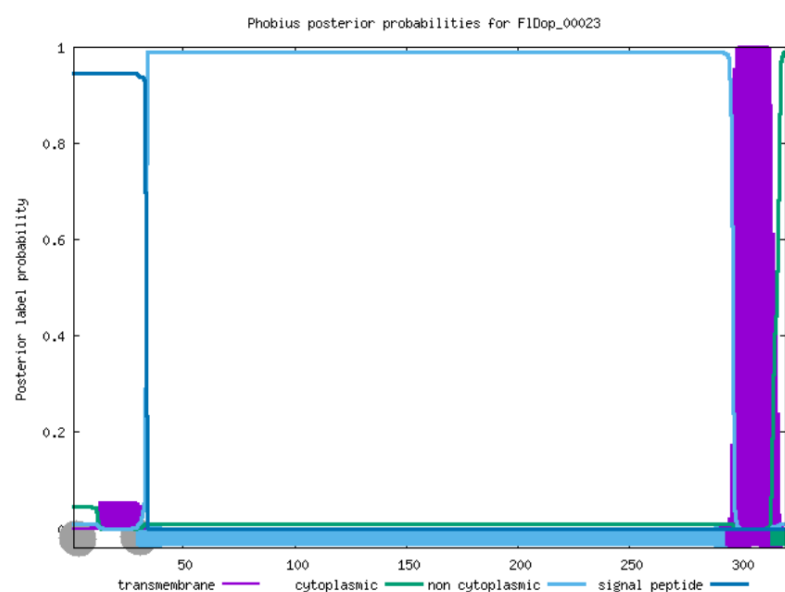


Supplementary Figure S1. Graphs from Phobius prediction of putative effectors

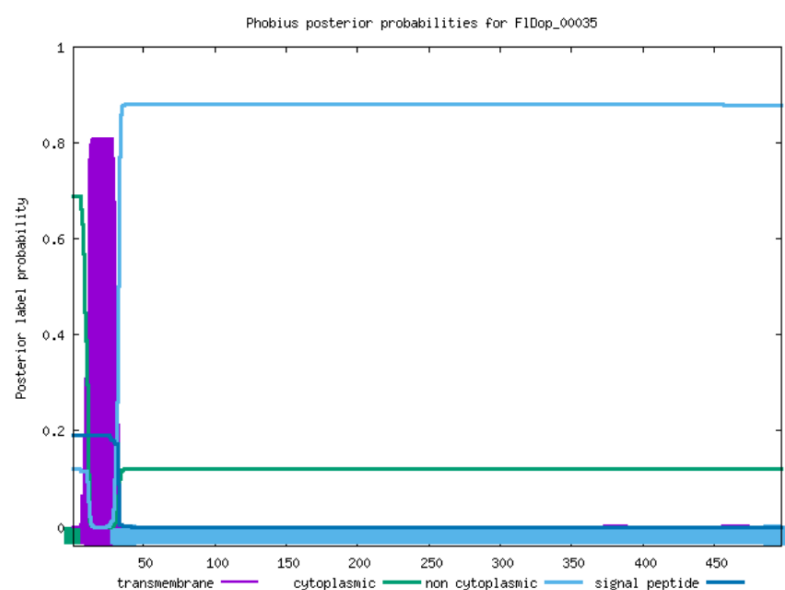
Prediction of FIDop_00023

```
ID FIDop_00023
FT SIGNAL      1    32
FT REGION      1    12      N-REGION.
FT REGION     13    24      H-REGION.
FT REGION     25    32      C-REGION.
FT TOPO_DOM    33   296      NON CYTOPLASMIC.
FT TRANSMEM   297   316
FT TOPO_DOM   317   320      CYTOPLASMIC.
//
```



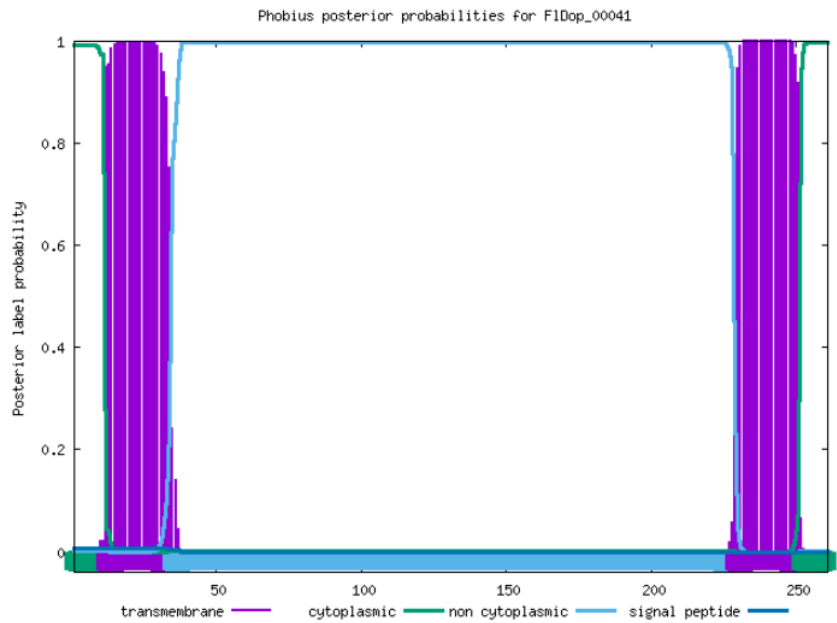
Prediction of FIDop_00035

```
ID FIDop_00035
FT TOPO_DOM    1    11      CYTOPLASMIC.
FT TRANSMEM   12    32
FT TOPO_DOM   33   497      NON CYTOPLASMIC.
//
```



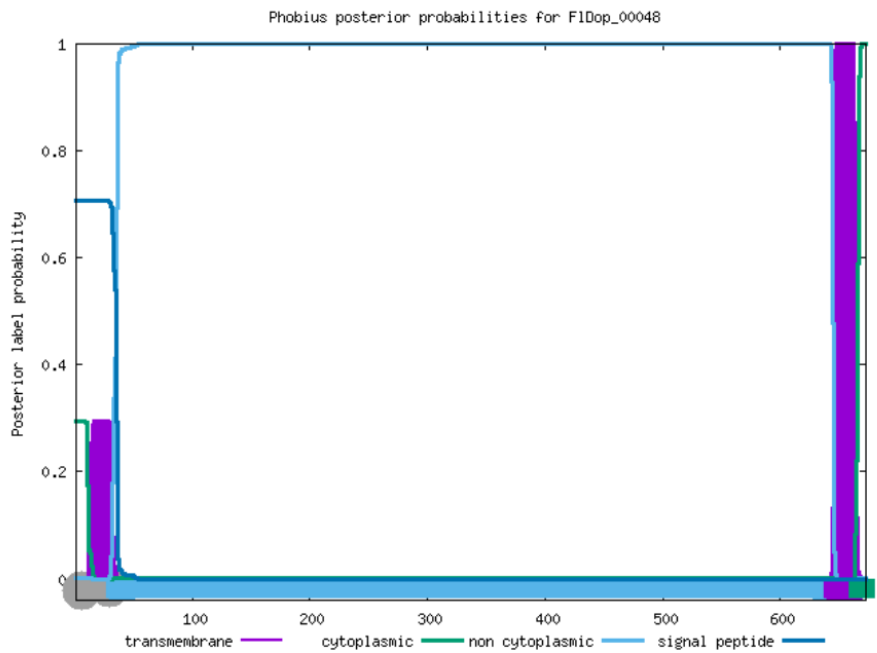
Prediction of FIDop_00041

ID	FIDop_00041			
FT	TOPO_DOM	1	11	CYTOPLASMIC.
FT	TRANSMEM	12	34	
FT	TOPO_DOM	35	228	NON CYTOPLASMIC.
FT	TRANSMEM	229	251	
FT	TOPO_DOM	252	261	CYTOPLASMIC.
//				



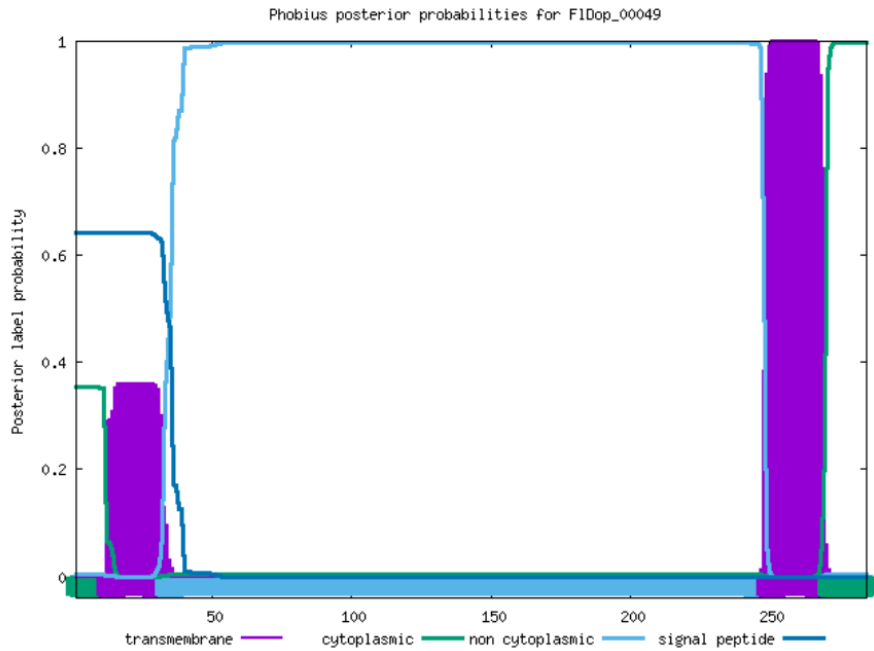
Prediction of FIDop_00048

ID	FIDop_00048				
FT	SIGNAL	1	34		
FT	REGION	1	14		N-REGION.
FT	REGION	15	30		H-REGION.
FT	REGION	31	34		C-REGION.
FT	TOPO_DOM	35	645		NON CYTOPLASMIC.
FT	TRANSMEM	646	666		
FT	TOPO_DOM	667	673		CYTOPLASMIC.
//					



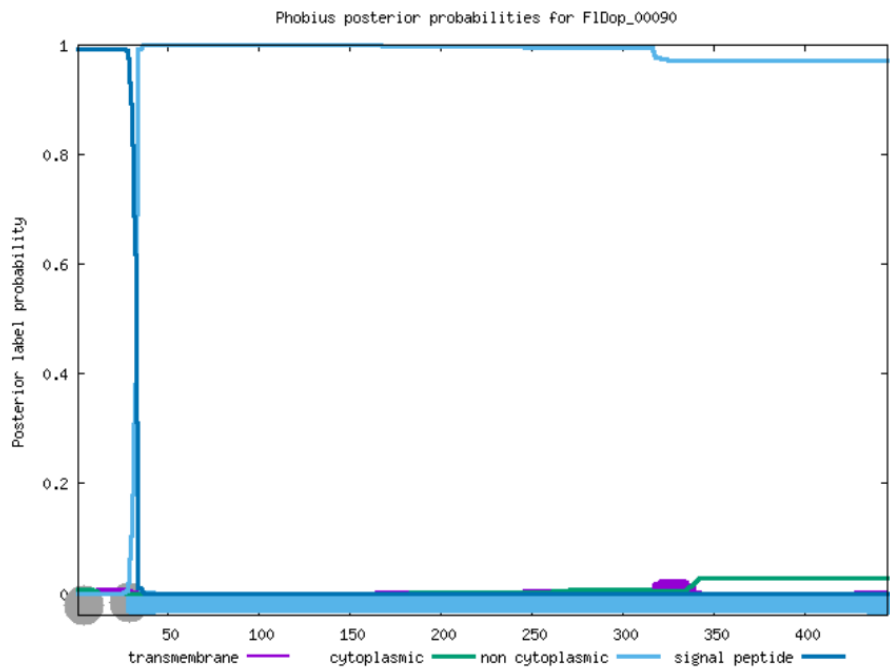
Prediction of FIDop_00049

```
ID FIDop_00049
FT TOPO_DOM 1 11 CYTOPLASMIC.
FT TRANSMEM 12 32
FT TOPO_DOM 33 248 NON CYTOPLASMIC.
FT TRANSMEM 249 270
FT TOPO_DOM 271 285 CYTOPLASMIC.
//
```



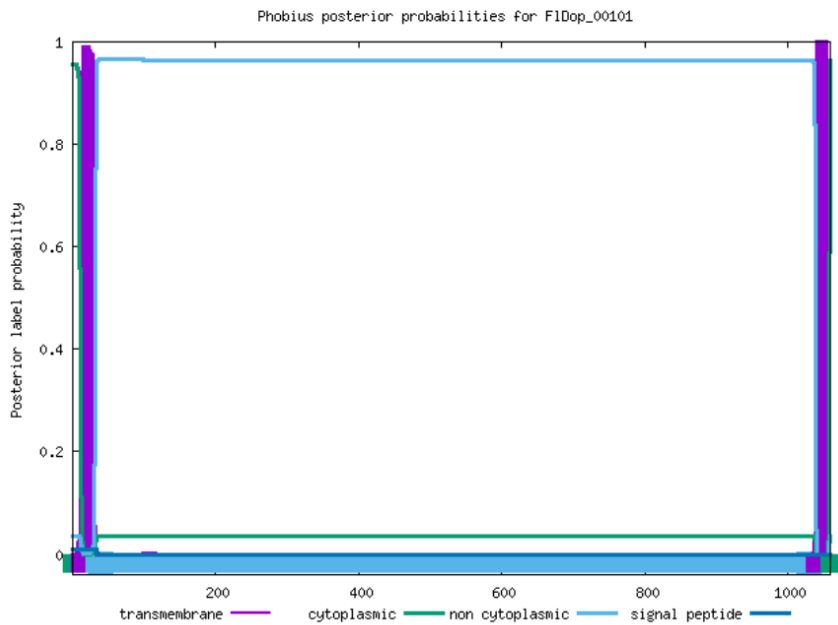
Prediction of FIDop_00090

```
ID FIDop_00090
FT SIGNAL 1 32
FT REGION 1 12 N-REGION.
FT REGION 13 24 H-REGION.
FT REGION 25 32 C-REGION.
FT TOPO_DOM 33 445 NON CYTOPLASMIC.
//
```



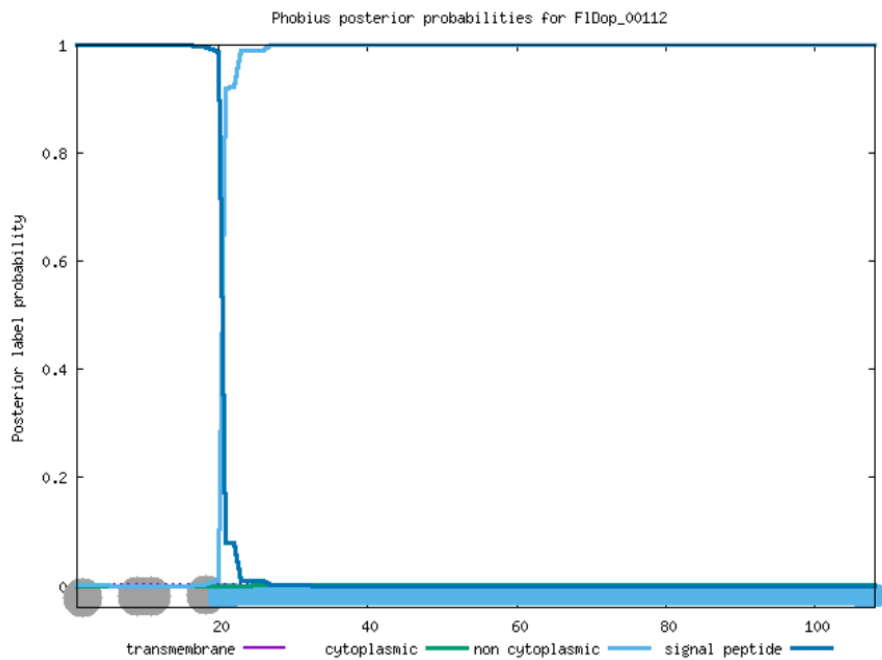
Prediction of FIDop_00101

```
ID FIDop_00101
FT TOPO_DOM 1 11 CYTOPLASMIC.
FT TRANSMEM 12 32
FT TOPO_DOM 33 1038 NON CYTOPLASMIC.
FT TRANSMEM 1039 1058
FT TOPO_DOM 1059 1059 CYTOPLASMIC.
//
```



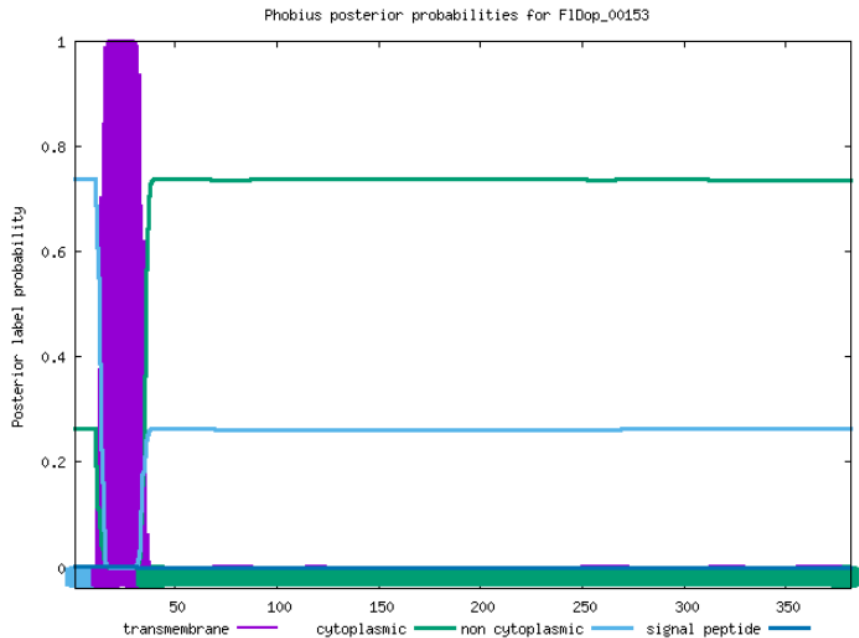
Prediction of FIDop_00112

```
ID FIDop_00112
FT SIGNAL 1 19
FT REGION 1 2 N-REGION.
FT REGION 3 14 H-REGION.
FT REGION 15 19 C-REGION.
FT TOPO_DOM 20 108 NON CYTOPLASMIC.
//
```



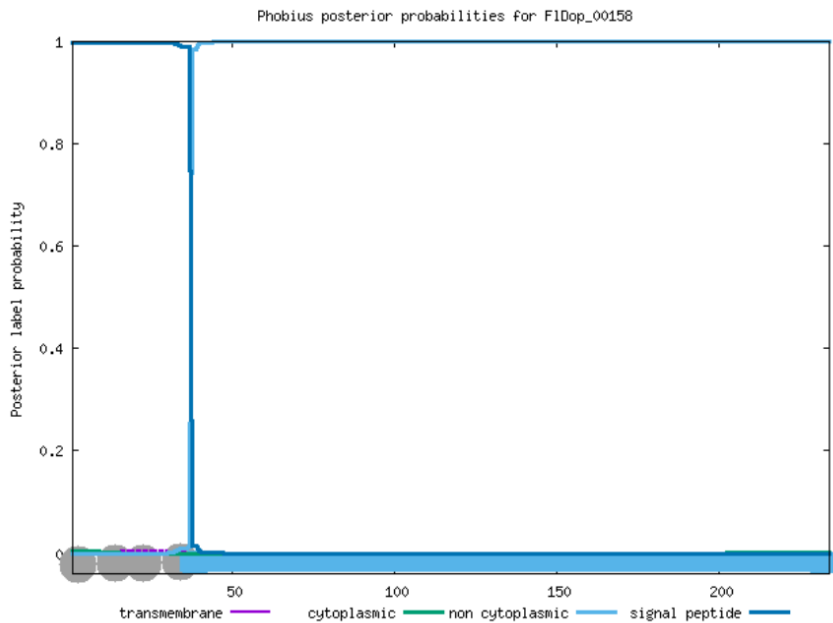
Prediction of FIDop_00153

```
ID FIDop_00153
FT TOPO_DOM 1 13 NON CYTOPLASMIC.
FT TRANSMEM 14 35
FT TOPO_DOM 36 382 CYTOPLASMIC.
//
```



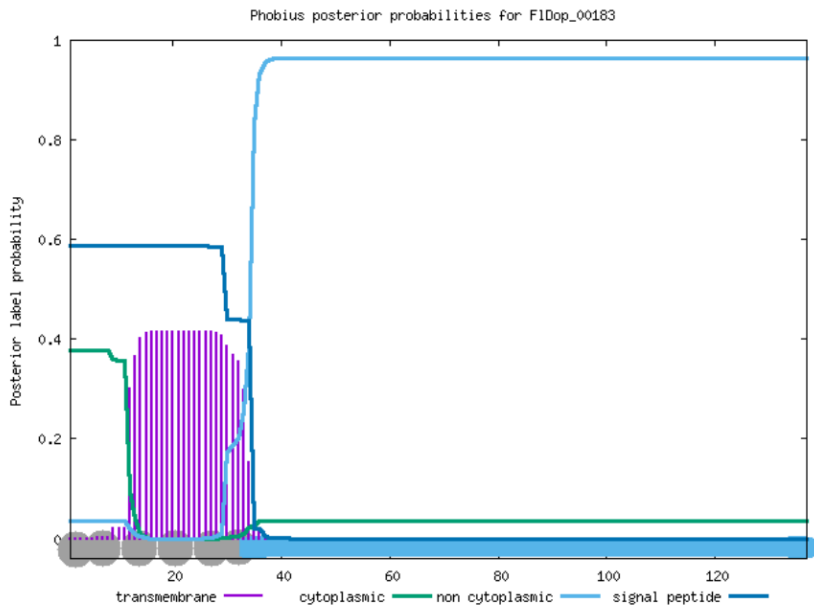
Prediction of FIDop_00158

```
ID FIDop_00158
FT SIGNAL 1 36
FT REGION 1 15 N-REGION.
FT REGION 16 28 H-REGION.
FT REGION 29 36 C-REGION.
FT TOPO_DOM 37 234 NON CYTOPLASMIC.
//
```



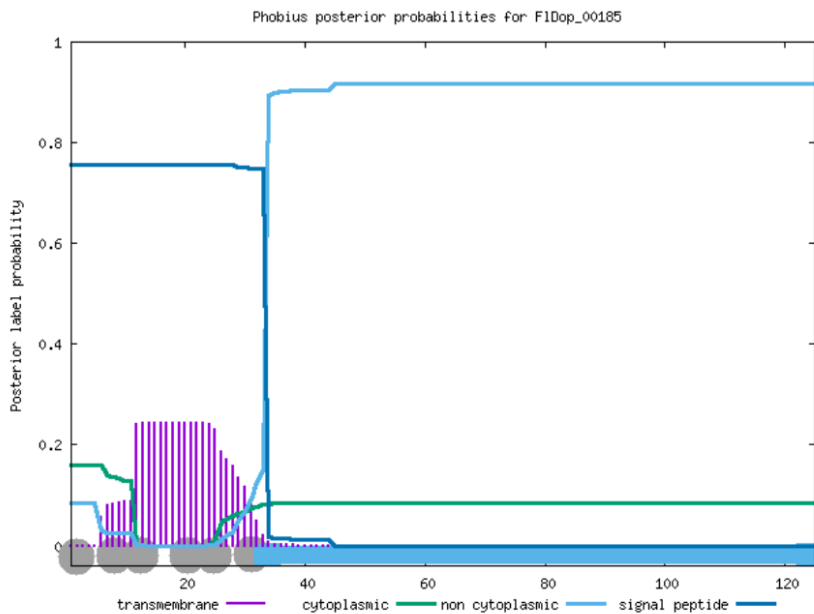
Prediction of FIDop_00183

ID	FIDop_00183			
FT	SIGNAL	1	33	
FT	REGION	1	12	N-REGION.
FT	REGION	13	24	H-REGION.
FT	REGION	25	33	C-REGION.
FT	TOPO_DOM	34	137	NON CYTOPLASMIC.
//				



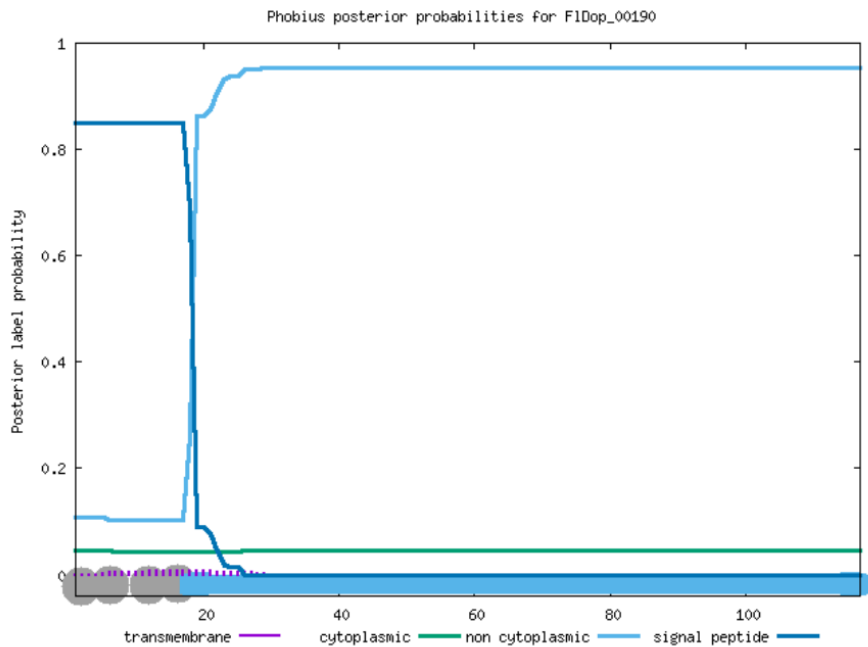
Prediction of FIDop_00185

ID	FIDop_00185			
FT	SIGNAL	1	32	
FT	REGION	1	9	N-REGION.
FT	REGION	10	25	H-REGION.
FT	REGION	26	32	C-REGION.
FT	TOPO_DOM	33	125	NON CYTOPLASMIC.
//				



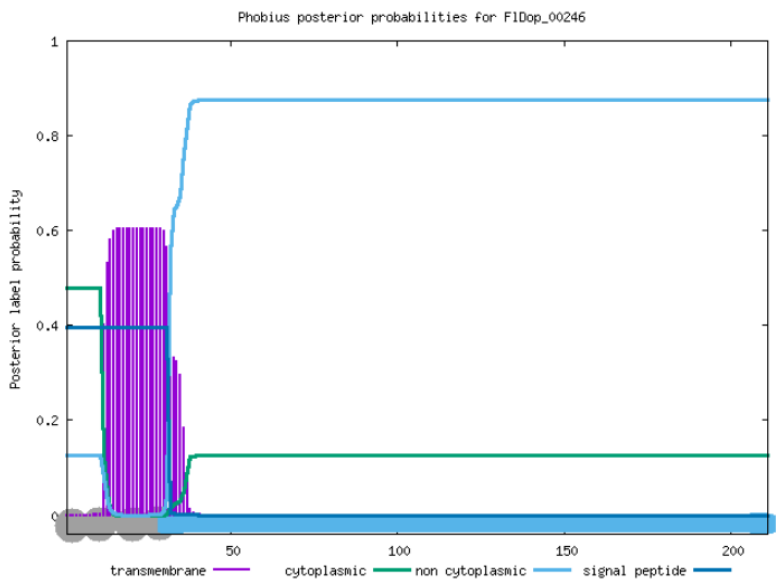
Prediction of FIDop_00190

```
ID FIDop_00190
FT SIGNAL      1    17
FT REGION      1     4      N-REGION.
FT REGION      5    12      H-REGION.
FT REGION     13    17      C-REGION.
FT TOPO_DOM    18   117      NON CYTOPLASMIC.
//
```



Prediction of FIDop_00246

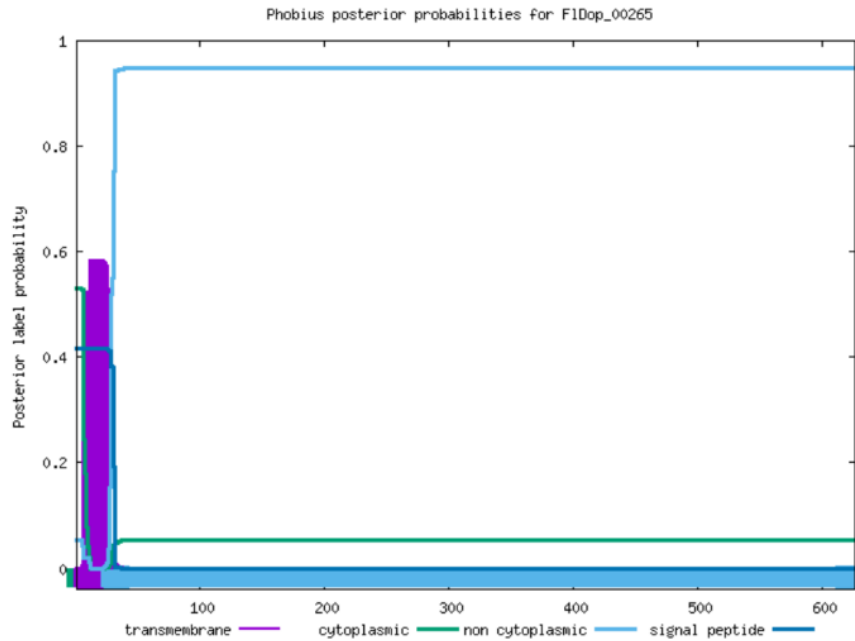
```
ID FIDop_00246
FT SIGNAL      1    30
FT REGION      1    13      N-REGION.
FT REGION     14    25      H-REGION.
FT REGION     26    30      C-REGION.
FT TOPO_DOM    31   211      NON CYTOPLASMIC.
//
```



Prediction of FIDop_00265

```

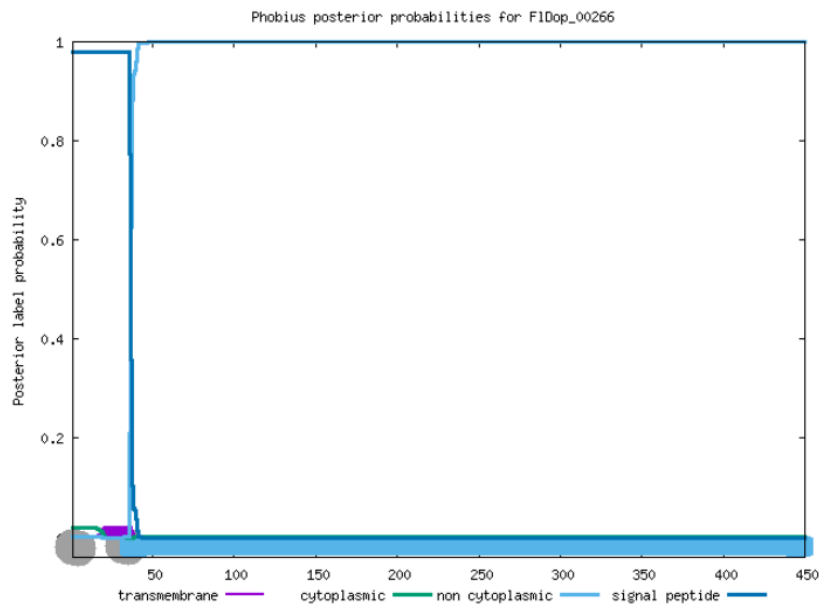
ID   FIDop_00265
FT   TOPO_DOM    1     6     CYTOPLASMIC.
FT   TRANSMEM    7    27
FT   TOPO_DOM    28   626    NON CYTOPLASMIC.
//
  
```



Prediction of FIDop_00266

```

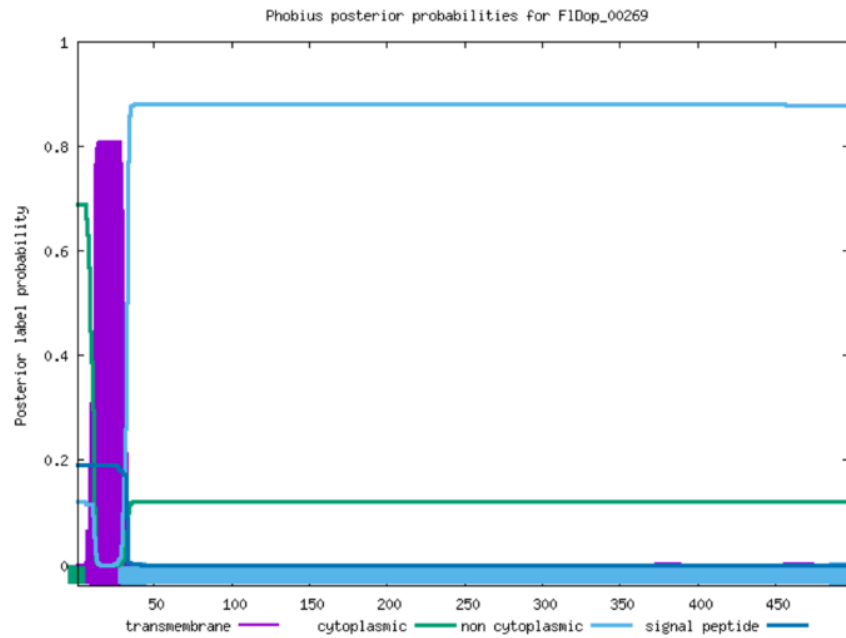
ID   FIDop_00266
FT   SIGNAL      1     35
FT   REGION      1     18    N-REGION.
FT   REGION     19     30    H-REGION.
FT   REGION     31     35    C-REGION.
FT   TOPO_DOM    36   450    NON CYTOPLASMIC.
//
  
```



Prediction of FIDop_00269

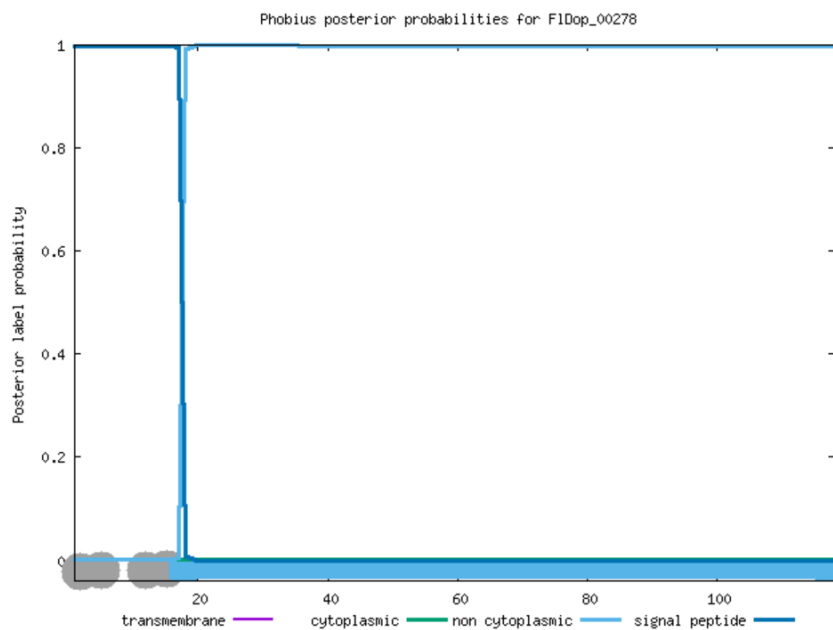
```

ID   FIDop_00269
FT   TOPO_DOM    1     11    CYTOPLASMIC.
FT   TRANSMEM    12    32
FT   TOPO_DOM    33    497    NON CYTOPLASMIC.
//
  
```



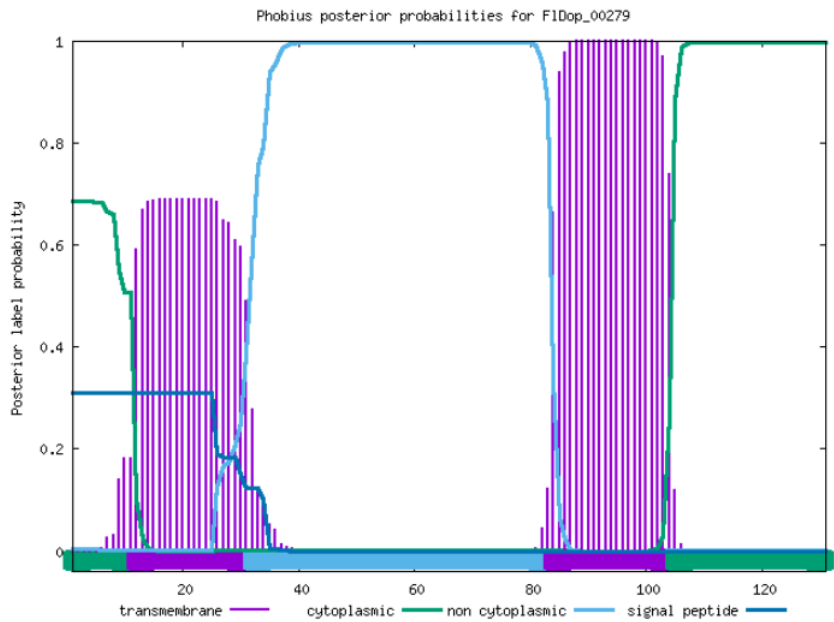
```

ID   FIDop_00278
FT   SIGNAL      1     16
FT   REGION      1     1    N-REGION.
FT   REGION      2     10    H-REGION.
FT   REGION      11    16    C-REGION.
FT   TOPO_DOM    17    118    NON CYTOPLASMIC.
//
  
```



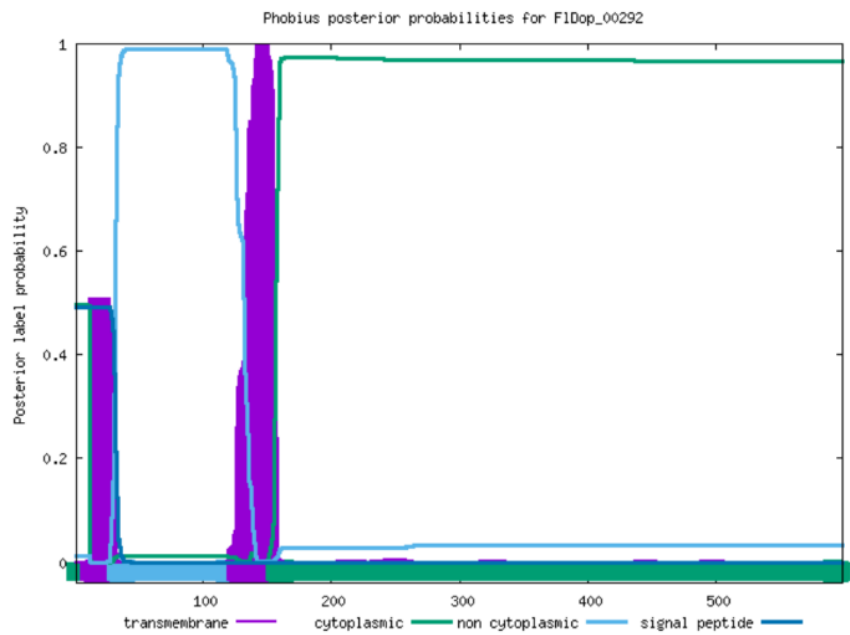
Prediction of FIDop_00279

```
ID FIDop_00279
FT TOPO_DOM 1 11 CYTOPLASMIC.
FT TRANSMEM 12 31
FT TOPO_DOM 32 83 NON CYTOPLASMIC.
FT TRANSMEM 84 104
FT TOPO_DOM 105 131 CYTOPLASMIC.
//
```



```
ID FIDop_00292
FT TOPO_DOM 1 11 CYTOPLASMIC.
FT TRANSMEM 12 32
FT TOPO_DOM 33 125 NON CYTOPLASMIC.
FT TRANSMEM 126 156
FT TOPO_DOM 157 599 CYTOPLASMIC.
//
```

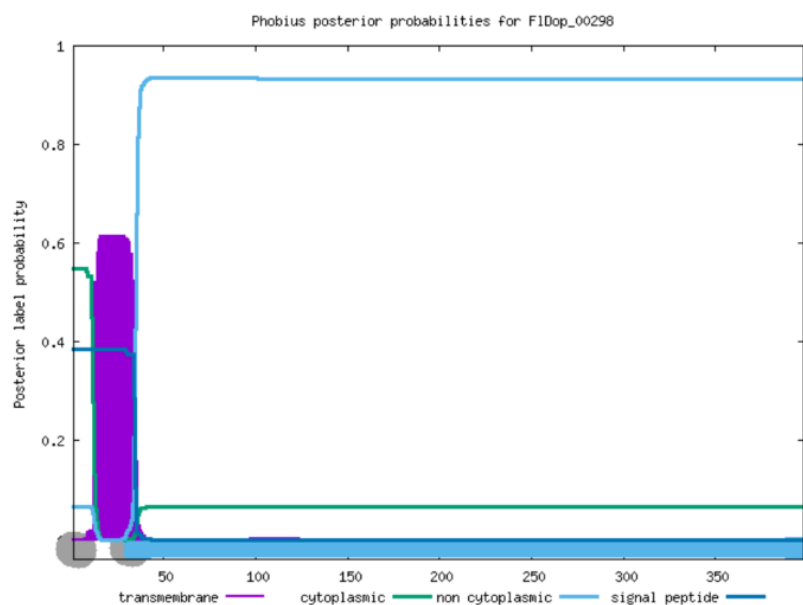
ftsH protein



```

ID FIDop_00298
FT SIGNAL 1 33
FT REGION 1 12 N-REGION.
FT REGION 13 24 H-REGION.
FT REGION 25 33 C-REGION.
FT TOPO_DOM 34 398 NON CYTOPLASMIC.
//

```

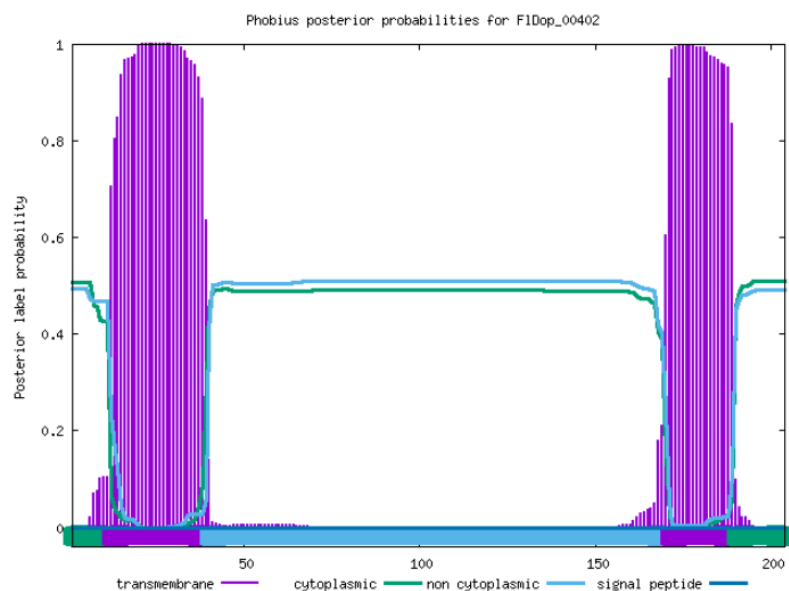


Prediction of FIDop_00402

```

ID FIDop_00402
FT TOPO_DOM 1 11 CYTOPLASMIC.
FT TRANSMEM 12 39
FT TOPO_DOM 40 170 NON CYTOPLASMIC.
FT TRANSMEM 171 189
FT TOPO_DOM 190 204 CYTOPLASMIC.
//

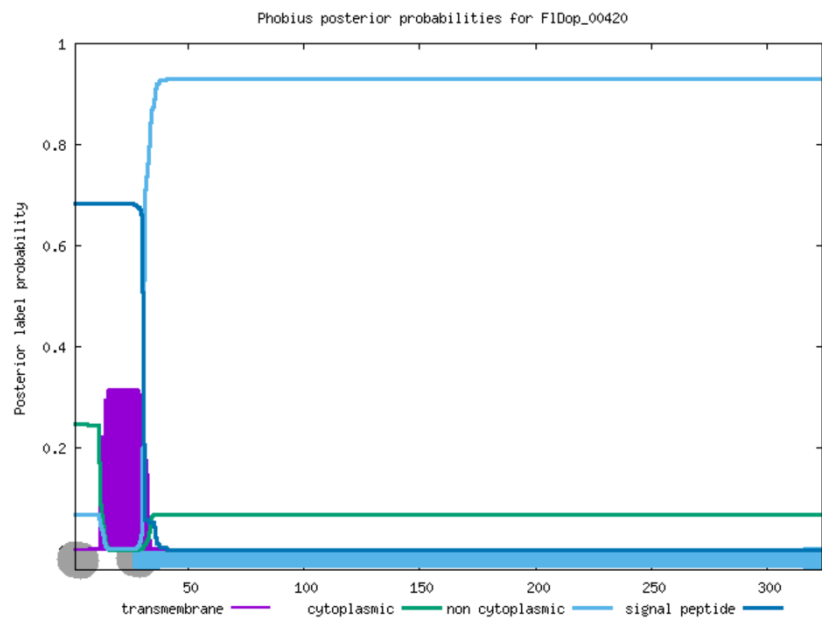
```



```

ID   FIDop_00420
FT   SIGNAL       1   29
FT   REGION       1   13   N-REGION.
FT   REGION      14   25   H-REGION.
FT   REGION      26   29   C-REGION.
FT   TOPO_DOM     30   324  NON CYTOPLASMIC.
//

```



Prediction of FIDop_00531

```

ID   FIDop_00531
FT   SIGNAL       1   32
FT   REGION       1   12   N-REGION.
FT   REGION      13   24   H-REGION.
FT   REGION      25   32   C-REGION.
FT   TOPO_DOM     33   122  NON CYTOPLASMIC.
//

```

