

1: At_CNA
2: bCh_Chut02001746
Matrix: EBLOSUM62
Gap_penalty: 15.0
Extend_penalty: 0.6

Length: 147
Identity: 35/147 (23.8%)
Similarity: 60/147 (40.8%)
Gaps: 14/147 (9.5%)
Score: 75.6

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At_CNA	1	WICQSYRGYMGVELLKSNSDGNESILKNLWHHTDAIICCSMKALPVFTFA	50
		.:.: ... :.:. . .: .:.:. . : .:.:.:.:.:.:. . .:.:	
bCh_Chut02001	1	MLIESYREITGIQLFDNTY-SDEYLSCLLYHAPFVVVSHGIEADPVFNYG	49
At_CNA	51	NQAGLDMLE-----TTLVALQDISLEKIFDDNGRKTLCSEFPQIMQOGF	94
	:.. :.:.:.:.:. . :.:. :.	
bCh_Chut02001	50	NLTAQQLWHIDWEQFTTTPSRLSAEPERAED---RQRLLE---AATHGF	93
At_CNA	95	ACLQGGICLSSMGRPVSYERAVAWKVLNEEENAHCICFVFINWSFV	140
	 :.:. :.:. . :.:. . :.:. . :.:. . :.:. . :.:. . :.:. . :.:. . :.:.	
bCh_Chut02001	94	ISNYTGIRISSKGQRFKIEQVLLWNLKNNSGEKIGQAALFRNWTNVK	140