

gi|4758204|ref|NP_004408.1| MV-MEVGILDAGGLRALLGERAAQCLLLDCRFFFAFNAGHIAGSVNVRFTIIVRRRAKAMGLEHIVPNAELRGRLLAGAYHAVVLLDERSAALDGAKRDTLALAAGALCREARAQVFFLKGGYEAFFSASCPPELCSKQSTPMGLSLPL 150
gi|114603388|ref|XP_001153609.1| MV-MEVGILDAGGLRALLGERAAQCLLLDCRFFFAFNAGHIAGSVNVRFTIIVRRRAKAMGLEHIVPNAELRGRLLAGAYHAVVLLDERSAALDGAKRDTLALAAGALCREARAQVFFLKGGYEAFFSASCPPELCSKQSTPMGLSLPL 150
gi|73953410|ref|XP_546235.2| MV-MDVGALDAGGLRLLRERAAQCLLLDCRFFFAFNAGHIAGSVNVRFTIIVRRRAKAMGLEHIVPNAELRGRLLAGAYHAVVLLDERSAALDGAKRDTLALAAGALCREARAARVFFLKGGYEAFFSASCPPELCSKQSTPMGLSLPL 150
gi|7305423|ref|NP_038670.1| MV-MEVGILDAGGLRALLRERAAQCLLLDCRFFFAFNAGHIAGSVNVRFTIIVRRRAKAMGLEHIVPNAELRGRLLAGAYHAVVLLDERSAALDGAKRDTLALAAGALCREARSTOVFFLKGGYEAFFSASCPPELCSKQSTPMGLSLPL 150
gi|16758604|ref|NP_446221.1| MV-MEVGILDAGGLRALLRERAAQCLLLDCRFFFAFNAGHIAGSVNVRFTIIVRRRAKAMGLEHIVPNAELRGRLLAGAYTTPVLLDERSAALDGRKRDGTLALAAGALCREARSTOVFFLKGGYEAFFSASCPPELCSKQSTPMGLSLPL 150
gi|114050891|ref|NP_001039917.1| MV-MEVVILDAGGLRLLRERAAQCLLLDCRFFFAFNAGHIAGSVNVRFTIIVRRRAKAMGLEHIVPNAELRGRLLAGAYHAVVLLDERSAALDGAKRDTLALAAGALCREARSAQIFLLKGGYEAFFSASCPPELCSKQSTPMGLSLPL 150
gi|50764356|ref|XP_422941.1| MVNLRVCALECEALRGLLQERAAQCLVLDRCRFFFAFNAGHIAGSVNVRFTIIVRRRAKAVLAHEVVPNEELRTRLRQGGVHTVLLDERSADLLEPKRDSIFLLALGTLCREARGARICFLKGGYEAFFSASCPPELCSKQSTPMGLSLPL 150
gi|47085789|ref|NP_998232.1| MVIMEVPIIDASLRDMLRGGDDPDCVLDRCRFFFAFNAGHIAGSVNVRFTIIVRRRAKAMGLEHIVPNAELRGRLLAGAYTTPVLLDERSAALDGRKRDGTLALAAGALCREARSTOVFFLKGGYEAFFSASCPPELCSKQSTPMGLSLPL 150



gi|4758204|ref|NP_004408.1| STS-VPD SAESGCCSSCSTPLYDQGGPVEILLPFLYLGSAYHASRKDMLDALGITALINVSANCPNHFEGHYQYKSIIPVEDNHKADISSWFNEAIDFIDSIKNAGGRVVFVHCQAGISRSATICLAYLMRTNRVKLDEAFEFVKORRSIISP 300
gi|114603388|ref|XP_001153609.1| STS-VPD SAESGCCSSCSTPLYDQGGPVEILLPFLYLGSAYHASRKDMLDALGITALINVSANCPNHFEGHYQYKSIIPVEDNHKADISSWFNEAIDFIDSIKNAGGRVVFVHCQAGISRSATICLAYLMRTNRVKLDEAFEFVKORRSIISP 300
gi|73953410|ref|XP_546235.2| STS-VPD STESGCCSSCSTPLYDQGGPVEILLPFLYLGSAYHASRKDMLDALGITALINVSANCPNHFEGHYQYKSIIPVEDNHKADISSWFNEAIDFIDSIKNAGGRVVFVHCQAGISRSATICLAYLMRTNRVKLDEAFEFVKORRSIISP 300
gi|7305423|ref|NP_038670.1| STS-VPD SAESGCCSSCSTPLYDQGGPVEILLPFLYLGSAYHASRKDMLDALGITALINVSANCPNHFEGHYQYKSIIPVEDNHKADISSWFNEAIDFIDSIKNAGGRVVFVHCQAGISRSATICLAYLMRTNRVKLDEAFEFVKORRSIISP 300
gi|16758604|ref|NP_446221.1| STS-VPD SAESGCCSSCSTPLYDQGGPVEILLPFLYLGSAYHASRKDMLDALGITALINVSANCPNHFEGHYQYKSIIPVEDNHKADISSWFNEAIDFIDSIKNAGGRVVFVHCQAGISRSATICLAYLMRTNRVKLDEAFEFVKORRSIISP 300
gi|114050891|ref|NP_001039917.1| STS-VPD SAESGCCSSCSTPLYDQGGPVEILLPFLYLGSAYHASRKDMLDALGITALINVSANCPNHFEGHYQYKSIIPVEDNHKADISSWFNEAIDFIDSIKNAGGRVVFVHCQAGISRSATICLAYLMRTNRVKLDEAFEFVKORRSIISP 300
gi|50764356|ref|XP_422941.1| SASSAPGSADSGCCSSCSTPLYDQGGPVEILLPFLYLGSAYHASRKDMLDALGITALINVSANCPNHFEGHYQYKSIIPVEDNHKADISSWFNEAIDFIDSIKNAGGRVVFVHCQAGISRSATICLAYLMRTNRVKLDEAFEFVKORRSIISP 300
gi|47085789|ref|NP_998232.1| SSNCHSNLADSSCNTCTPLYDQGGPVEILLPFLYLGSAYHASRKDMLDALGITALINVSANCPNHFEGHYQYKSIIPVEDNHKADISSWFNEAIDFIDSIRNKGGRVVFVHCQAGISRSATICLAYLMRTNRVKLDEAFEFVKORRSIISP 300



gi|4758204|ref|NP_004408.1| FFSFMGQLLOFESQVLAP-HCSAEAGSPAMAVLDRGTSITTVFNFPVSIIPVHSTNSALS YLQSPIITSPSC 370
gi|114603388|ref|XP_001153609.1| FFSFMGQLLOFESQVLAP-HCSAEAGSPAMAVLDRGTSITTVFNFPVSIIPVHSTNSALS YLQSPIITSPSC 370
gi|73953410|ref|XP_546235.2| FFSFMGQLLOFESQVLAP-HCSAEAGSPAMAVLDRGTSITTVFNFPVSIIPVHSTNSALS YLQSPIITSPSC 370
gi|7305423|ref|NP_038670.1| FFSFMGQLLOFESQVLAP-HCSAEAGSPAMAVLDRGTSITTVFNFPVSIIPVHSTNSALS YLQSPIITSPSC 370
gi|16758604|ref|NP_446221.1| FFSFMGQLLOFESQVLAP-HCSAEAGSPAMAVLDRGTSITTVFNFPVSIIPVHSTNSALS YLQSPIITSPSC 370
gi|114050891|ref|NP_001039917.1| FFSFMGQLLOFESQVLAP-HCSAEAGSPAMAVLDCSTITTVFNFPVSIIPVHSTNSALS YLQSPIITSPSC 370
gi|50764356|ref|XP_422941.1| FFSFMGQLLOFESQVLAP-HCSAEAGSPAMSVLDRGASTITTVFNFPVSIIPVHTSSALS YLQSPIITSPSC 370
gi|47085789|ref|NP_998232.1| FFSFMGQLLOFESQVLASSTCSSEAGSPAIG-----KNSITVFN-----PVHTAASPLSFLQSPIITSPSC 370

