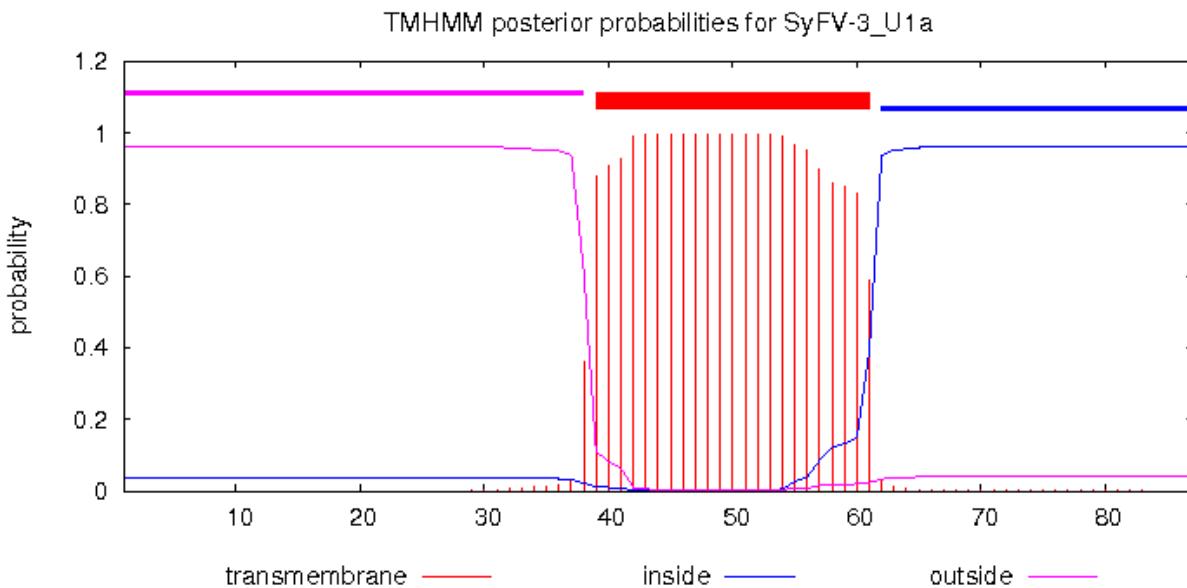


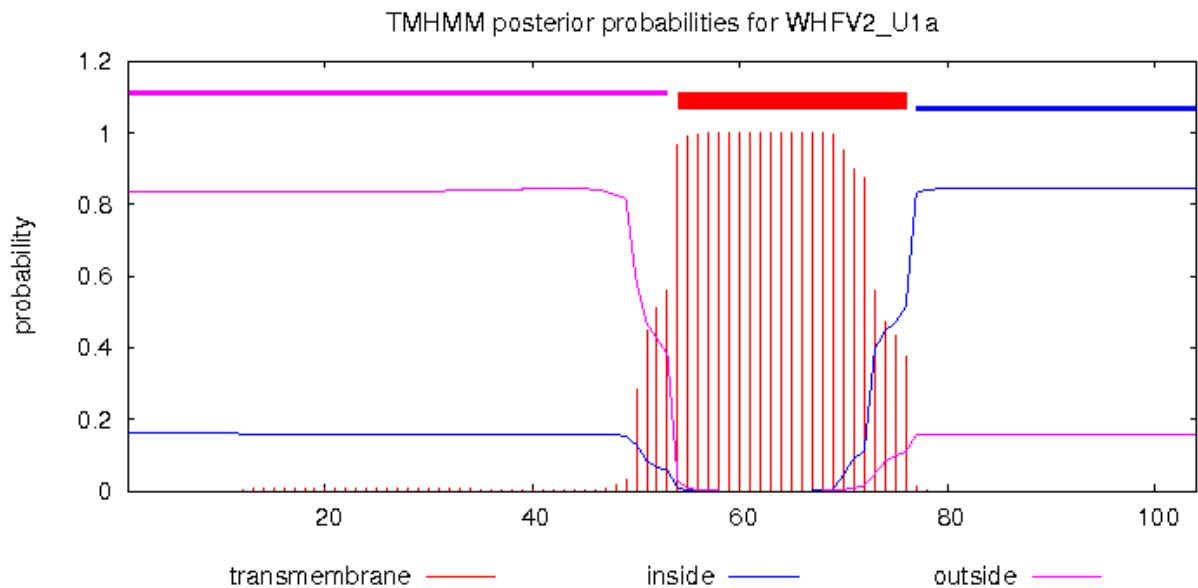
CLUSTAL 2.1 multiple sequence alignment

```
SyFV-3_U1a      MEIPLD----DFIQ--YSDQSFFDPLYNPLL-----QNHSSYS-HSSKAVTHIIILIIIFLL
WHFV2_U1a      MNSLYDQEAEIDLFSNLVSNGTHSVPLDTSFVNISVSGGSSTELPSSHNLNFNIFLVFLI
*: * *::: *: . ** ..::.. . ** . **: . *:::**:
SyFV-3_U1a      IV-FHLIQISNTLILQSSCTTIMD-TLS-LLTNQTLR-ILHNLI
WHFV2_U1a      IVTFLAIWTIQILAKSKKVRVDGSRSEQGGVRSRNQDVNVH
** *: * : . : . * .. :* : * . * * : 

# SyFV-3_U1a Length: 87
# SyFV-3_U1a Number of predicted TMHs: 1
# SyFV-3_U1a Exp number of AAs in TMHs: 22.10657
# SyFV-3_U1a Exp number, first 60 AAs: 21.4583
# SyFV-3_U1a Total prob of N-in: 0.03607
# SyFV-3_U1a POSSIBLE N-term signal sequence
SyFV-3_U1a      TMHMM2.0      outside      1      38
SyFV-3_U1a      TMHMM2.0      TMhelix     39      61
SyFV-3_U1a      TMHMM2.0      inside       62      87
```



```
# WHFV2_U1a Length: 104
# WHFV2_U1a Number of predicted TMHs: 1
# WHFV2_U1a Exp number of AAs in TMHs: 22.53844
# WHFV2_U1a Exp number, first 60 AAs: 8.96397
# WHFV2_U1a Total prob of N-in: 0.16222
WHFV2_U1a      TMHMM2.0      outside      1      53
WHFV2_U1a      TMHMM2.0      TMhelix     54      76
WHFV2_U1a      TMHMM2.0      inside       77      104
```



CLUSTAL 2.1 multiple sequence alignment

```

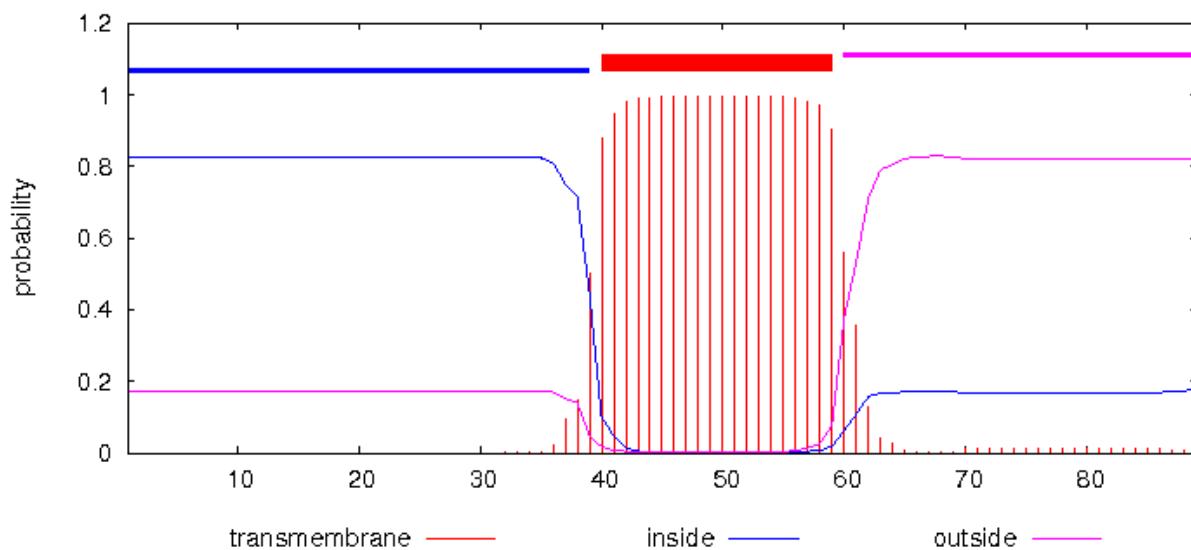
WHFV2_U1b      M-DYHSNLSQVEKGESG----WVEKLGTGWSKVPKPGCQSCMKWVIINLLIDIIILLT
SyFV-3_U1b      MHNYHGHSEPLNKPNQAQNTAQSYLKILHIIWIII----KFCFQLIIVFCVVGSFVIDCL
* :**.: . ::* ::    ::: *   * :    : *:: :*:: :: .:::

WHFV2_U1b      AYHVIGPGSASWNKDASSFVPAICGINSTIHMTPK
SyFV-3_U1b      VFKVVAAP--EDWR---DLIKNKS-LFTWQNMVT-
. ::*:. * .*.    .::    . : :  **

# WHFV2_U1b Length: 89
# WHFV2_U1b Number of predicted TMHs: 1
# WHFV2_U1b Exp number of AAs in TMHs: 21.72033
# WHFV2_U1b Exp number, first 60 AAs: 20.90298
# WHFV2_U1b Total prob of N-in: 0.82740
# WHFV2_U1b POSSIBLE N-term signal sequence
WHFV2_U1b      TMHMM2.0      inside     1      39
WHFV2_U1b      TMHMM2.0      TMhelix   40      59
WHFV2_U1b      TMHMM2.0      outside    60      89

```

TMHMM posterior probabilities for WHFV2\_U1b



```
# SyFV-3_U1b Length: 82
# SyFV-3_U1b Number of predicted TMHs: 1
# SyFV-3_U1b Exp number of AAs in TMHs: 23.2496
# SyFV-3_U1b Exp number, first 60 AAs: 23.22388
# SyFV-3_U1b Total prob of N-in: 0.15539
# SyFV-3_U1b POSSIBLE N-term signal sequence
SyFV-3_U1b      TMHMM2.0      outside      1      28
SyFV-3_U1b      TMHMM2.0      TMhelix     29      51
SyFV-3_U1b      TMHMM2.0      inside       52      82
```

TMHMM posterior probabilities for SyFV-3\_U1b

