Phyre²

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Description	P13518
Date	Thu Jan 5 11:33:38 GMT 2012
Unique Job ID	0fba59284080d6c1

Secondary structure and disorder prediction


430
440
450
460
470
480
Sequence
TREGQVHHRELMCRIFDGNEEVSSAEYMPMVLQFGLSEEYDRLQISRLIPLLRYWPEENL
Secondary structure
SS confidence
Disorder
? ? ? ?
Disorder confidence
490
500
510
520
530
540
Sequence
AIQVTVESLIRPRFQRWLRDTLMQCEKSQRKRIIIELAEADVGQHISRLQPVIRLVNALG
Secondary structure
SS confidence
Disorder
? ?
Disorder confidence
550
560
570
580
590
600
Sequence
VRVAVNQAGLTLVSTSWIKELNVELLKLHPGLVRNIEKRTENQLLVQSLVEACSGTSTQV
Secondary structure
SS confidence
Disorder
? ? ?
Disorder confidence
610
620
630
640
Sequence
YATGVRSRSEWQTLIQRGVTGGQGDFFASSQPLDTNVKKYSQRYSV
Secondary structure
SS confidence
Disorder
? ? ?
Disorder confidence
Confidence Key
High(9)
Low (0)
? Disordered
Alpha helix
Beta strand