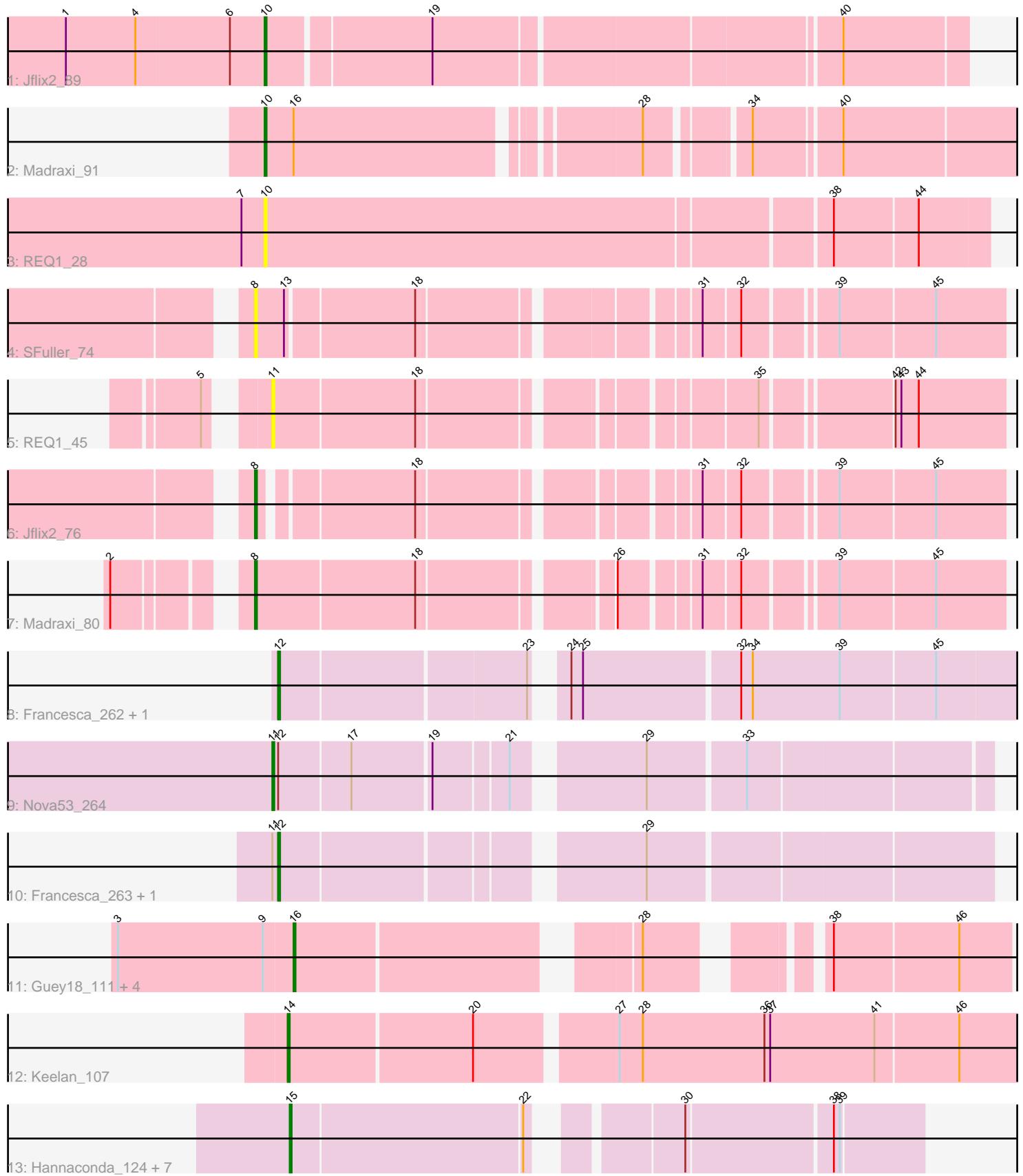


# Pham 280792



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 280792 Report

This analysis was run 02/07/26 on database version 634.

Pham number 280792 has 26 members, 6 are drafts.

Phages represented in each track:

- Track 1 : Jflix2\_89
- Track 2 : Madraxi\_91
- Track 3 : REQ1\_28
- Track 4 : SFuller\_74
- Track 5 : REQ1\_45
- Track 6 : Jflix2\_76
- Track 7 : Madraxi\_80
- Track 8 : Francesca\_262, Dorin\_262
- Track 9 : Nova53\_264
- Track 10 : Francesca\_263, Dorin\_263
- Track 11 : Guey18\_111, Volt\_110, Ziko\_109, Fryberger\_106, Ronaldo\_108
- Track 12 : Keelan\_107
- Track 13 : Hannaconda\_124, Yeet\_125, Odette\_135, Superphikiman\_128, HokkenD\_123, Courthouse\_126, Rearden\_129, Gonephishing\_125

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

The start number called the most often in the published annotations is 15, it was called in 5 of the 20 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Courthouse\_126, Gonephishing\_125, Hannaconda\_124, HokkenD\_123, Odette\_135, Rearden\_129, Superphikiman\_128, Yeet\_125,

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

- 

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

- Dorin\_262, Dorin\_263, Francesca\_262, Francesca\_263, Fryberger\_106, Guey18\_111, Jflix2\_76, Jflix2\_89, Keelan\_107, Madraxi\_80, Madraxi\_91, Nova53\_264, REQ1\_28, REQ1\_45, Ronaldo\_108, SFuller\_74, Volt\_110, Ziko\_109,

### **Summary by start number:**

Start 8:

- Found in 3 of 26 ( 11.5% ) of genes in pham
- Manual Annotations of this start: 2 of 20
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Jflix2\_76 (CF), Madraxi\_80 (CF), SFuller\_74 (CF),

Start 10:

- Found in 3 of 26 ( 11.5% ) of genes in pham
- Manual Annotations of this start: 2 of 20
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Jflix2\_89 (CF), Madraxi\_91 (CF), REQ1\_28 (CF),

Start 11:

- Found in 4 of 26 ( 15.4% ) of genes in pham
- Manual Annotations of this start: 1 of 20
- Called 50.0% of time when present
- Phage (with cluster) where this start called: Nova53\_264 (CG), REQ1\_45 (CF),

Start 12:

- Found in 5 of 26 ( 19.2% ) of genes in pham
- Manual Annotations of this start: 4 of 20
- Called 80.0% of time when present
- Phage (with cluster) where this start called: Dorin\_262 (CG), Dorin\_263 (CG), Francesca\_262 (CG), Francesca\_263 (CG),

Start 14:

- Found in 1 of 26 ( 3.8% ) of genes in pham
- Manual Annotations of this start: 1 of 20
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Keelan\_107 (DP),

Start 15:

- Found in 8 of 26 ( 30.8% ) of genes in pham
- Manual Annotations of this start: 5 of 20
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Courthouse\_126 (J), Gonephishing\_125 (J), Hannaconda\_124 (J), HokkenD\_123 (J), Odette\_135 (J), Rearden\_129 (J), Superphikiman\_128 (J), Yeet\_125 (J),

Start 16:

- Found in 6 of 26 ( 23.1% ) of genes in pham
- Manual Annotations of this start: 5 of 20
- Called 83.3% of time when present
- Phage (with cluster) where this start called: Fryberger\_106 (DP), Guey18\_111 (DP), Ronaldo\_108 (DP), Volt\_110 (DP), Ziko\_109 (DP),

**Summary by clusters:**

There are 4 clusters represented in this pham: J, CG, CF, DP,

Info for manual annotations of cluster CF:

- Start number 8 was manually annotated 2 times for cluster CF.
- Start number 10 was manually annotated 2 times for cluster CF.

Info for manual annotations of cluster CG:

- Start number 11 was manually annotated 1 time for cluster CG.
- Start number 12 was manually annotated 4 times for cluster CG.

Info for manual annotations of cluster DP:

- Start number 14 was manually annotated 1 time for cluster DP.
- Start number 16 was manually annotated 5 times for cluster DP.

Info for manual annotations of cluster J:

- Start number 15 was manually annotated 5 times for cluster J.

### ***Gene Information:***

Gene: Courthouse\_126 Start: 68736, Stop: 69029, Start Num: 15

Candidate Starts for Courthouse\_126:

(Start: 15 @68736 has 5 MA's), (22, 68853), (30, 68913), (38, 68985), (39, 68988),

Gene: Dorin\_262 Start: 128550, Stop: 128915, Start Num: 12

Candidate Starts for Dorin\_262:

(Start: 12 @128550 has 4 MA's), (23, 128673), (24, 128682), (25, 128688), (32, 128766), (34, 128772), (39, 128817), (45, 128865),

Gene: Dorin\_263 Start: 128893, Stop: 129231, Start Num: 12

Candidate Starts for Dorin\_263:

(Start: 11 @128890 has 1 MA's), (Start: 12 @128893 has 4 MA's), (29, 129061),

Gene: Francesca\_262 Start: 129192, Stop: 129557, Start Num: 12

Candidate Starts for Francesca\_262:

(Start: 12 @129192 has 4 MA's), (23, 129315), (24, 129324), (25, 129330), (32, 129408), (34, 129414), (39, 129459), (45, 129507),

Gene: Francesca\_263 Start: 129535, Stop: 129873, Start Num: 12

Candidate Starts for Francesca\_263:

(Start: 11 @129532 has 1 MA's), (Start: 12 @129535 has 4 MA's), (29, 129703),

Gene: Fryberger\_106 Start: 52384, Stop: 52698, Start Num: 16

Candidate Starts for Fryberger\_106:

(3, 52294), (9, 52369), (Start: 16 @52384 has 5 MA's), (28, 52540), (38, 52609), (46, 52672),

Gene: Gonephishing\_125 Start: 68379, Stop: 68672, Start Num: 15

Candidate Starts for Gonephishing\_125:

(Start: 15 @68379 has 5 MA's), (22, 68496), (30, 68556), (38, 68628), (39, 68631),

Gene: Guey18\_111 Start: 53707, Stop: 54021, Start Num: 16

Candidate Starts for Guey18\_111:

(3, 53617), (9, 53692), (Start: 16 @53707 has 5 MA's), (28, 53863), (38, 53932), (46, 53995),

Gene: Hannaconda\_124 Start: 69032, Stop: 69325, Start Num: 15

Candidate Starts for Hannaconda\_124:

(Start: 15 @69032 has 5 MA's), (22, 69149), (30, 69209), (38, 69281), (39, 69284),

Gene: HokkenD\_123 Start: 71913, Stop: 72206, Start Num: 15

Candidate Starts for HokkenD\_123:

(Start: 15 @71913 has 5 MA's), (22, 72030), (30, 72090), (38, 72162), (39, 72165),

Gene: Jflix2\_89 Start: 50660, Stop: 51001, Start Num: 10

Candidate Starts for Jflix2\_89:

(1, 50558), (4, 50594), (6, 50642), (Start: 10 @50660 has 2 MA's), (19, 50741), (40, 50939),

Gene: Jflix2\_76 Start: 46571, Stop: 46909, Start Num: 8

Candidate Starts for Jflix2\_76:

(Start: 8 @46571 has 2 MA's), (18, 46643), (31, 46766), (32, 46784), (39, 46826), (45, 46874),

Gene: Keelan\_107 Start: 53290, Stop: 53652, Start Num: 14

Candidate Starts for Keelan\_107:

(Start: 14 @53290 has 1 MA's), (20, 53383), (27, 53449), (28, 53461), (36, 53524), (37, 53527), (41, 53581), (46, 53623),

Gene: Madraxi\_91 Start: 52811, Stop: 53170, Start Num: 10

Candidate Starts for Madraxi\_91:

(Start: 10 @52811 has 2 MA's), (Start: 16 @52826 has 5 MA's), (28, 52988), (34, 53033), (40, 53075),

Gene: Madraxi\_80 Start: 49350, Stop: 49697, Start Num: 8

Candidate Starts for Madraxi\_80:

(2, 49296), (Start: 8 @49350 has 2 MA's), (18, 49431), (26, 49518), (31, 49554), (32, 49572), (39, 49614), (45, 49662),

Gene: Nova53\_264 Start: 130197, Stop: 130535, Start Num: 11

Candidate Starts for Nova53\_264:

(Start: 11 @130197 has 1 MA's), (Start: 12 @130200 has 4 MA's), (17, 130236), (19, 130275), (21, 130311), (29, 130368), (33, 130416),

Gene: Odette\_135 Start: 73830, Stop: 74123, Start Num: 15

Candidate Starts for Odette\_135:

(Start: 15 @73830 has 5 MA's), (22, 73947), (30, 74007), (38, 74079), (39, 74082),

Gene: REQ1\_28 Start: 11269, Stop: 11631, Start Num: 10

Candidate Starts for REQ1\_28:

(7, 11257), (Start: 10 @11269 has 2 MA's), (38, 11554), (44, 11596),

Gene: REQ1\_45 Start: 18020, Stop: 18358, Start Num: 11

Candidate Starts for REQ1\_45:

(5, 17999), (Start: 11 @18020 has 1 MA's), (18, 18092), (35, 18242), (42, 18302), (43, 18305), (44, 18314),

Gene: Rearden\_129 Start: 69937, Stop: 70230, Start Num: 15

Candidate Starts for Rearden\_129:

(Start: 15 @69937 has 5 MA's), (22, 70054), (30, 70114), (38, 70186), (39, 70189),

Gene: Ronaldo\_108 Start: 53289, Stop: 53603, Start Num: 16

Candidate Starts for Ronaldo\_108:

(3, 53199), (9, 53274), (Start: 16 @53289 has 5 MA's), (28, 53445), (38, 53514), (46, 53577),

Gene: SFuller\_74 Start: 46761, Stop: 47108, Start Num: 8

Candidate Starts for SFuller\_74:

(Start: 8 @46761 has 2 MA's), (13, 46776), (18, 46839), (31, 46965), (32, 46983), (39, 47025), (45, 47073),

Gene: Superphikiman\_128 Start: 69018, Stop: 69311, Start Num: 15

Candidate Starts for Superphikiman\_128:

(Start: 15 @69018 has 5 MA's), (22, 69135), (30, 69195), (38, 69267), (39, 69270),

Gene: Volt\_110 Start: 53453, Stop: 53767, Start Num: 16

Candidate Starts for Volt\_110:

(3, 53363), (9, 53438), (Start: 16 @53453 has 5 MA's), (28, 53609), (38, 53678), (46, 53741),

Gene: Yeet\_125 Start: 71942, Stop: 72235, Start Num: 15

Candidate Starts for Yeet\_125:

(Start: 15 @71942 has 5 MA's), (22, 72059), (30, 72119), (38, 72191), (39, 72194),

Gene: Ziko\_109 Start: 53295, Stop: 53609, Start Num: 16

Candidate Starts for Ziko\_109:

(3, 53205), (9, 53280), (Start: 16 @53295 has 5 MA's), (28, 53451), (38, 53520), (46, 53583),