

fset \$\frac{1}{2}\$	↓ 1 ↓ 25	↓ 50	↓ 75	\$ 100	\$ 125	↓ 150	0	↓ 175	\$ 200	\$ 225	\$ 250	\$ 275	\$ 300	\$ 325	\$ 350		1 375	↓ 400	↓ 425	\$ 450	↓ 475	\$ 500	\$ 525	\$ 550	\$ 5	.575	1 600	↓625	1 650	↓ 675	↓ 700	1 725	\$ 750	\$ 775	\$800	\$ 825	\$ 850
omo sapiens	M F S M R I V C L V L S V V G T A W T A D S G E G D F L A E G G G V	R G P R V V E R H Q S A C K D S D W P F C S D E	CDWNYKCPSGCRMKGLIDEVNQDFT	NRINKLKNSLFEYQKNNKDSHSL	TTNIMEILRGDFSSANN <mark>RDNTY</mark> N <mark>R</mark> V	R V S E D L R S R I E V L K R K V I E K V Q H I	I Q L L Q K N V R A Q L V D M K R L E V D	D I D I K I R S C R G S C S R A L A R E V D L	D L K D Y E D Q Q K Q L E Q V I A K D L L P S I	R D R Q H L P L I K M K P V P D L V P (N F K S Q L Q K V P P E W K A L T D M P Q M R	MELERPGGNEITRGGSTSTGTGTES	R N P S S A G S W N S G S S G P G S S	T G N R N P G S S G T G G T A T W K P G S S G P G S	T G S W N S G S S G T G S T G N Q N P G S P	PRPGSTGTWNPGSSERGSAGH	HWTSESSVSGSTGQWHSESG	FRPDSGSGNARPDWGTF	E V S G N V S P G T R R E Y H T E K L V T S K G	DKELRTGKVTSGSTTTRRRSCS	K T V T K T V I G P D G H K E V T K E V V T S E D (G S D C P E A M D L G T L S G I G T L D G F R H	HR HP DE A A F F D T A S T G K T F P G	P G F F S P M L G E F V S E T E S R G S E S G I	I F T N T K E S S S H H P G I A E F P S R G K	SSSYSKQFTSSTSYNRGDSTF	E S K S Y K M A D E A G S E A D H E G T H S T K R G H A	K S R P V R D C D D V L Q T H P S G T Q S G I	I F N I K L P G S S K I F S V Y C D Q E T S L	L G G W L L I Q Q R M D G S L N F N R T W Q D Y K R G	F G S L N D E G E G E F W L G N D Y L H L L T Q	Q R G S V L R V E L E D W A G N E A Y A I	E Y H F R V G S E A E G Y A L Q V S S Y E G	T A G D A L I E G S V E E G A E Y T S H N N M Q F	F S T F D R D A D Q W E E N C A E V Y G G G W W Y N N	C Q A A N L N G I Y Y P G G S Y D P R N N S P Y	E I E N G V V W V S F R G A D Y S L R A V R M K I R P L
	ADSGEGDFLAEGGGV	R		DSHSL	TTNIMEILR					KPVPDLVPC	N F A L T D M P Q M R	ELERPGGNEIT GGSTSYGTGSETESP	R	NPGSSGTGGTATWK		GSAGH	HWTSESSV SESG	S F R P D S P G S G N A P N N P D W G T F	EV REYHTEKLVTSKG	DKEL TGKEKVTSGSTTTTR	TVIGPDGHKEVT	DLGTLSGIGTLDGFR		LGEFVSETESR	ESSSHHPGIAEFPSR	SSSYSKQFT NRGDSTF	ESKS MADEAGSEADHEGTHSTKR										
	SGEGDFLAEGGGV	R		SI	TTNIMEILR					VPDLVPO	N F	MELERPGGNEIT YGTGSETESP	R			SAGE	HWISESSV		REYHTEKL V	EKVTSGSTTTTR	TVIGPDGHKEV	TLSGIGTLDGFR		EFVSETESR	ESSSHHPGIAEFPSKG	QFISSISYNK	ADEAGSEADHEGTHSTKR										
	GEGDFLAEGGGV	R			TTNIMEILR							MELERPGGNEI GTGSETESP	RNPS						REYHTEKLVTS		TVIGPDGH	LSGIGTLDGFR		SETESRGSESGI	IFT SSSHHPGIAEFPSRG		ADEAGSEADHEGTHST										
	EGDFLAEGGGV	R										ELERPGGNEITR							REYHTEKLVT		IGPDGHKEVT	SGIGTLDGFR		GSESGI	I FTNTKESSSHHPG I A E FPSRG		DEAGSEADHEGTHSTKRGH										
	G D F L A E G G G V	R										LERPGGNEITB							HTEKLVTSKG	DKEL	IGPDGHKEV	l-	HRHPDEAAFFDTASTGK	GSESGI	IFTNTKESS HHPGIAEFPS		DEAGSEADHEGTHST										
	DIEXEGGGV	"										ERPGGNEITR							LVTSKG	DKEL		· · · · · · · · · · · · · · · · · · ·	THE BEART		SSHHPGIAEFPS		DEAGSEADHEGTHSTKR										
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omology model																																					
eptide	signal peptide																																				
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etal ion-binding site		_			_																														Ci Ci Ci		
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m regex DE	DEG N TRG LysEnd MOD Glc LIG SH3	3 MOD Glc DOC USP7	LIG LIR Apic 2 TRG LvsEnd LIG FH	HA 1 MOD NEK2 1 MOD GSK3 1	CLV Separi MOD Gld LIG LIR Gen 1	n 1 MOD CK2 1 TRG NLS Mono LIG CaM	aM IQ 9 DOC CYCL	MOD GSK3 1 LIG WD	WD4 LIG SH2 DOC CYCL T	DOC MAPK 1 LIG MYND	MOD PIKK 1 LIG LIR Ne	MOD GSK3 1 MOD GSK3 1	MOD GSK3 1 MOD Glo	DOC USP7 MOD GSK3 1	MOD GIC LIG SH3 3	MOD CK1 1 MOD PKA 2	MOD PLK MOD GSK3 1	MOD ProDKin 1 LIG SH3 3 LIG FHA	DOC WW Pir CLV N CLV PCSK	MOD CK2 1 MOD Gld MOD PKA 2	LIG SH3 3 CLV PCSK LIG FHA 2	MOD GSK3 1	ER MOD NEK2 1 LIG LIR	LIR Ne MOD GSK3 1 MOD GSK3	1 MOD GSK3 1	LIG SH2 MOD CK1 1 MOD SUI	JMO rev 2	MOD GSK3 1	LIG UBA3 DOC USP7 LIG SH2	LIG 14-3-3 2 LIG FHA 2 LIG	14-3-3 2 LIG LIR G¢ M	MOD PKA 2	Nem MOD CK2 1 LIG SH2 MOD	PLK MOD CK2 1 LIG SH2 MOD N	NEK2 1 LIG WD40 WDF LIG S	DOC USP7 MOD N-GLC	MOD NEK2 2 DOC C
	Disordered DEG_N TRG_LysEnd_ MOD_Gld LIG_SH3 MOD_NEK2_1		MOD_Glc DOC_CY	LIG_PTB_Apo_2 LIG_	FHA_1 LIG_R(LIG_LIR_Nem_	m_3 LIG_WD40_WDR5_ CLV_N TRG_L	G_LysEnd_ DEG_APCC_DBOX_1 LIG_W	WD40_W MOD_PKA_2	CLV_PCSK	DEG_APCC_DBOX_1	LIG_LIR_Ge	MOD_CK2_1	MOD_GIG MOD_GSK3_1	DOC_USP7 MOD_Glc N	OD_CK1_1 MOD_CK1_1 MOD_ProDK	DKin_1 MOD_GSK3_1 MO	DD_GSK3_1 LIG_14-3-3	B DOC_WW_Pir MOD_CK2_1	MOD_ProDKin_1 LIG_FHA_1	LIG_FHA_2 MOD_CK1_1 LIG_14-3	DOC_USP7 MOD_GSK3_1	MOD_Glc	LIG_TRAF6 LIG_P	G_Pex14_ MOD_CK2_1 MOD_GI	Lic	LIG_14-3-3_3 MOD_NEK2_1 MOD_GSK3_1	1 CLV_P	MOD_PIKK_1	MOD_GSK3_1 LIG_SH2	MOD_NEK2_1 LIG_LIR_Ne	LIG_LIR_Ne	LIG_WD40_WDF	MOD_LATS_1 MOD_CK2_1	CLV_C14_C MOD_PLK	MO	MOD_ProDKin	n_1 MOD_PLK
	MOD_NEK2_1			MOD_GK2_1 MOL	_CKI_I LIG_H(IRG_EN MOD_	_EINI	THG_NES_CHMT_1 LIG_WD40_WDF			LIG_UBA3_	CLV_PCSK_	MOD_CK1_1 LIG_FHA_2	MOD_CK1_1	MOD_Glc	MOD_GSK3_1 DOC_WW_P	_PII DOC_USP/ MOD_GIÇ	MOD_GIC LIG_BI	CT_ MOD_CK1_1 LIG_LIR_N¢ LIG_LIR_N¢ LIG_LIR_N¢	R_Ge MOD_CK2_1	LIG_14-3-3_3 CLV_N	DOC_USP7	G_WD40_WDK5_	MOD_GSK3_1 M	DOC_WW_Pir MOD_PKA_2		MOD_CK1_1 MOD_GSK3_1 MOD_PLK	CLV_P	MOD_Gic LIG_WI	WD40_WDH5_ LIG_BHCT_ MOD_PLK Glc LIG_LIR_Nem_3	LIG_BRCT_	TRG_ENI		MOD_NEK2_1 MOD_GSK3_1	LIG_TRA		MOD_CK2_1 DOC_WW_Pir	r LIG_WD40_W
						CLV_P	CLV_P LI	LIG_WD40_W				MOD_ProDk	Kin_1 Pir					LIG_SH3_3	MOD_CK1_1 OD_GICNHGI LIG_FHA_2	CLV_P	CLV_PCSK_			LIG_BRO	CT_	LIG_BRCT_	LIG	_14-3-3_3	MOD_PK_1	LIG_Pex14_	[LIG_SH2]	-3 3	LIG_	FHA_2			
							DOC_MAPK_	_1				LIG_SH3_	_3						OD_GSK3_1 LIG_elF4E_1		LIG_I II/L_I					OLV_I OOIÇ				MOD_NEK2_1	MOD_GSK:	SK3_1					
							MOD_	_SUMO_rev_2																							MOD_PIKK	KK_1					