

logo

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NTDB id 93443 AMD29 RS01255 WP 000923270.1
NTDB id 434 SGO RS09485 WP 012130954.1
NTDB id 437 SPR RS10075 WP 000950018.1
consensus
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MLNLSYDFMQRAFLAVIAMSLFSPILGTFLILRRQSLMSDTLSHVS LAGVAFGLVLGLSPTSTLWVLA AVLEYLRTVYK
 MLNLSYDFMQRAFLAVIAMSLFSPILGTFLILRRQSLMSDTLSHVS LAGVAFGLVLGLSPTSTLWVLA AVLEYLRTVYK
 .MLNLSYDFMQRAFLAVIAMSLFSPILGTFLILRRQSLMSDTLSHVS LAGVAFGLVLGLSPTSTLWVLA AVLEYLRTVYK
 .MLSLLSYDFMQRAFLAVIAMSLFSPILGTFLILRRQSLMSDTLSHVS LAGVAFGLVLGLSPTSTLWVLA AVLEYLRTVYK
 * * * * *

85
84
84

logo

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consensus
```

[illegible]

170
169
169

logo

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consensus
```

[illegible]

255
254
254

logo

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consensus

Sequence logo for the 12 amino acid motif. The logo shows the conservation of residues across three rows. The top row has T, F, L, L, V, S, L, V, R, R, F, Y, K. The middle row has G, T, F, L, L, V, S, L, V, G, L, L, R, K, R. The bottom row has S, L, F, L, V, V, N, L, I, K, K, F, V, K. Below the sequences are asterisks indicating conservation: ** ! * * * * * ! * * * * * !

270
268
268

-  non conserved
-  similar
-  >50% conserved