

EFE_M_stipitatus	MIELETFQLPQSVSGREADIALGLTMVRAWRRDGIQVRMSPAQAQAEKSQRAFELSRHFFR	60
EFE_P_syringae	MTNLQTFELPTEVTGCAADISLGRALIQAQKDGIFQIKTDSEQDRKTQEAMAASKQFCK	60
	* :*:***:** .*: * **:* ** :::**::*****: . * .*:*. * : **:* :	
EFE_M_stipitatus	QSLETKARCVSDLTYSGYIASGQELTASEADLSEVFTVCRDVPLTDPRVQSKWPCGPGP	120
EFE_P_syringae	EPLTFKSSCVSDLTYSGYVASGEEVTAGKPDFPEIFTVCKDLSVGDQRVKAGWPCHGVP	120
	: * * : *****:* **:* ** . : * : *****:* : * ** : : ***** *	
EFE_M_stipitatus	WPDESWRQGMQAHAEELGSVGERLLRLIALGLGLDIDALTTLTHDGWHHMRVLRFPARSP	180
EFE_P_syringae	WPNNTYQKSMKTFMEELGLAGERLLKLTALGFELPINTFTDLTRDGWHHMRVLRFPQP	180
	:::*.***. **** .*****:* **:* * *::* **:****** ** : :	
EFE_M_stipitatus	TTTRGIGAHTDYGLLVIAAQDDVGGLYVRPPVEGEKRPRNWLPHESSAGMYEHDEPWTVV	240
EFE_P_syringae	TLSRGIGAHTDYGLLVIAAQDDVGGLYIRPPVEGEKRNRNWLPGESSAGMFEHDEPWTFV	240
	* :*****:*****:***** ***** *****:*****:*	
EFE_M_stipitatus	KPVPGLTVFPGDILQFLTRGYLLSTPHKVVLNTRERFALAYFHEPQFEACVRPLSAPTR	300
EFE_P_syringae	TPTPGVWTVFPGDILQFMTGGQLLSTPHKVKLNTRERFACAYFHEPNFEASAYPLFEP	300
	.*.*** *****:* * ***** ***** *****:***. ** * :	
EFE_M_stipitatus	DEYIHYGTHFTNMFMRSPDRVTTQRILDESRLTTLNWLRLQEAFLRTAPLEAVPLQRAAG	360
EFE_P_syringae	NERIHYGEHFTNMFMRSPDRITTTQRINKENRLAHLLEDLKKYSRTRATGS-----	350
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