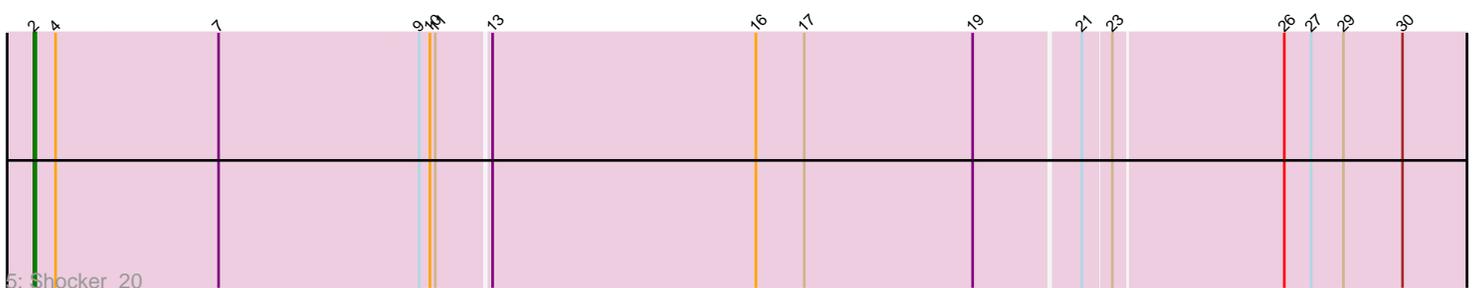
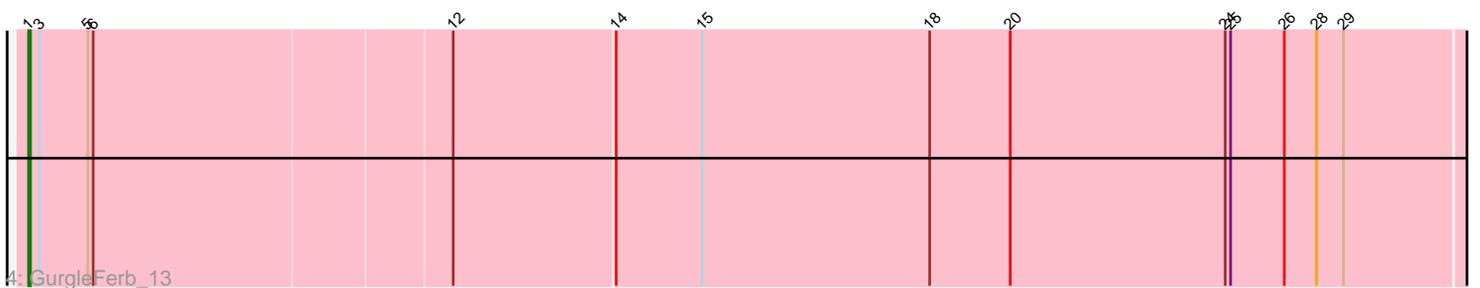
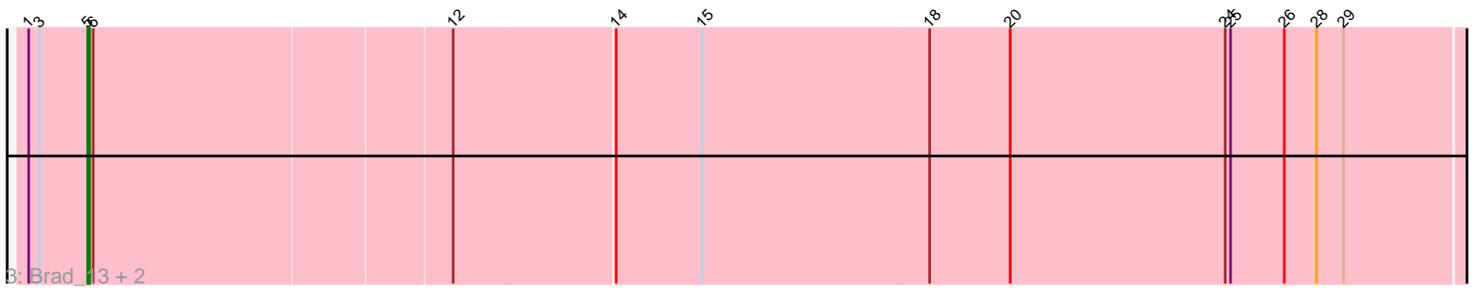
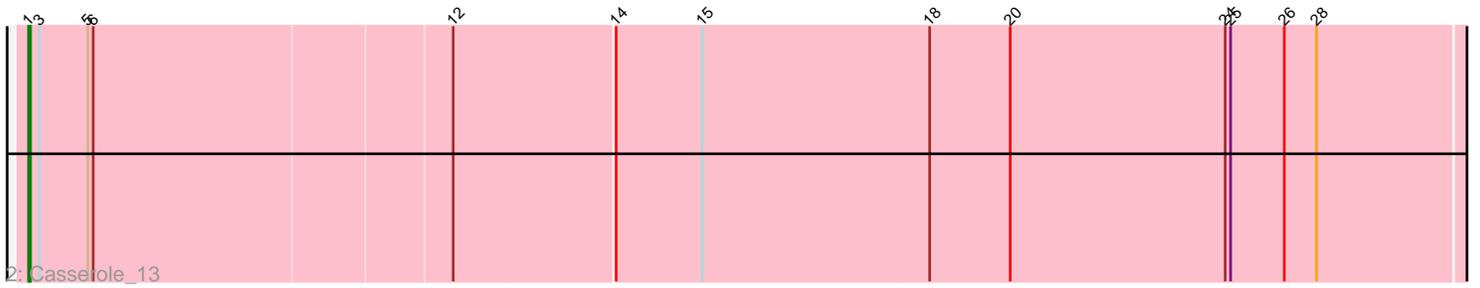
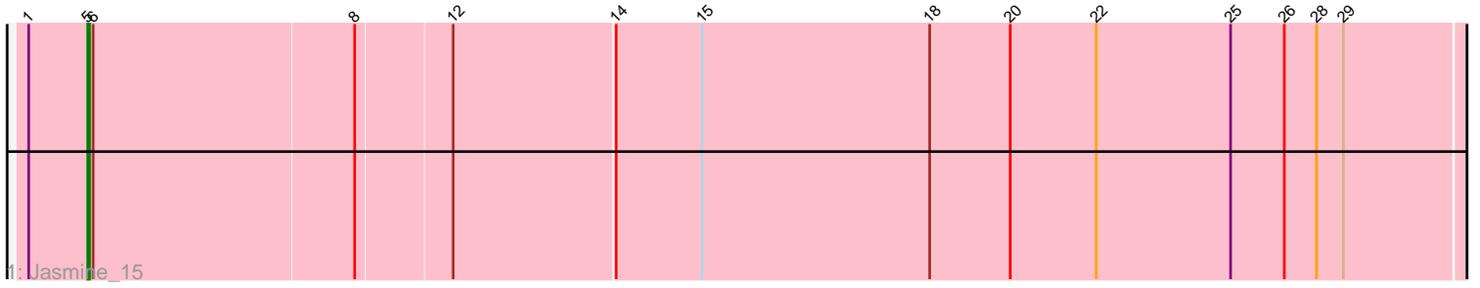


Zoomed Pham 7115



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 7115 Report

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

Pham number 7115 has 7 members, 0 are drafts.

Phages represented in each track:

- Track 1 : Jasmine_15
- Track 2 : Casserole_13
- Track 3 : Brad_13, Nellie_13, Adat_13
- Track 4 : GurgleFerb_13
- Track 5 : Shocker_20

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

The start number called the most often in the published annotations is 5, it was called in 4 of the 7 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Adat_13, Brad_13, Jasmine_15, Nellie_13,

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

- Casserole_13, GurgleFerb_13,

Genes that do not have the "Most Annotated" start:

- Shocker_20,

Summary by start number:

Start 1:

- Found in 6 of 7 (85.7%) of genes in pham
- Manual Annotations of this start: 2 of 7
- Called 33.3% of time when present
- Phage (with cluster) where this start called: Casserole_13 (AV), GurgleFerb_13 (AV),

Start 2:

- Found in 1 of 7 (14.3%) of genes in pham
- Manual Annotations of this start: 1 of 7
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Shocker_20 (singleton),

Start 5:

- Found in 6 of 7 (85.7%) of genes in pham
- Manual Annotations of this start: 4 of 7
- Called 66.7% of time when present
- Phage (with cluster) where this start called: Adat_13 (AV), Brad_13 (AV), Jasmine_15 (AV), Nellie_13 (AV),

Summary by clusters:

There are 2 clusters represented in this pham: singleton, AV,

Info for manual annotations of cluster AV:

- Start number 1 was manually annotated 2 times for cluster AV.
- Start number 5 was manually annotated 4 times for cluster AV.

Gene Information:

Gene: Adat_13 Start: 7343, Stop: 9493, Start Num: 5

Candidate Starts for Adat_13:

(Start: 1 @7310 has 2 MA's), (3, 7316), (Start: 5 @7343 has 4 MA's), (6, 7346), (12, 7541), (14, 7628), (15, 7676), (18, 7802), (20, 7847), (24, 7967), (25, 7970), (26, 8000), (28, 8018), (29, 8033), (34, 8162), (37, 8219), (42, 8357), (50, 8540), (51, 8576), (54, 8666), (55, 8690), (57, 8747), (58, 8753), (62, 8795), (63, 8873), (64, 8885), (65, 8906), (66, 8930), (67, 8969), (68, 8987), (70, 9044), (75, 9113), (77, 9146), (80, 9179), (81, 9200), (84, 9245), (85, 9260), (89, 9344), (90, 9347), (92, 9365), (93, 9368), (96, 9395), (97, 9404), (98, 9416), (99, 9431), (101, 9452), (102, 9461),

Gene: Brad_13 Start: 7341, Stop: 9491, Start Num: 5

Candidate Starts for Brad_13:

(Start: 1 @7308 has 2 MA's), (3, 7314), (Start: 5 @7341 has 4 MA's), (6, 7344), (12, 7539), (14, 7626), (15, 7674), (18, 7800), (20, 7845), (24, 7965), (25, 7968), (26, 7998), (28, 8016), (29, 8031), (34, 8160), (37, 8217), (42, 8355), (50, 8538), (51, 8574), (54, 8664), (55, 8688), (57, 8745), (58, 8751), (62, 8793), (63, 8871), (64, 8883), (65, 8904), (66, 8928), (67, 8967), (68, 8985), (70, 9042), (75, 9111), (77, 9144), (80, 9177), (81, 9198), (84, 9243), (85, 9258), (89, 9342), (90, 9345), (92, 9363), (93, 9366), (96, 9393), (97, 9402), (98, 9414), (99, 9429), (101, 9450), (102, 9459),

Gene: Casserole_13 Start: 7374, Stop: 9557, Start Num: 1

Candidate Starts for Casserole_13:

(Start: 1 @7374 has 2 MA's), (3, 7380), (Start: 5 @7407 has 4 MA's), (6, 7410), (12, 7605), (14, 7692), (15, 7740), (18, 7866), (20, 7911), (24, 8031), (25, 8034), (26, 8064), (28, 8082), (34, 8226), (37, 8283), (42, 8421), (49, 8550), (50, 8604), (51, 8640), (54, 8730), (55, 8754), (57, 8811), (62, 8859), (63, 8937), (64, 8949), (65, 8970), (66, 8994), (67, 9033), (68, 9051), (70, 9108), (75, 9177), (77, 9210), (80, 9243), (81, 9264), (84, 9309), (85, 9324), (89, 9408), (90, 9411), (92, 9429), (93, 9432), (96, 9459), (97, 9468), (98, 9480), (99, 9495), (101, 9516), (102, 9525),

Gene: GurpleFerb_13 Start: 7309, Stop: 9492, Start Num: 1

Candidate Starts for GurpleFerb_13:

(Start: 1 @7309 has 2 MA's), (3, 7315), (Start: 5 @7342 has 4 MA's), (6, 7345), (12, 7540), (14, 7627), (15, 7675), (18, 7801), (20, 7846), (24, 7966), (25, 7969), (26, 7999), (28, 8017), (29, 8032), (34, 8161), (37, 8218), (42, 8356), (50, 8539), (51, 8575), (54, 8665), (55, 8689), (57, 8746), (58, 8752), (62, 8794), (63, 8872), (64, 8884), (65, 8905), (66, 8929), (67, 8968), (68, 8986), (70, 9043), (75,

9112), (77, 9145), (80, 9178), (81, 9199), (84, 9244), (85, 9259), (89, 9343), (90, 9346), (92, 9364), (93, 9367), (96, 9394), (97, 9403), (98, 9415), (99, 9430), (101, 9451), (102, 9460),

Gene: Jasmine_15 Start: 7429, Stop: 9579, Start Num: 5

Candidate Starts for Jasmine_15:

(Start: 1 @7396 has 2 MA's), (Start: 5 @7429 has 4 MA's), (6, 7432), (8, 7576), (12, 7627), (14, 7714), (15, 7762), (18, 7888), (20, 7933), (22, 7981), (25, 8056), (26, 8086), (28, 8104), (29, 8119), (34, 8248), (35, 8251), (37, 8305), (42, 8443), (46, 8536), (50, 8626), (51, 8662), (54, 8752), (55, 8776), (57, 8833), (58, 8839), (60, 8860), (62, 8881), (63, 8959), (64, 8971), (65, 8992), (66, 9016), (67, 9055), (68, 9073), (70, 9130), (71, 9136), (72, 9178), (75, 9199), (76, 9220), (80, 9265), (81, 9286), (84, 9331), (85, 9346), (89, 9430), (90, 9433), (92, 9451), (93, 9454), (96, 9481), (97, 9490), (98, 9502), (99, 9517), (101, 9538), (102, 9547),

Gene: Nellie_13 Start: 7343, Stop: 9493, Start Num: 5

Candidate Starts for Nellie_13:

(Start: 1 @7310 has 2 MA's), (3, 7316), (Start: 5 @7343 has 4 MA's), (6, 7346), (12, 7541), (14, 7628), (15, 7676), (18, 7802), (20, 7847), (24, 7967), (25, 7970), (26, 8000), (28, 8018), (29, 8033), (34, 8162), (37, 8219), (42, 8357), (50, 8540), (51, 8576), (54, 8666), (55, 8690), (57, 8747), (58, 8753), (62, 8795), (63, 8873), (64, 8885), (65, 8906), (66, 8930), (67, 8969), (68, 8987), (70, 9044), (75, 9113), (77, 9146), (80, 9179), (81, 9200), (84, 9245), (85, 9260), (89, 9344), (90, 9347), (92, 9365), (93, 9368), (96, 9395), (97, 9404), (98, 9416), (99, 9431), (101, 9452), (102, 9461),

Gene: Shocker_20 Start: 8991, Stop: 11159, Start Num: 2

Candidate Starts for Shocker_20:

(Start: 2 @8991 has 1 MA's), (4, 9003), (7, 9093), (9, 9204), (10, 9210), (11, 9213), (13, 9240), (16, 9387), (17, 9414), (19, 9507), (21, 9564), (23, 9579), (26, 9672), (27, 9687), (29, 9705), (30, 9738), (31, 9777), (32, 9828), (33, 9831), (36, 9849), (38, 9897), (39, 9909), (40, 9915), (41, 10014), (43, 10068), (44, 10083), (45, 10098), (47, 10122), (48, 10134), (52, 10281), (53, 10308), (56, 10392), (59, 10437), (60, 10443), (61, 10461), (67, 10638), (69, 10677), (71, 10719), (73, 10773), (74, 10785), (75, 10788), (76, 10809), (78, 10830), (79, 10842), (82, 10878), (83, 10884), (86, 10938), (87, 10971), (88, 11013), (91, 11028), (93, 11043), (94, 11052), (95, 11064), (100, 11118),