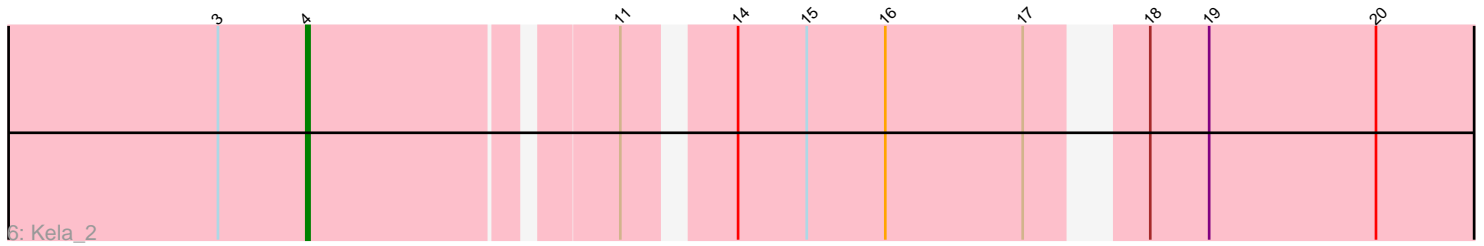
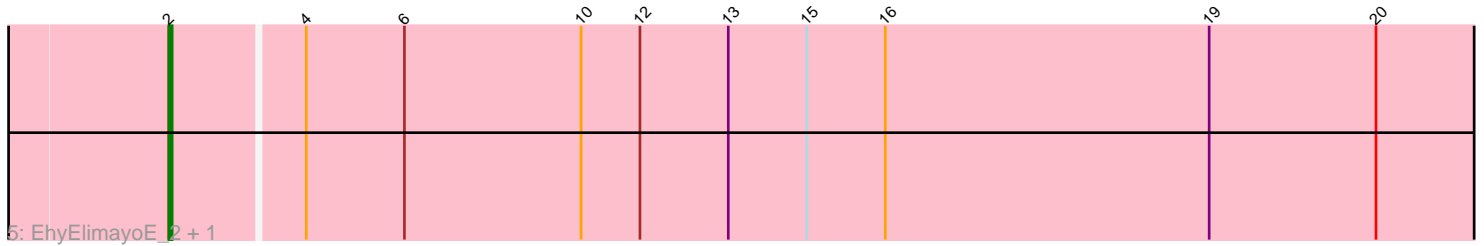
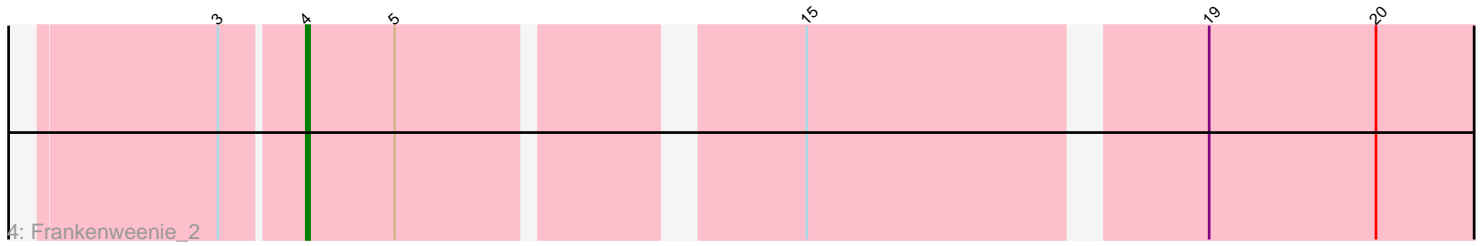
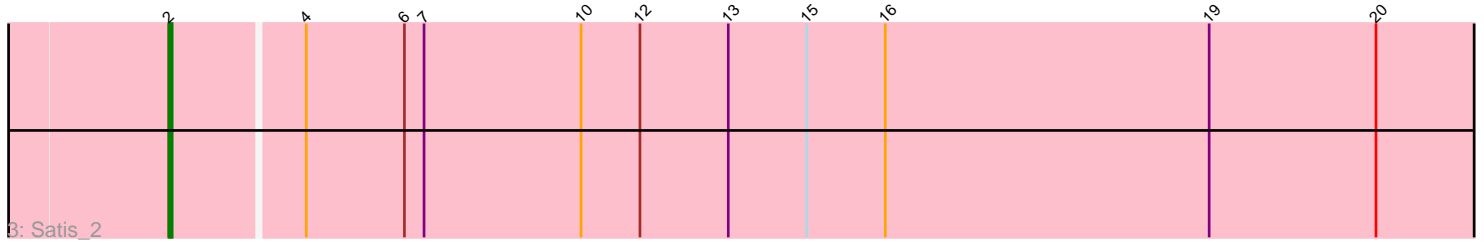
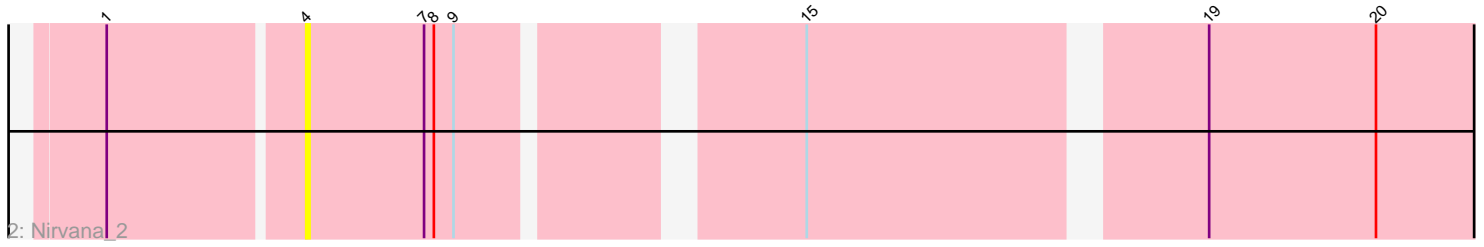
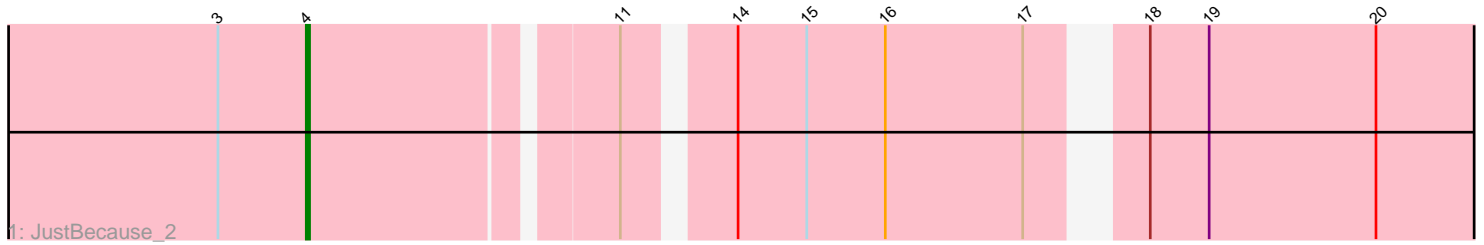


Pham 7414



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 7414 Report

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

Pham number 7414 has 7 members, 1 are drafts.

Phages represented in each track:

- Track 1 : JustBecause_2
- Track 2 : Nirvana_2
- Track 3 : Satis_2
- Track 4 : Frankenweenie_2
- Track 5 : EhyElimayoE_2, Kradal_2
- Track 6 : Kela_2

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

The start number called the most often in the published annotations is 2, it was called in 3 of the 6 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- EhyElimayoE_2, Kradal_2, Satis_2,

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

-

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

- Frankenweenie_2, JustBecause_2, Kela_2, Nirvana_2,

Summary by start number:

Start 2:

- Found in 3 of 7 (42.9%) of genes in pham
- Manual Annotations of this start: 3 of 6
- Called 100.0% of time when present
- Phage (with cluster) where this start called: EhyElimayoE_2 (BM), Kradal_2 (BM), Satis_2 (BM),

Start 4:

- Found in 7 of 7 (100.0%) of genes in pham
- Manual Annotations of this start: 3 of 6
- Called 57.1% of time when present

- Phage (with cluster) where this start called: Frankenweenie_2 (BM), JustBecause_2 (BM), Kela_2 (BM), Nirvana_2 (BM),

Summary by clusters:

There is one cluster represented in this pham: BM

Info for manual annotations of cluster BM:

- Start number 2 was manually annotated 3 times for cluster BM.
- Start number 4 was manually annotated 3 times for cluster BM.

Gene Information:

Gene: EhyElimayoE_2 Start: 1829, Stop: 1431, Start Num: 2

Candidate Starts for EhyElimayoE_2:

(Start: 2 @1829 has 3 MA's), (Start: 4 @1790 has 3 MA's), (6, 1760), (10, 1706), (12, 1688), (13, 1661), (15, 1637), (16, 1613), (19, 1514), (20, 1463),

Gene: Frankenweenie_2 Start: 1707, Stop: 1378, Start Num: 4

Candidate Starts for Frankenweenie_2:

(3, 1731), (Start: 4 @1707 has 3 MA's), (5, 1680), (15, 1572), (19, 1461), (20, 1410),

Gene: JustBecause_2 Start: 1676, Stop: 1350, Start Num: 4

Candidate Starts for JustBecause_2:

(3, 1703), (Start: 4 @1676 has 3 MA's), (11, 1589), (14, 1562), (15, 1541), (16, 1517), (17, 1475), (18, 1451), (19, 1433), (20, 1382),

Gene: Kela_2 Start: 1676, Stop: 1350, Start Num: 4

Candidate Starts for Kela_2:

(3, 1703), (Start: 4 @1676 has 3 MA's), (11, 1589), (14, 1562), (15, 1541), (16, 1517), (17, 1475), (18, 1451), (19, 1433), (20, 1382),

Gene: Kradal_2 Start: 1829, Stop: 1431, Start Num: 2

Candidate Starts for Kradal_2:

(Start: 2 @1829 has 3 MA's), (Start: 4 @1790 has 3 MA's), (6, 1760), (10, 1706), (12, 1688), (13, 1661), (15, 1637), (16, 1613), (19, 1514), (20, 1463),

Gene: Nirvana_2 Start: 1764, Stop: 1435, Start Num: 4

Candidate Starts for Nirvana_2:

(1, 1821), (Start: 4 @1764 has 3 MA's), (7, 1728), (8, 1725), (9, 1719), (15, 1629), (19, 1518), (20, 1467),

Gene: Satis_2 Start: 1825, Stop: 1427, Start Num: 2

Candidate Starts for Satis_2:

(Start: 2 @1825 has 3 MA's), (Start: 4 @1786 has 3 MA's), (6, 1756), (7, 1750), (10, 1702), (12, 1684), (13, 1657), (15, 1633), (16, 1609), (19, 1510), (20, 1459),