

		(***)		(**)
AtCLF	(151)	NCSFLFNLNDQ---FVLDA	AYRKGDKLKFANHSPE	PNCYAKVIMVAGDHRV
PpCLF	(147)	NSSFLLFNLNDQASLYVLD	ACRKGDKLKFANHSPT	PNCYAKVIMVSGDHRV
DmEz	(57)	MCSFLFNLNND---FVVD	ATRKGDKIRFANHSIN	PNCYAKVMMVTGDHRI
HsEZH2	(57)	MCSFLFNLNND---FVVD	ATRKGDKIRFANHSVN	PNCYAKVMMVNGDHRV
vSET	(47)	LEDYLF	SRKNMS-----AM-ALGFGAIF	NHSKDP--NARHELTA
Consensus	(151)	MCSFLFNLNN	FVLDA RKG KIKFANHS	PNCYAKVIMVAGDHRI

(\*)

AtCLF	(198)	GIFAKERI	ILAGEELFYDYRYE	PDRA	PAWAKKPEAPGSKKDENVTPSVGRP
PpCLF	(197)	GIFAKERI	IGAGEELFYDYQYE	PDRA	PVWARKPDDPNNK-RDDMPSTGGRA
DmEz	(104)	GIFAKRAI	IQPGEELFFDYRYG	PTEQLKFVGI	EREMEIV-----
HsEZH2	(104)	GIFAKRAI	IQTGEELFFDYRYS	QADALKYVGI	EREMEIP-----
vSET	(88)	RIFTIKPI	IAIGEEITISYG	DDYWLSRP	RLTQN-----
Consensus	(201)	GIFAKR I	GEELFFDYRYEP	A W V	E