

```

agcttccggcacggccttcaagcgcgggacgcgacaaagtc59atggaccgcaaccctcg 59 start
                                     M D R N P S* 6 exp phos|pred yoy
ccgccgccgcccgggtcgcgacaaggaggaggaggaggagggtggccgggtggagactgcata 119
P P P P G R D K E E E E E V A G G D C I 26 neg charge cluster
gggagcacgggtctacagcaaacactggctcttcggcgtcctcagcggactcatccagatt 179 ex1|ex2 bound
G S T V Y S K H W L F G V L S G L I Q I 46
gttagcctgaaaacaccaaactagctcagatgatgaggagcagctgacggagcttgat 239
V S P E N T K S S S D D E E Q L T E L D 66 neg charge cluster
gaagaaatggagaatgaaatttgcagagtatgggatgtcaatggatgaggacgtggct 299 ex2|ex3 bound
E E M E N E I C R V W D M S M D E D V A 86 ARM-like pred NES
Ttatttctccaagaatttaatgctcctgatataattcatgggagtactggccaagtccaag 359
L F L Q E F N A P D I F M G V L A K S K 106 Sympotin-1_like
tgtcctcgattaagagaaatctgtgtgggaatttttaggtaatatggcctgtttccaggag 419 ex3|ex4 bound
C P R L R E I C V G I L G N M A C F Q E 126 pred MAPK
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I C V S I S S D K N L G Q V L L H C L Y 146
gattcagaccacactactctgctggaacaagcaggttgttgccttacttgcctttcccag 539 ex5|ex6 bound
D S D P P T* L L E T S R L L L T C L S* Q 166 *pred phos
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A E V A S V W V E R I Q E H P A I Y D S 186
atttgcttcattatgtcaagttcaacaaatgttgacttgcctggtgaagggtgggggaggtt 659 ex6|ex7 bound
I C F I M S S S T N V D L L V K V G E V 206
gtggacaagctcttttgatttggatgagaaactaatgttagaatgggtcagaaatggggct 719
V D K* L F D L D E K L M L E W V R N G A 226 *pred ubiq
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A Q P L D Q P Q E E S E E Q P V F R L V 246
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P C I L E A A K Q V R S E N P E W L D V 266
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Y M H I L Q L L T T V D D G I Q A I V H 286
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C P D T G K D I W N L L F D L V* C H E F 306 *pred ubiq
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C Q S D D P P I I L Q E Q K T V L A S V 326
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T Q D D F H L K I L K D I L C E F L S N 406 exp phos
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