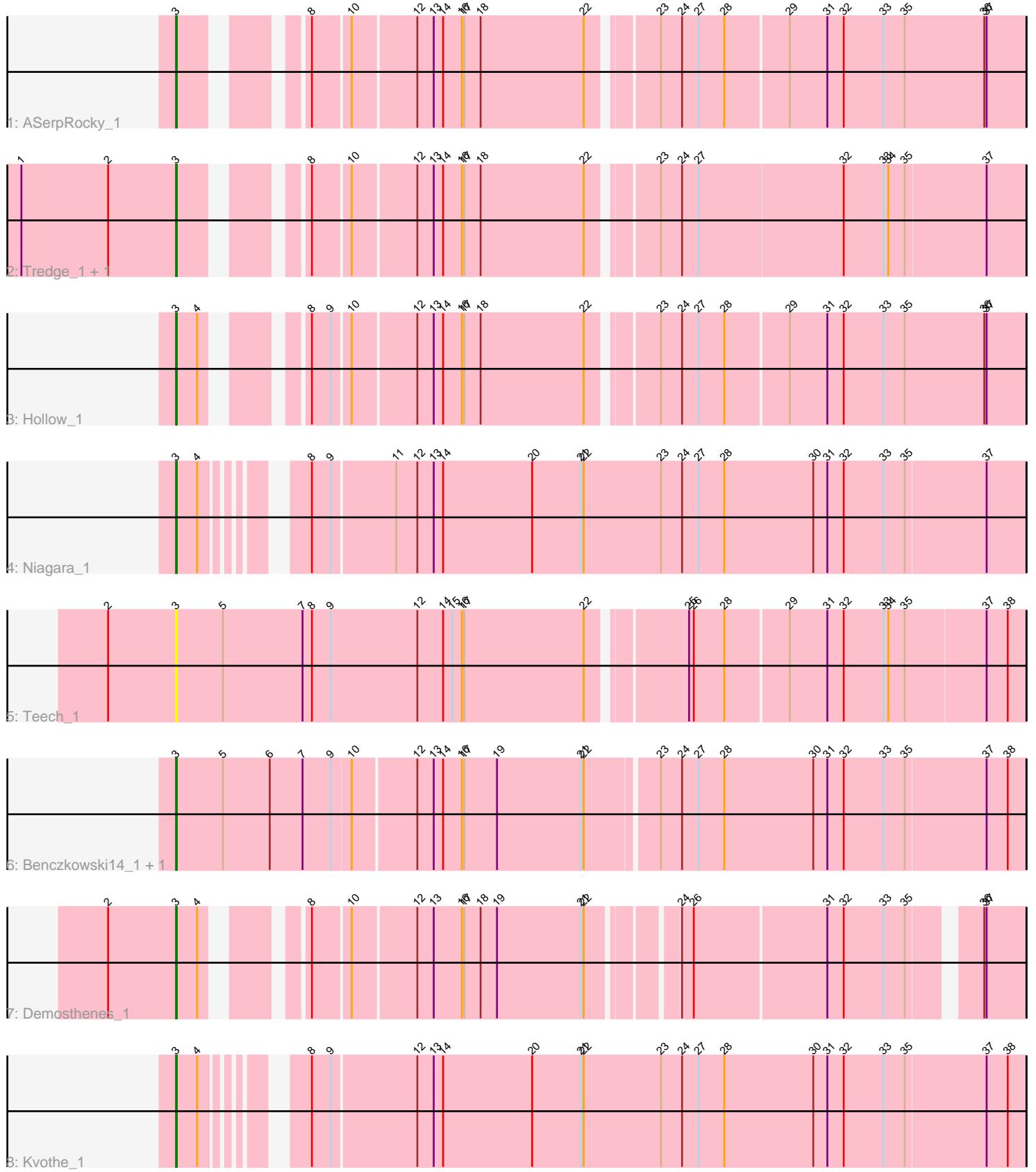


Pham 6120



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 6120 Report

This analysis was run 04/05/24 on database version 557.

Pham number 6120 has 10 members, 1 are drafts.

Phages represented in each track:

- Track 1 : ASerpRocky_1
- Track 2 : Tredge_1, Teatealatte_1
- Track 3 : Hollow_1
- Track 4 : Niagara_1
- Track 5 : Teech_1
- Track 6 : Benczkowski14_1, Katyusha_1
- Track 7 : Demosthenes_1
- Track 8 : Kvothe_1

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 9 of the 9 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- ASerpRocky_1, Benczkowski14_1, Demosthenes_1, Hollow_1, Katyusha_1, Kvothe_1, Niagara_1, Teatealatte_1, Teech_1, Tredge_1,

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 10 of 10 (100.0%) of genes in pham
- Manual Annotations of this start: 9 of 9
- Called 100.0% of time when present
- Phage (with cluster) where this start called: ASerpRocky_1 (CS4), Benczkowski14_1 (CS4), Demosthenes_1 (CS4), Hollow_1 (CS4), Katyusha_1 (CS4), Kvothe_1 (CS4), Niagara_1 (CS4), Teatealatte_1 (CS4), Teech_1 (CS4), Tredge_1 (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 9 times for cluster CS4.

Gene Information:

Gene: ASerpRocky_1 Start: 2858, Stop: 1872, Start Num: 3

Candidate Starts for ASerpRocky_1:

(Start: 3 @2858 has 9 MA's), (8, 2744), (10, 2699), (12, 2621), (13, 2600), (14, 2588), (16, 2564), (17, 2561), (18, 2540), (22, 2408), (23, 2333), (24, 2306), (27, 2285), (28, 2252), (29, 2174), (31, 2126), (32, 2105), (33, 2054), (35, 2027), (36, 1925), (37, 1922),

Gene: Benczkowski14_1 Start: 2921, Stop: 1857, Start Num: 3

Candidate Starts for Benczkowski14_1:

(Start: 3 @2921 has 9 MA's), (5, 2861), (6, 2801), (7, 2759), (9, 2723), (10, 2699), (12, 2621), (13, 2600), (14, 2588), (16, 2564), (17, 2561), (19, 2519), (21, 2411), (22, 2408), (23, 2321), (24, 2294), (27, 2273), (28, 2240), (30, 2126), (31, 2108), (32, 2087), (33, 2036), (35, 2009), (37, 1907), (38, 1880),

Gene: Demosthenes_1 Start: 2818, Stop: 1865, Start Num: 3

Candidate Starts for Demosthenes_1:

(2, 2905), (Start: 3 @2818 has 9 MA's), (4, 2791), (8, 2704), (10, 2659), (12, 2581), (13, 2560), (16, 2524), (17, 2521), (18, 2500), (19, 2479), (21, 2371), (22, 2368), (24, 2272), (26, 2257), (31, 2092), (32, 2071), (33, 2020), (35, 1993), (36, 1918), (37, 1915),

Gene: Hollow_1 Start: 2856, Stop: 1870, Start Num: 3

Candidate Starts for Hollow_1:

(Start: 3 @2856 has 9 MA's), (4, 2829), (8, 2742), (9, 2718), (10, 2697), (12, 2619), (13, 2598), (14, 2586), (16, 2562), (17, 2559), (18, 2538), (22, 2406), (23, 2331), (24, 2304), (27, 2283), (28, 2250), (29, 2172), (31, 2124), (32, 2103), (33, 2052), (35, 2025), (36, 1923), (37, 1920),

Gene: Katyusha_1 Start: 2921, Stop: 1857, Start Num: 3

Candidate Starts for Katyusha_1:

(Start: 3 @2921 has 9 MA's), (5, 2861), (6, 2801), (7, 2759), (9, 2723), (10, 2699), (12, 2621), (13, 2600), (14, 2588), (16, 2564), (17, 2561), (19, 2519), (21, 2411), (22, 2408), (23, 2321), (24, 2294), (27, 2273), (28, 2240), (30, 2126), (31, 2108), (32, 2087), (33, 2036), (35, 2009), (37, 1907), (38, 1880),

Gene: Kvothe_1 Start: 2876, Stop: 1857, Start Num: 3

Candidate Starts for Kvothe_1:

(Start: 3 @2876 has 9 MA's), (4, 2849), (8, 2762), (9, 2738), (12, 2633), (13, 2612), (14, 2600), (20, 2486), (21, 2423), (22, 2420), (23, 2321), (24, 2294), (27, 2273), (28, 2240), (30, 2126), (31, 2108), (32, 2087), (33, 2036), (35, 2009), (37, 1907), (38, 1880),

Gene: Niagara_1 Start: 2878, Stop: 1859, Start Num: 3

Candidate Starts for Niagara_1:

(Start: 3 @2878 has 9 MA's), (4, 2851), (8, 2764), (9, 2740), (11, 2662), (12, 2635), (13, 2614), (14, 2602), (20, 2488), (21, 2425), (22, 2422), (23, 2323), (24, 2296), (27, 2275), (28, 2242), (30, 2128), (31, 2110), (32, 2089), (33, 2038), (35, 2011), (37, 1909),

Gene: Teatealatte_1 Start: 2841, Stop: 1855, Start Num: 3

Candidate Starts for Teatealatte_1:

(1, 3039), (2, 2928), (Start: 3 @2841 has 9 MA's), (8, 2727), (10, 2682), (12, 2604), (13, 2583), (14, 2571), (16, 2547), (17, 2544), (18, 2523), (22, 2391), (23, 2316), (24, 2289), (27, 2268), (32, 2085), (33, 2034), (34, 2028), (35, 2007), (37, 1905),

Gene: Teech_1 Start: 2913, Stop: 1858, Start Num: 3

Candidate Starts for Teech_1:

(2, 3000), (Start: 3 @2913 has 9 MA's), (5, 2853), (7, 2751), (8, 2739), (9, 2715), (12, 2604), (14, 2571), (15, 2559), (16, 2547), (17, 2544), (22, 2391), (25, 2280), (26, 2274), (28, 2235), (29, 2157), (31, 2109), (32, 2088), (33, 2037), (34, 2031), (35, 2010), (37, 1908), (38, 1881),

Gene: Tredge_1 Start: 2841, Stop: 1855, Start Num: 3

Candidate Starts for Tredge_1:

(1, 3039), (2, 2928), (Start: 3 @2841 has 9 MA's), (8, 2727), (10, 2682), (12, 2604), (13, 2583), (14, 2571), (16, 2547), (17, 2544), (18, 2523), (22, 2391), (23, 2316), (24, 2289), (27, 2268), (32, 2085), (33, 2034), (34, 2028), (35, 2007), (37, 1905),