

TEM2016_SelAC/1-263	1	HPETLVKVKDAEDQLGARVGYIELDLNSGKI	LESFRPEERFPMMS	TFKVL	LCG	53					
TEM2016_SelAC_Simulated/1-263	1	HPETRVRVKVGAEECLGAGRGYIELDLNSGKI	LESFRPEERFPMR	STFKVL	LCG	53					
TEM2016_Consensus/1-263	1	HPETLVKVKDAEDQLGARVGYIELDLNSGKI	LESFRPEERFPMMS	TFKVL	LCG	53					
TEM2016_DMS/1-263	1	SEKVKMAVQQMEWRMG	GHVGYFQIDIMDGDV	LEAWRSKERFPMMS	TMKVILCG	53					
TEM2016_DMS_Simulated/1-263	1	HEKTKTKVRDAERRMG	GRVGYLQIDIHGDV	LESFRQKERFPMMS	TFKVL	LCG	53				
TEM2016_SelAC/1-263	54	AVLSRVDAGQEQLGRRRIHYSQNDLVEYSPV	TEKHLTDGMTVRE	LC	SAAITMSD	106					
TEM2016_SelAC_Simulated/1-263	54	AELSRGDAGQEQLGRRRIHYSQADEVEYSPV	TEKHLTDGMTVRE	LC	SAAVTMDD	106					
TEM2016_Consensus/1-263	54	AVLSRVDAGQEQLGRRRIHYSQNDLVEYSPV	TEKHLTDGMTVRE	LC	SAAITMSD	106					
TEM2016_DMS/1-263	54	CILERVDNGFLKLQRKVKFQVNDLVAWSP	ITMMYIITGMTIQD	LCDAAITLSD		106					
TEM2016_DMS_Simulated/1-263	54	AIIYRVDAAGTEKLGRRVHF	TVNDLVAWSP	ITSQYINDGMTIAD	LCDAAITLSD	106					
TEM2016_SelAC/1-263	107	NTAANLLLTITGGPKELTAF	LHNMGDHSVTRLDRWEPE	LN	EAI	PNDERD	TTMPA	159			
TEM2016_SelAC_Simulated/1-263	107	NTAADLLLTITGGRGELTAF	LHNMTDHSVTRLARGAPE	LN	GEAI	PGDECD	TTMPI	159			
TEM2016_Consensus/1-263	107	NTAANLLLTITGGPKELTAF	LHNMGDHSVTRLDRWEPE	LN	EAI	PNDERD	TTMPA	159			
TEM2016_DMS/1-263	107	NTAANI	LLKELGGPIMLT	MWMNMMGDMY	TRLDRWEPY	LN	MAYEQDERD	TTTPK	159		
TEM2016_DMS_Simulated/1-263	107	NTAANI	LLKSLGGPIELTE	YMN	NMGDNV	TRLDRWEPY	LN	AATPAD	DERD	TTTPK	159
TEM2016_SelAC/1-263	160	AMATTLRKLLTGELLTLASRQQ	LIDWMEADKVAGPLL	RSAL	PAGWF	IADKSGA	212				
TEM2016_SelAC_Simulated/1-263	160	AMAQTLRGLL	TEELLTLASRARL	IDWMEADKVAGPLL	RSCL	PAGWF	IADKSGA	212			
TEM2016_Consensus/1-263	160	AMATTLRKLLTGELLTLASRQQ	LIDWMEADKVAGPLL	RSAL	PAGWF	IADKSGA	212				
TEM2016_DMS/1-263	160	SMADTIKQMLKTHHLSFNS	QQILISW	MYMDKVAGPLL	RQKIPADWY	IADKSGA	212				
TEM2016_DMS_Simulated/1-263	160	VMAKTIHEL	LLKDHRLSKGS	QQILIE	WMKLDKVAGPLL	RQAI	PADWY	IADKSGA	212		
TEM2016_SelAC/1-263	213	GERGSRGI	IAALGPDGKPSRIVV	IYMTGSQATMDERNRQ	IAEIGASLI	KHW	263				
TEM2016_SelAC_Simulated/1-263	213	EVRGSGGI	IAALGPDGKPSRIVV	IYVTGRQATMDERSRQGE	EIGASLI	KRW	263				
TEM2016_Consensus/1-263	213	GERGSRGI	IAALGPDGKPSRIVV	IYTTGSQATMDERNRQ	IAEIGASLI	KHW	263				
TEM2016_DMS/1-263	213	GDHGSRGI	VALMGPNKHMERV	IIYMTGSNANMI	QRNQWFKEIGKNI	IKNW	263				
TEM2016_DMS_Simulated/1-263	213	GKHGSRGI	VAAIGPAGVASRVI	IIYLTGSNNNMD	ARNQWF	AEIGKNI	IKNW	263			