

AtCLO1	MGSKTEMMERDAMATVAPYAPVTYHRRARVDLDDRLPKPYMPRALQAPDREHPYGT PGHK	60
EgCLO1	-----MTDAKDAMQTVAAQAPVTAERRVREDMEEYLPKPYMARALAAPDVNHPEGT KDHM	55
	:*** ** *	
AtCLO1	NYGLSVLQQHVVSFFDIDDNGIIYPWETYSGLRMLGFNIIGSLIIAAVINLTLSYATLPGW	120
EgCLO1	HYNMSVLQQHVAFFDQDDNGIVYPWETYSASRAIGFNVIISLLMAIVINGAMS WRTLPGW	115
	:*. :*****:*** *****:*****. * :***:* **:*: * *** :*: *:*****	
AtCLO1	LPSPPFPFIYIHNIHKSKHGSDSKTYDNEGRFMPVNLELIFSKYAKTLPDKLSLGELWEMT	180
EgCLO1	IPSPFPFIYVHNHVKKHGSDSGTYDTEGRFLPANFENIFSKYARTVPDRLTFGEMWRMT	175
	:*****:***** ***** ** *.****:*.*:* *****:*:**:*::**:*.* **	
AtCLO1	EGNRDAWDIFGWIAGKIEWGLLYLLARDEEGFLSKEAIRRCFDGSLFEYCAKIYAGISED	240
EgCLO1	ESNRVAFDFFGWTAAKLEWIILYILARDEDGFLSKEAIRRCFDGSLFEYCARQRKSNDKT	235
	*.** *:*:*** *.*:** :*:*****:*****:*****:*****:	
AtCLO1	KTAYY	245
EgCLO1	S----	236
	.	