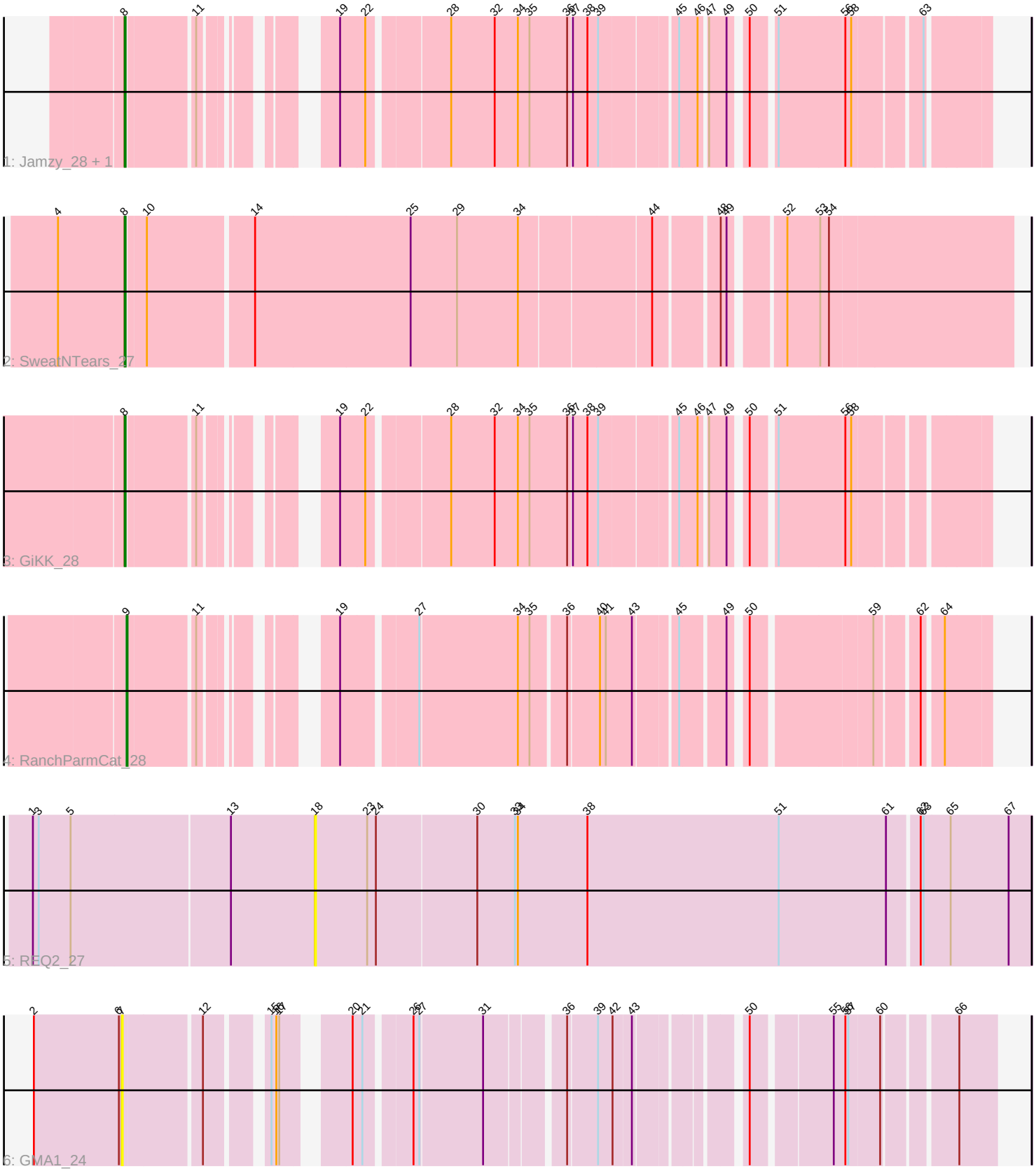
# Pham 348345


1: Jamzy_28 + 1
2: SweatNTears_27
3: GiKK_28
4: RanchParmCat_28
5: REQ2_27
6: GMA1_24

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 348345 Report

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

Pham number 348345 has 7 members, 2 are drafts.

Phages represented in each track:
• Track 1 : Jamzy_28, Button_26
• Track 2 : SweatNTears_27
• Track 3 : GiKK_28
• Track 4 : RanchParmCat_28
• Track 5 : REQ2_27
• Track 6 : GMA1_24

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

The start number called the most often in the published annotations is 8, it was called in 4 of the 5 non-draft genes in the pham.

Genes that call this "Most Annotated" start:
• Button_26, GiKK_28, Jamzy_28, SweatNTears_27,

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

Genes that do not have the "Most Annotated" start:
• GMA1_24, REQ2_27, RanchParmCat_28,

**Summary by start number:**

Start 7:
• Found in 1 of 7 ( 14.3% ) of genes in pham
• No Manual Annotations of this start.
• Called 100.0% of time when present
• Phage (with cluster) where this start called: GMA1_24 (singleton),

Start 8:
• Found in 4 of 7 ( 57.1% ) of genes in pham
• Manual Annotations of this start: 4 of 5
• Called 100.0% of time when present

• Phage (with cluster) where this start called: Button_26 (CT), GiKK_28 (CT), Jamzy_28 (CT), SweatNTears_27 (CT),

Start 9:
• Found in 1 of 7 ( 14.3% ) of genes in pham
• Manual Annotations of this start: 1 of 5
• Called 100.0% of time when present
• Phage (with cluster) where this start called: RanchParmCat_28 (CT),

Start 18:
• Found in 1 of 7 ( 14.3% ) of genes in pham
• No Manual Annotations of this start.
• Called 100.0% of time when present
• Phage (with cluster) where this start called: REQ2_27 (singleton),

**Summary by clusters:**

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

Info for manual annotations of cluster CT:
•Start number 8 was manually annotated 4 times for cluster CT.
•Start number 9 was manually annotated 1 time for cluster CT.

***Gene Information:***

Gene: Button_26 Start: 19490, Stop: 20224, Start Num: 8
Candidate Starts for Button_26:
(Start: 8 @19490 has 4 MA's), (11, 19553), (19, 19640), (22, 19664), (28, 19739), (32, 19784), (34, 19808), (35, 19820), (36, 19859), (37, 19862), (38, 19877), (39, 19886), (45, 19958), (46, 19976), (47, 19982), (49, 20000), (50, 20012), (51, 20033), (56, 20099), (58, 20105), (63, 20165),

Gene: GMA1_24 Start: 19617, Stop: 20363, Start Num: 7
Candidate Starts for GMA1_24:
(2, 19527), (6, 19614), (7, 19617), (12, 19692), (15, 19740), (16, 19743), (17, 19746), (20, 19797), (21, 19806), (26, 19848), (27, 19854), (31, 19917), (36, 19989), (39, 20016), (42, 20031), (43, 20049), (50, 20139), (55, 20214), (56, 20226), (57, 20229), (60, 20259), (66, 20325),

Gene: GiKK_28 Start: 19789, Stop: 20523, Start Num: 8
Candidate Starts for GiKK_28:
(Start: 8 @19789 has 4 MA's), (11, 19852), (19, 19939), (22, 19963), (28, 20038), (32, 20083), (34, 20107), (35, 20119), (36, 20158), (37, 20161), (38, 20176), (39, 20185), (45, 20257), (46, 20275), (47, 20281), (49, 20299), (50, 20311), (51, 20332), (56, 20398), (58, 20404),

Gene: Jamzy_28 Start: 19803, Stop: 20537, Start Num: 8
Candidate Starts for Jamzy_28:
(Start: 8 @19803 has 4 MA's), (11, 19866), (19, 19953), (22, 19977), (28, 20052), (32, 20097), (34, 20121), (35, 20133), (36, 20172), (37, 20175), (38, 20190), (39, 20199), (45, 20271), (46, 20289), (47, 20295), (49, 20313), (50, 20325), (51, 20346), (56, 20412), (58, 20418), (63, 20478),

Gene: REQ2_27 Start: 22841, Stop: 23572, Start Num: 18
Candidate Starts for REQ2_27:

(1, 22556), (3, 22562), (5, 22595), (13, 22754), (18, 22841), (23, 22895), (24, 22904), (30, 23006), (33, 23045), (34, 23048), (38, 23120), (51, 23318), (61, 23429), (62, 23459), (63, 23462), (65, 23489), (67, 23549),

Gene: RanchParmCat_28 Start: 20169, Stop: 20900, Start Num: 9
Candidate Starts for RanchParmCat_28:
(Start: 9 @20169 has 1 MA's), (11, 20232), (19, 20319), (27, 20388), (34, 20487), (35, 20499), (36, 20532), (40, 20562), (41, 20568), (43, 20595), (45, 20634), (49, 20676), (50, 20688), (59, 20802), (62, 20838), (64, 20856),

Gene: SweatNTears_27 Start: 20679, Stop: 21539, Start Num: 8
Candidate Starts for SweatNTears_27:
(4, 20610), (Start: 8 @20679 has 4 MA's), (10, 20697), (14, 20802), (25, 20961), (29, 21009), (34, 21072), (44, 21201), (48, 21261), (49, 21267), (52, 21312), (53, 21345), (54, 21354),