

**Table S1.** Force constants and equilibrium distances of coordinating atoms to metal centres, as calculated from B3LYP/ Def2-TZVP optimisation of the metal sites.

State	Ligating Bond	Force constant (kCal/mol·Å <sup>-2</sup> )	Equilibrium Distance (Å)
FU-a <sub>oxy</sub>	HIS43 (N) – CU2	90.1	1.990
	HIS62 (N) – CU2	82.7	1.993
	HIS71 (N) – CU2	7.4	2.372
	HIS184 (N) – CU1	93.1	1.984
	HIS188 (N) – CU1	89.3	1.982
	HIS215 (N) – CU1	11.5	2.352
FU-a <sub>deoxy</sub>	HIS43 (N) – CU2	49.9	2.014
	HIS62 (N) – CU2	50.6	2.011
	HIS71 (N) – CU2	49.3	2.018
	HIS184 (N) – CU1	50.4	2.013
	HIS188 (N) – CU1	49.0	2.019
	HIS215 (N) – CU1	50.4	2.012

**Table S2.** Angles of the metal ion centres and ligating atoms, including force constants, calculated using B3LYP/ Def2-TZVP.

Oxygenation State	Ligating atoms	Force constant (kCal/mol·rad <sup>-2</sup> )	Angle (°)
FU-a <sub>oxy</sub>	HIS43 (N) – CU2 – HIS62 (N)	102.61	97.22
	HIS43 (N) – CU2 – HIS71 (N)	32.44	91.30
	HIS62 (N) – CU2 – HIS71 (N)	44.07	93.73
	HIS184 (N) – CU1 – HIS188 (N)	100.00	97.30
	HIS184 (N) – CU1 – HIS215 (N)	25.83	91.99
	HIS188 (N) – CU1 – HIS215 (N)	42.50	93.51
FU-a <sub>deoxy</sub>	HIS43 (N) – CU2 – HIS62 (N)	2.87	121.86
	HIS43 (N) – CU2 – HIS71 (N)	2.96	118.16
	HIS62 (N) – CU2 – HIS71 (N)	2.98	119.98
	HIS184 (N) – CU1 – HIS188 (N)	2.99	118.33
	HIS184 (N) – CU1 – HIS215 (N)	2.86	122.09
	HIS188 (N) – CU1 – HIS215 (N)	2.95	119.58

**Table S3.** Mean equilibrium distances and associated variance of coordinating distances from ligating atoms to metal centres, as calculated from molecular dynamic trajectories.

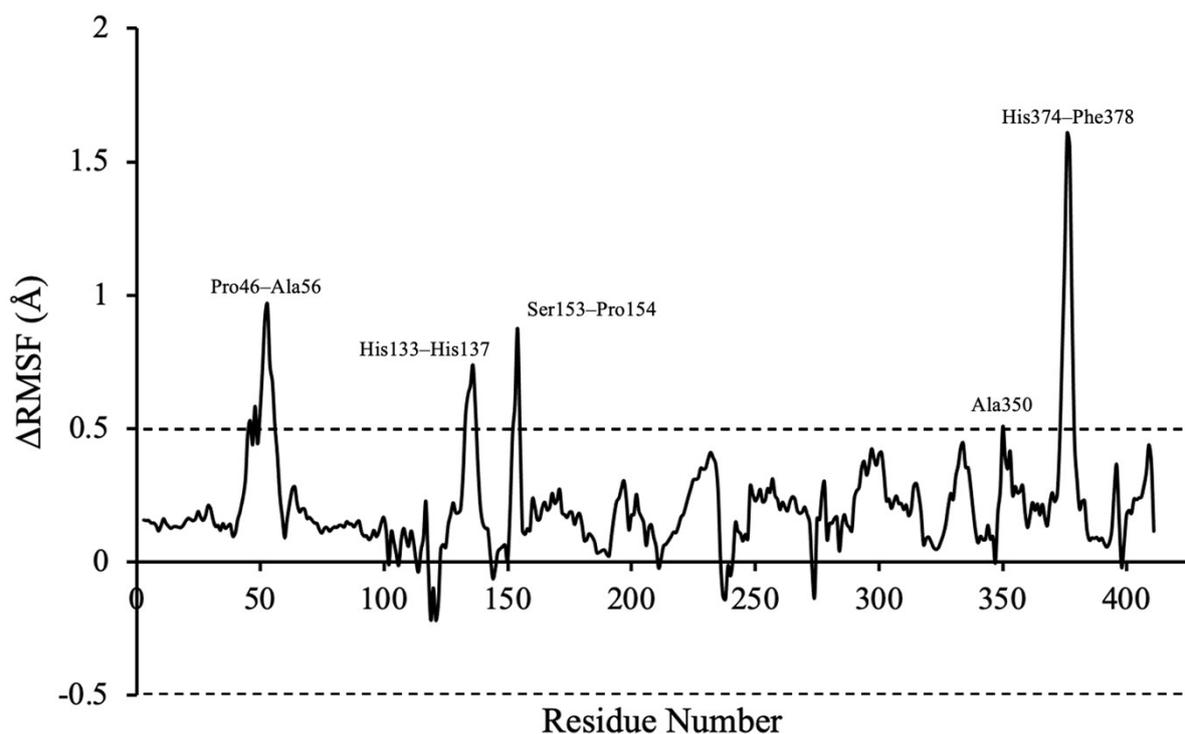
Oxygenation State	Ligating Bond	Equilibrium Distance (Å)	SD (±)
FU-a <sub>oxy</sub>	HIS43 (N) – CU2	2.03	0.06
	HIS62 (N) – CU2	2.00	0.06
	HIS71 (N) – CU2	2.49	0.15
	HIS184 (N) – CU1	1.97	0.05
	HIS188 (N) – CU1	2.01	0.05
	HIS215 (N) – CU1	2.44	0.14
FU-a <sub>deoxy</sub>	HIS43 (N) – CU2	2.40	0.14
	HIS62 (N) – CU2	2.33	0.08
	HIS71 (N) – CU2	2.20	0.02
	HIS184 (N) – CU1	2.20	0.02
	HIS188 (N) – CU1	2.27	0.15
	HIS215 (N) – CU1	2.20	0.02

**Table S4.** Mean binding free energy ( $\Delta G_{\text{bind}}$ ) values for FU-a<sub>deoxy</sub> and FU-a<sub>oxy</sub> systems calculated using MM/GBSA across internal dielectric constant values ranging from 1 to 5.

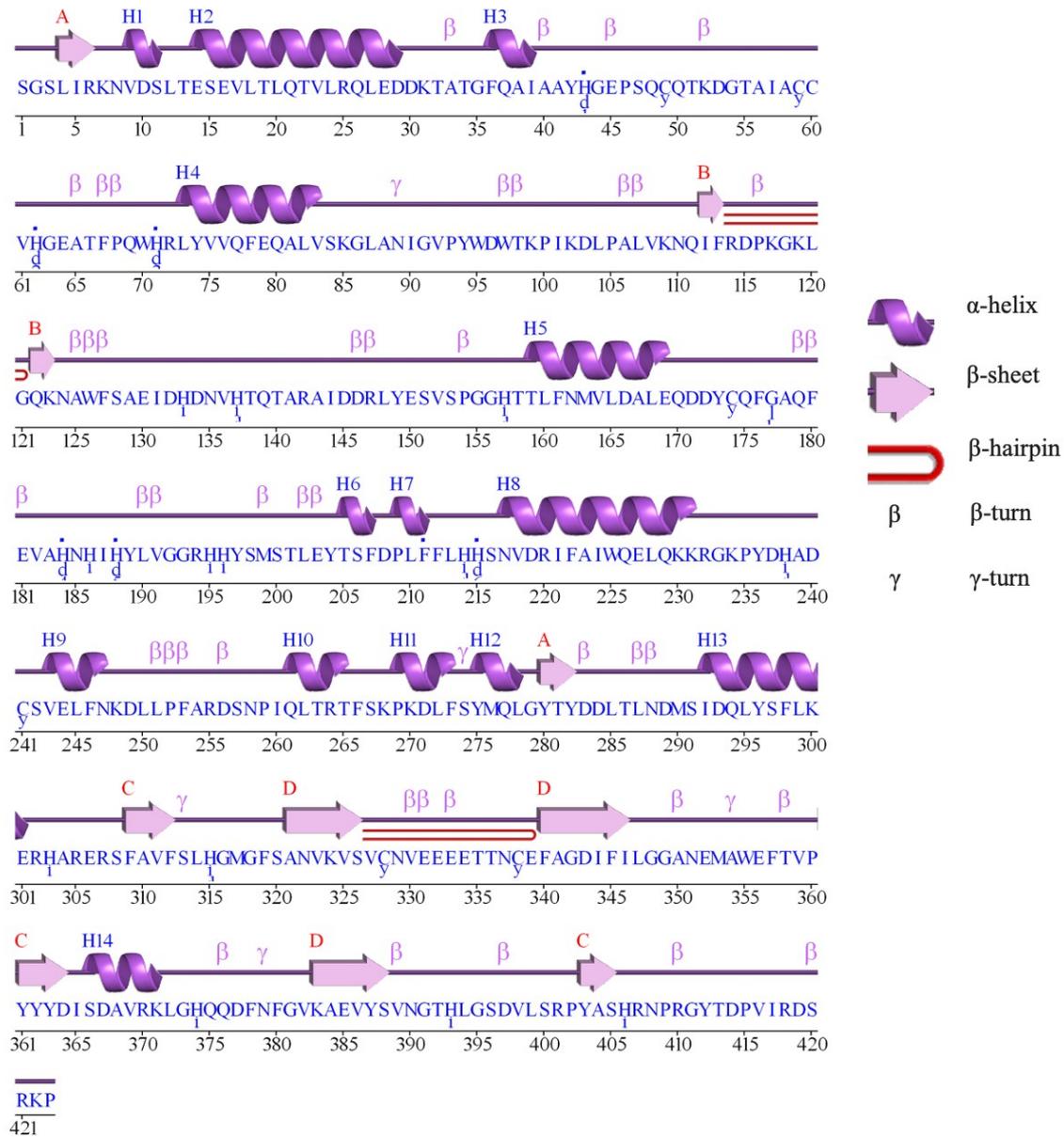
Oxygenation State	IDC	$\Delta G_{\text{bind}}$ (kcal mol <sup>-1</sup> )	SD (±)
FU-a <sub>oxy</sub>	1	-72.77	1.98
	2	-38.36	1.56
	3	-26.89	1.44
	4	-21.16	1.38
	5	-17.72	1.35
FU-a <sub>deoxy</sub>	1	-0.92	3.04
	2	-1.15	3.04
	3	-1.23	3.04
	4	-1.27	3.04
	5	-1.29	3.04

**Table S5.** Mean binding free energy ( $\Delta G_{\text{bind}}$ ) values for FU-a<sub>deoxy</sub> and FU-a<sub>oxy</sub> systems calculated using MM/PBSA across internal dielectric constant values ranging from 1 to 5.

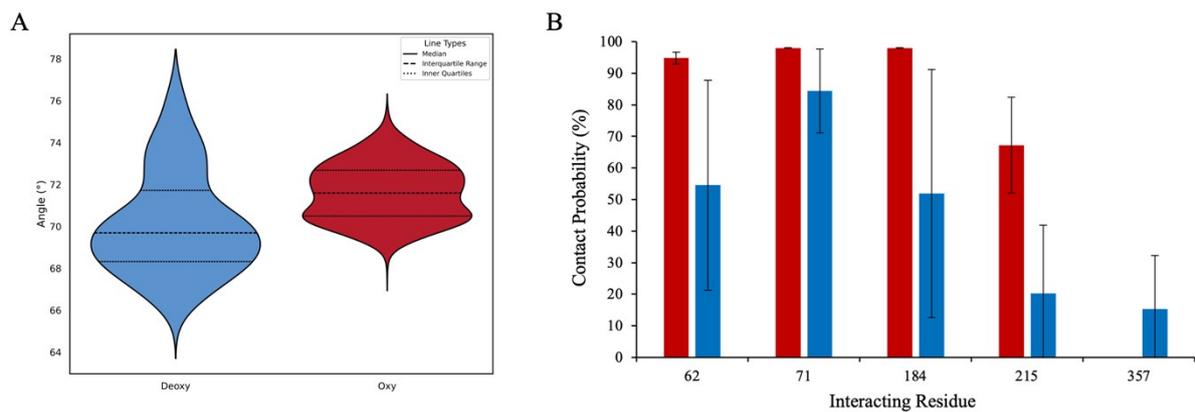
Oxygenation State	IDC	$\Delta G_{\text{bind}}$ (kcal mol <sup>-1</sup> )	SD ( $\pm$ )
FU-a <sub>oxy</sub>	1	-80.31	2.112
	2	-42.53	1.649
	3	-29.93	1.514
	4	-23.63	1.451
	5	-19.85	1.415
FU-a <sub>deoxy</sub>	1	-4.29	2.986
	2	-4.60	3.003
	3	-4.70	3.009
	4	-4.75	3.012
	5	-4.78	3.014



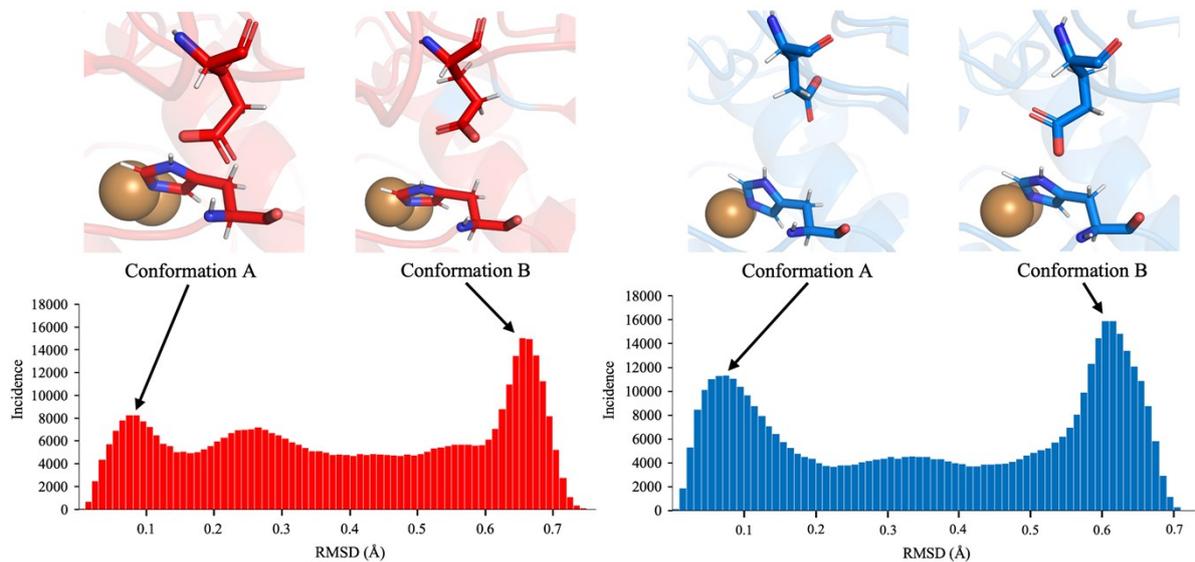
**Figure S1.** Per-residue differences in root-mean-square fluctuation (RMSF) between FU-a<sub>deoxy</sub> and FU-a<sub>oxy</sub> systems, calculated as  $\Delta\text{RMSF}$  (FU-a<sub>deoxy</sub> – FU-a<sub>oxy</sub>). Residues with absolute RMSF differences exceeding  $\pm 0.5$  Å are highlighted. The dotted line denotes the  $\pm 0.5$  Å threshold used to define significant deviations. Major fluctuations are observed in labelled regions.



**Figure S2.** Two-dimensional secondary structure map of the FU-a hemocyanin domain. The structure is divided into two domains: Domain 1 (residues 1–305) and Domain 2 (residues 306–423), as indicated along the sequence.  $\alpha$ -helices are depicted as coils,  $\beta$ -sheets as arrows,  $\beta$ -hairpins as loops, and  $\beta$ -turns ( $\beta$ ) and  $\gamma$ -turns ( $\gamma$ ) are explicitly labelled.



**Figure S3.** Structural and Functional Insights into Domain Rotation and Phe67 Residue Interactions. (A) Violin plot illustrating the inward rotation of domain II relative to domain I in the oxygenated (FU-a<sub>oxy</sub>) and deoxygenated (FU-a<sub>deoxy</sub>) states. (B) Interaction data showing the  $\pi$ - $\pi$  stacking interactions of Phe67 with coordinating histidines, highlighting the differences between the oxygenated (FU-a<sub>oxy</sub>, red) and deoxygenated (FU-a<sub>deoxy</sub>, blue) states.



**Figure S4.** Comparison of Glu352 side-chain conformations in FU-a<sub>oxy</sub> (red) and FU-a<sub>deoxy</sub> (blue). Conformation A (low RMSD) and Conformation B (high RMSD) represent the dominant states identified from side-chain heavy-atom RMSD analysis.

**Table S6.** Sequence conservation in the interaction network shown in Figure 6. \*From visual inspection of the 3D structure alignment of octopus (PDB 1JS8) and horseshoe crab (PDB 1OXY) hemocyanin.

Amino acid residue	Conservation in functional units of SLH and KLH	Conservation in FU-a of molluscan hemocyanins	Notes
His43	Full	Full	Coordinates Cu1
His62	Full	Full	Coordinates Cu1
Phe67	Full	Full	Structurally conserved in horseshoe crab*
His71	Full	Full	Coordinates Cu1
His184	Full	Full	Coordinates Cu2
Leu202	Full	Full	
Glu203	Glu, Asp, His, Arg	Glu or Asp	
Gly316	No	No	
Gly318	No	No	
Gly348	Full	Full	
Gly349	Gly, Asp	Full	Asp in FU-c only
Glu352	Full	Full	
Trp355	Trp, Phe	Full	Phe in FU-d only

**Table S7.** Protein sequence alignment (Clustal Omega) of the eight functional units (FUa – FUh) of slipper limpet hemocyanin (*Crepidula fornicata*; SLH1; GenBank accession number WCA44164.1) and keyhole limpet hemocyanin (*Megathura crenulata*; MecerH1; UniProtKB/Swiss-Prot accession number Q10583.2), along with the protein sequence of *Enteroctopus dofleini* Odg from PDB 1JS8. FUa of SLH is shown in bold, and the key residues participating in the interaction network shown in Figure 6 are highlighted in yellow.

SLH1_FUh	-----TGDVTRYEVSALELAQVTNLRDALYKLNQNDHGPNGFEEIASFHGA	45
MecerH1_FUh	-----EDILVRKNIHSLSHHEAEELRDALYKLNQNDESHGGYEHIAAGFHGY	45
<b>SLH1_FUa</b>	----- <b>SGSLIRKNVDSLTESEVLTQTVLRQLEDDKATATGFQAI</b> <b>AAAY</b> <b>GE</b>	<b>45</b>
MecerH1_FUa	MLSVRLLIVVLALANAENLVRKSVHEHLTQEETLDLQAALRELQMDSSSIGFQKIAAAHGA	60
SLH1_FUd	-----EVQASSHVRRNLEDLTTGEAESLRSALRDMEDG---SFDAIARFHGY	45
MecerH1_FUd	-----EVTSANRIRKNIENLSLGELESLRAAFLEIENDG---TYESIAKFHGS	45
SLH1_FUf	-----EVAPNRIRHDLTHLSERDIRSLKSSIRDLQLDDSDNGYQNIASFHGA	47
MecerH1_FUf	-----HLSLNKVRHDLSTLSERDIGSLKYALSSSQADTSADGFAAIASFHGL	47
SLH1_FUg	-----IAGVGVKRDINTLTAEMENLRDALGRVQAGTGRLTYDNVVS AHGY	46
OcdoH1	-----AIIRKNVNSLTPSDEIKELRDAMAKVQADTS DNGYQKIASYHGI	43
MecerH1_FUg	-----MAGHGVRKEINTLTAEVDNLKDMRAVMADHGPNGYQAI AAFHGN	46
SLH1_FUb	-----SNVVRKDIELLTDEEVYALRQAMHRFQNDTSIDGYQAVAEFHGL	44
MecerH1_FUb	-----INDLTREEVNLREAFHKFQEDRSVDGYQATAEYHGL	37
SLH1_FUc	-----EDDDHVRNDIVTLTAEQEQNVRFDAFQGLKNDKSVAGYNQIAAFHGO	46
MecerH1_FUc	-----PRSRIRKNVDRLSPEEMNELRKALALLKEDKSAGGFQQLGAFHGE	46
SLH1_FUe	-----RLVRKSVLSLSPAERRSLVLAMRSLQEDSSADGFQSLASFHAL	43
MecerH1_FUe	-----VMIRKIDITQLDKRQQLSLVKALESMDKADHSSDGFQAIASFHAL	43
	: * : : : . . : *	
SLH1_FUh	PGLCPENA-TDHYACCRHGMPAFPHWHRLLTQVFERALKDKG-AVVGVPYDWT RPAKAM	103
MecerH1_FUh	PNLCPEKG-DEKYPCCVHGMSIFPHWHRLHTIQFERALKKHG-SHLGIPYDWTQTISSL	103
<b>SLH1_FUa</b>	<b>PSQCQTKDGT-AIACCVHGEATFPQWHRLYVVFQEQALVSKGLANIGVPYDWT</b> <b>KPIKDL</b>	<b>104</b>
MecerH1_FUa	PASCVHKD-T-SIACCIHGMPTFPWHRRAYVVHMERALQTKR-RTSGLPYDWT EPI TQL	117
SLH1_FUd	PGLCEHD-GH-KAACCVHGS PAFPHWHRLYVEQVENSLLSHG-SAVSVPYDWT QPIKKL	102
MecerH1_FUd	PGLCQLN-GN-PISCCVHGMPTFPWHRLYVVVENALLKKG-SSVAVPYDWT KRIEHL	102
SLH1_FUf	PALCPSPEAA-EYACCVHGMPTFPWHRLYTVEMEDAMVRHG-SSVALPYDWT MPITQL	105
MecerH1_FUf	PAKCNDSHNN-EVACCIHGMPTFPWHRLYTLQFEQALRRHG-SSVAVPYDWT KPIHNI	105
SLH1_FUg	PPQCTH-DGH-KVACCVHGMASFPWHRLFTRQMEVALSWEG-AKVGIPYDWT EAFTEL	103
OcdoH1	PLSCHYENG-TAYACCVHGMVTFPNWHRLLTQKQMEDALVAKG-SHVGIPYDWT TTFANL	101
MecerH1_FUg	PMLCPMPDGK-NYSCCTHGMATFPWHRLYTVQMEDALTAHG-ARVGLPYDWT GTAFTAL	104
SLH1_FUb	PAKCPHPDAAVRYACCVHGMATFPWHRLFVVEVEDELRSRG-LEIGIPYDWT RPNNAHI	103
MecerH1_FUb	PARCPRDPAKDRYACCVHGMPIFPHWHRLFVTVQVEDALVGRG-ATIGIPYDWT EPMTHI	96
SLH1_FUc	PNWCPSTQAERYACCTHGMVFPWHRLLTQVAENGLIANG-LHSGLPYDWT LPMTSL	105
MecerH1_FUc	KWCPSPPEASKKFACCVHGMVFPWHRLYTVQSENALRRHG-YDGLPYDWT SPNLHL	105
SLH1_FUe	PPLCPYPEATKRFACCIHGMATFPWHRLYTVQFEDALRRHG-ALVGIPYDWT VVNSRL	102
MecerH1_FUe	PPLCPSPAASKRFACCVHGMATFPQWHRLYTVQFQDSLRRKHG-AVVGIPYDWT LPRSEL	102
	* * * * * : : . : * * * * :	
SLH1_FUh	PSLFTDSY-----DNNPFQTYRMTFNDQY--IQRDV-SEELFNHPSEGDVESL FHQ	151
MecerH1_FUh	PTFFADSG-----NPNPFKYHIRSINQD--TVRDV-NEAIFQQTKFGEFSSIFYL	151
<b>SLH1_FUa</b>	<b>PALVKNQIFRDPKQKLGQKNWAFSAEIDHDNVHTQTARAI-DDRLYESVSPGGHTTLFNM</b>	<b>163</b>
MecerH1_FUa	PSLAADPVYIDSQGGKAHTNYWYRGNIDFL--DKTNRVAV-DDRLFEKVKPGQHTHLSM	174
SLH1_FUd	PALINDATYYDSRAHAKLENPFFRWKIPGTDY--TSRDP-RPDLFDS-----DYFLNN	153
MecerH1_FUd	PHLISDATYYNSRQHHYETNPFHHGKI THENEI--TTRDP-KDSL FHS-----DYFYEQ	153
SLH1_FUf	PDLFTSETYYDAWRDEVANPFARAYIKVAGGY--TVRDP-QAQLKQLSRDQGSALFDL	162
MecerH1_FUf	PHLFTDKEYYDVRNKMVNPFPARGYVPSHDTY--TVRDV-QEGLFHLTSTGEHSALLNQ	162
SLH1_FUg	PALVREEE-----NPNFHHGRI PGTDTV--TTRAP-RPQLFRDPEHGDESFFYRQ	150
OcdoH1	PVLVTEEK-----DNSFHHAHIDVANTD--TTRSP-RAQLFDDPKGDKSFFYRQ	148
MecerH1_FUg	PTFVTDEE-----DNPFFHHGHIDYLGVD--TTRSP-RDKLFNDPERGSE SFFYRQ	151
SLH1_FUb	PALANDVSYEDANTHQQLHNPFLHDAIFLGE--KTTTRQV-QDDLSESPFEGDHTSLFDG	160
MecerH1_FUb	PGLAGNKTYVDSHGA-SHTNPFHSSVIAFEENAPHTKRQI-QDRLFKPATFGHHTDLFNQ	154
SLH1_FUc	PSIVLNATYVHPKTGESVANPLYSGQVDGHD---TTRSV-RQELFEQPKFGHMTKIAEK	160
MecerH1_FUc	PELADHEKYVDPEDGVEKHNPWFDGHIDTVDKT--TTRSV-QNKLFEQPEFGHYT SIAEK	162
SLH1_FUe	PDFIAESVWDDPLPHANFSPWAGADIEFDNSAV--VRDVMNDRISQKGPKYD TWSWKQ	160
MecerH1_FUe	PELLTVSTIHDPE TGRDIPNPFHIGSKIEFEGENVHTKRDINRDLFQGSTKTHHWFIEQ	162
	* : * : *	
SLH1_FUh	ALETLEETNYCDFEVQYEMLHNAVHELIGGNTYSMSTLEYSAFDPFFMVHHSIDRIWQ	211
MecerH1_FUh	ALQALEEDNYCDFEVQYELHNEVHALIGGAEKYSMSTLEYSAFDPYFMHHSALDKIWI	211
<b>SLH1_FUa</b>	<b>VLDALQDDYQCQFEAQFEVAHNIHYLVGGRHHYSMSTLEYSDFLFLHHSNVDRIFA</b>	<b>223</b>
MecerH1_FUa	VLDALQDQDFCKFEIQFELAHNAIHYLVGGKHDYSMANLEYTAYDPIFFLHHSNVDRIFA	234
SLH1_FUd	ALLALEQTSYCDFEVQFEILHNLHLSFLGGRGKYSLSLSDYSAFDPVFFLHHSNVDRIFA	213
MecerH1_FUd	VLYALEQDNFCDFEIQLEILHNLHLSLGLGKYSMSNLDYAAFDPVFFLHHSNVDRIFA	213
SLH1_FUf	VLLVLEQTDYCDFEVQFEVVHNAIHYLVGGRQLYSMSSLEYTSYDPIFFVHHSFVDKIWA	222
MecerH1_FUf	ALLALEQHDYCDFAVQFEVMHNTIHYLVGGPQVYSLSSLHYASYDPIFFI HHSFVDKVWA	222
SLH1_FUg	VQLALEQRDYCDFEVQFEVIHNAIHSWIGGTSYGMSTLEYSAYDPVFFI HHSNVDRQFA	210

Ocdoh1	IALLALEQTDFCDFEIQFEIGHNAIHSWVGSSPYGMSTLHYTSYDPLFLYHHSNTDRIWS	208
MecrH1_FUg	VLLALEQTDFCQFEVQFEITHNAIHSWVGGLTPYGMSTLEYTTYDPLFLWHANTDRIWA	211
SLH1_FUa	ILLAFEQTDFCDFEVQFEVFNHAPHFLVGGFAGYTLATLHYSAFDPIFYLLHHSNVDRLWA	220
MecrH1_FUa	ILYAFEQEDYCDFEVQFEITHNTIHAWTGGSEHFSMSSLHYTAFDPLFYFHHSNVDRLWA	214
SLH1_FUc	VMLAFEQDNFCDFEIQYEIAHNYIHALVGGNETYSMASLRYTAYDPIFFLHHSNTDRIWA	220
MecrH1_FUc	VLLALEQDNFCDFEIQYEIAHNYIHALVGGAQPYGMASLRYTAFDPLFLYHHSNTDRIWA	222
SLH1_FUe	YLVALEQENFCDFEVQFEIAHNNALHAWMGSEVHSMALHLYASYDVPFLLHHSNTDRIWA	220
MecrH1_FUe	ALLALEQNTNYCDFEVQFEIMHNGVHTWVGKPEYIGLHLYASYDPLFYIHHSQTDRIWA	222
	. : * : . : * : * : * : * : * : . : * : * : * : * : * : * : * : * : * : :	
SLH1_FUa	IWQSLQKLRHKPFNYASCASRSLYKPLEPFSYTSL-NADPLTLNNAQPVHIFDTA-KFHY	269
MecrH1_FUa	IWQELQKRRVKAHAGSCAGDIMHVPLHPFNYESV-NNDFFTRENSLPNAVVDSH-RFNY	269
<b>SLH1_FUa</b>	<b>IWQELQKRRKPYDHADCSVELFNKDLLPFARDSN--PIQLTRTFSKPKDLFSYM-QLGY</b>	<b>280</b>
MecrH1_FUa	IWQRLQELRNKDKPKAMDCAQELLHQKMEFFSWEDN--DIPLTNEHSTPADLFDYC-ELHY	291
SLH1_FUd	IWQALQKIRGLPFDESDCALNMGTPHLPFDKDEE-NQFDLTNKSRI DAFDYSNHFDY	272
MecrH1_FUd	IWQDLQRFKRPRYREANCAIQLMHTPLQPFDKSDN-ND-EATKTHATPHDGFYQNSFGY	271
SLH1_FUf	VWQELQKRRGMNYDRADCAVNFNMNERMHPFDWEAL-NPDVTRTRESLQSVWKYE-DLGY	280
MecrH1_FUf	VWQALQEKRRGLPSDRADCAVSLMTQNMRFPHYEIN--HNQFTKKHAVPNDFVKYE-LLGY	279
SLH1_FUg	IWQALQKYRGLDYNTANCIQELRQVQEPFDRDDN--PVVTRHYSKAI DAFNYD-QYGY	267
Ocdoh1	VWQALQKYRGLPYNTANCEINKLVKPLKPFNLDTN--PNAVTKAHSTGATSFDYH-KLGY	265
MecrH1_FUg	IWQALQEYRGLPYDHANCEIQAMKRPLRPFSDPIN--HNAFTHSNAKPTDVFEYS-RFNF	268
SLH1_FUa	IWQQLQIKRGLPYK-AHCASSTQEVLPKFPFPFNWNNHDKTFENARPTNIYDYEAVLDY	279
MecrH1_FUa	VWQALQMRHRKPYR-AHCAISLEHMHKLPFAFSSPLNNEKTHANAMPNKIYDYENVLHY	273
SLH1_FUc	IWQALQKYRGLPYNSANCAIAEMRKLQPFQATSLTNPDPYVTRDRHSTPFDPVFNYSFHY	280
MecrH1_FUc	IWQALQKYRGLPYNVANCAVTSMREPLQPFGLSANINTDHVTKEHSVPFNVDYKTNFNY	282
SLH1_FUe	LWQELQKYRGRDANEVNCALMLKPELKFPSFGSPYLNPTTTLYSKPEDAFDYKGFHY	280
MecrH1_FUe	IWQSLQRFRLGSGSEANCAVNLMTKPLKFPFSGAPYLNLDHDFSKPEDTFDYQ-KFGY	281
	: * * * * * : * * * * * : * * * * * : * * * * * : * * * * * :	
SLH1_FUa	HYDSLNLNGHGVHELHDMIEDMQATSRI FAGFVLSGISTSARVHVDVTRGED--T-----	322
MecrH1_FUa	KYDNLNLNGHNIEELEEVLRSLRSLKSRVFAFVLSGIRTTAVVKVYIKSGTDSDD-----	324
<b>SLH1_FUa</b>	<b>TYDDLTLNDSIDQLYSFLKERHARERSFAGVSLHGMCFSANVKVSVCNVEEE--ETTNC</b>	<b>338</b>
MecrH1_FUa	DYDTLNLNGMTPEELKTYLDERSRRAFASFRLKGFSSANVVFVVCIPDDNDRNDHDC	351
SLH1_FUd	HYDTLKFNSWTIPQLEQVLEKQSRDRVFAGFLLHNIIGTSADVEIDVCVATGNGA-KSCN	331
MecrH1_FUd	HYDNLNLNHYSIPQLDHLQERKRHRDRVFAGFLLHNIIGTSADGHVFCVCLPTGEHT-KDCS	330
SLH1_FUf	AYDNLEIGGKSEIEELEIKKQSHPRVFAGFQLHGLGTSADVELSVCKSQNSC-----	334
MecrH1_FUf	RYDNLEIGGMNLHEIEKEIKDKQHVRVVFAGFLLHGIRTSADVQFQICKTSEDC-----	333
SLH1_FUg	QYDNLNFHGMTISQLDMELEKQEDHVFANFMLHGIQTSADVVDLDCDAKGC-----	321
Ocdoh1	QYDNLNFHGMTIPELEELKEIQHEDRVFAGFLLRTIGQSADVNFVCTKDGC-----	319
MecrH1_FUg	DYDNLRFHGMTIKKLEHELEKQKEEDRTPFAAFLHGIKKSADVSVFCVNDHDC-----	322
SLH1_FUa	TYDSLQFGGMTVDQLEHYLHERKTQNRDFVGIALHNIIGVSAWVTLVSLQVEGGEP-----	333
MecrH1_FUa	TYEDLTFGGISLENIEMKHENQEDRIYAGFLLAGIRTSANVDIFIKTDSVQ-----	327
SLH1_FUc	HYDNLDFNGMSVAQLQREVIRRRGLERAFAGFMLHGKKSCLVDFDICKPDGTC-----	334
MecrH1_FUc	EYDTLEFNGLSISQLNKKLEAIKQSPRVFAGFLLSGFKKSLVKNICTDSSNC-----	336
SLH1_FUe	EYDTLELQGLDQRLQDYINKEKEHDRVAGFLLSGIGQSAHASFSVCKANGEC-----	334
MecrH1_FUe	IYDTLEFAGWSIRGIDHIVRNREQHSRVFAGFLLGFGTTSATVDFQVCRTAGDC-----	335
	* : : : : : : : : * : . : . : . :	
SLH1_FUa	VSVGNFYVLGGSSEMPWAYERIYKLDMTEAASKLGLSSE--STF-----HFKLTVTKYD	374
MecrH1_FUa	EYAGSFVILGGAKEMPWAYERYRFDITETVHNLNLTDD---HV-----KFRFDLKKYD	375
<b>SLH1_FUa</b>	<b>EFAGDIFILGGANEWAEFTVPPYYDISDAVRKLGHQD--FNF-----GVKAEVYSVN</b>	<b>390</b>
MecrH1_FUa	EKAGDFVVLGGPSEMKWQFYRPLFLDLSDTVHKMGMKLD--GHY-----TVKAEVYSVN	403
SLH1_FUd	HPAGKFAILGGEYEMPFTFDRLYKYDISDTRVRLGLRLDLSAADF-----DVQIKIFAYN	385
MecrH1_FUd	HEAGMFSILGGQTEMSFVDFRLYKLDITKALKKNGVHLQ--GDF-----DLEIBITAVN	382
SLH1_FUf	VSAGVIFILGGKLEMPWAFDRFLKLDITDTHLDMGIEPE--DVFDTPAPFFLSYEVHAVN	392
MecrH1_FUf	HSGGTQIFVLGGTKEMAWAYNRFKYDITHALHDAHITPE--DVFPSEPFPIKVSVTAVN	391
SLH1_FUg	NFAGTFAILGGPLEMPWAFDRFLKYDVTSVFKQMLRDP--SEY-----SFRVSLTAVN	373
Ocdoh1	TFGGTFCILGGEHEMFWAFDRFLKYDITSLKHLRLDAH--DDF-----DIKVTIKGID	371
MecrH1_FUg	HFAGTFAILGGEHEMFWAFDRFLRYDITQVLKQMHLEYD--SDF-----TFHMRIIDTS	374
SLH1_FUa	YQVGKIAVLGGEKEMPWFDRFLFKVEITSAQKLGLSYD--DDF-----NVHLDITDVT	385
MecrH1_FUa	HKAGTFAVLGGSKEMKWGDRVFKFDITHVLKDLDTAD--GDF-----EVTVDITEVD	379
SLH1_FUc	TKSGEFYLLGDENELPWEYDRLYKYEITHELEDMLGEPQ--DRF-----DVQYHVYDLK	386
MecrH1_FUc	HPAGEFYLGDENEMPWAYDRVFKYDITEKLHDLKLHAE--DHF-----YIDYEVFDLK	388
SLH1_FUe	TAAGDFDILGGSAMPWFDRFLKYEITDVLKGLDVH--DSF-----NITVSLTALD	386
MecrH1_FUe	EDAGYFTVLGGEKEMPWAFDRFLKYDITETLTKMNLRH--EIF-----QIEVTITSYD	387
	* : : * . * : : : : : : : . : . : . :	
SLH1_FUa	GTALNV-TFPDPVIVKRAANSQHDVVLVPLSVANQLPPKIVVRQGTQVVFHASESGV-SS	432
MecrH1_FUa	HTELDASVLPAPIIVRRPNAVFDIIEIPIGKDVNLPPKVVVRGRTKIMFMSVDEAVTTP	435
<b>SLH1_FUa</b>	<b>GTHLGSVLSRYPYASHRNRPRGYTDPVIRDSR-----KP-----</b>	<b>423</b>
MecrH1_FUa	GTALPDDLLPHPVVVVHHPKGFDPVVKHHQ-----SANLLVRKN-----	443
SLH1_FUd	GSYIDASLLHRPTIIFEPGGQQTDDVGHVE-----R-----	417
MecrH1_FUd	GSHLDSHVIHSPTILFEAGTDSAHTDDGHT-----P-----	414
SLH1_FUf	GTTLPLSTISPPTLVFPQPAEGAAEHSSYS-----	422
MecrH1_FUf	GTVLPASILHAPTIIYEPGLDHHEDHSSS-----	421
SLH1_FUg	GTQLDSRLIEAPSVSFVPGNKPGSKTSAHE-----DPVPLD-----	410
Ocdoh1	GHVLSNKYLSPTTVFLAPAKTTH-----	394

MecrH1_FUg	GKQLPSDLIKMPTVEHSPGGKHHHEKHEDHH-----	405
SLH1_FU <b>b</b>	GKKWAEDTFHHNTIIHVPGEGHEDPG-----	411
MecrH1_FU <b>b</b>	GTKLASSLIPHASVIREHARVKFDKV-----	405
SLH1_FU <b>c</b>	NTDLGDDVFGKAVIIYSAGQGHQKGHEEDYL-----E-----	418
MecrH1_FU <b>c</b>	PASLGKDLFKQPSVIEPRIGHHEGEVYQA-----	418
SLH1_FU <b>e</b>	GSALSSSLLPTPSVIFEPKSRTELH-----	412
MecrH1_FU <b>e</b>	GTVLDSGLIPTPSIIYDPAHHDISSH-----	413

:

SLH1_FU <b>h</b>	LREVGSYTNSVHCAIPPGQANLYDLDVAYSLEAGDYFTSSDKTKCQQG-SRIQITVDDE	491
MecrH1_FU <b>h</b>	MLNLGSYTAMFKCKVPPFSFHAFELGKMYSVESGDYFMTASTTELCNDNNLRIHVHVDDE	495
<b>SLH1_FU<b>a</b></b>	-----	<b>423</b>
MecrH1_FU <b>a</b>	-----	443
SLH1_FU <b>d</b>	-----	417
MecrH1_FU <b>d</b>	-----	414
SLH1_FU <b>f</b>	-----	422
MecrH1_FU <b>f</b>	-----	421
SLH1_FU <b>g</b>	-----	410
Ocdoh1	-----	394
MecrH1_FU <b>g</b>	-----	405
SLH1_FU <b>b</b>	-----	411
MecrH1_FU <b>b</b>	-----	405
SLH1_FU <b>c</b>	-----	418
MecrH1_FU <b>c</b>	-----	418
SLH1_FU <b>e</b>	-----	412
MecrH1_FU <b>e</b>	-----	413

Table S8. Sequence alignment of selected molluscan hemocyanins. Slipper limpet hemocyanin type 1 (*Crepidula fornicata*; CrfoH1; GenBank accession number WCA44164.1) is shown in bold text with the key residues participating in the interaction network (FUa only) shown in Figure 6 highlighted in yellow. Key (sequence accession number and organism): APCA; CAD88977.1; *Aplysia californica*. Coas; AYO86684.1; *Cornu aspersum*. EuscH1; AIL00900.1; *Euprymna scolopes*. HaasH1; XP\_067664054.1; *Haliotis asinina*. HacRH1; XP\_071088435.1; *Haliotis cracherodii*. HadishaH1; AYD59980.1; *Haliotis discus hannai*. HadivH1; ACU00134.1; *Haliotis diversicolor*. HarubH1; ANE23704.1; *Haliotis rubra*. HarufH1; XP\_046359783.2; *Haliotis rufescens*. HeluHaD; AEO51767.1; *Helix lucorum*. HepoHaN; AYO86687.1; *Helix pomatia*. LijaH2; XP\_064608055.1; *Liolophura japonica*. LisaH1; DAZ89886.1; *Littorina saxatilis*. LystH1; AYO86691.1; *Lymnaea stagnalis*. MecrH1; Q10583.2; *Megathura crenulata*. MetuMH; AGX25261.1; *Melanoides tuberculata*. NunuH1; AH10286.1; *Nucula nucleus*. OcvuHA; CAI9741816.1; *Octopus vulgaris*. PhacH1; WGL47534.1; *Physella acuta*. PocaH1; DAZ89889.1; *Pomacea canaliculata*. RaveH1; DAD54775.1; *Rapana venosa*. SeofH1; ABD47515.1; *Sepia officinalis*. SemaH; AGV74427.1; *Sepiella maindroni*. TopaH1; BAS69907.1; *Todarodes pacificus*. H1; type 1 or type 1-like. H2; type 2 or type 2-like. HA; A-type. MH; mega-hemocyanin. HaD; alphaD. HaN; alphaN.

CoasHaN	-----MSQ--LWF--LISLGLLVVWC-QATLIRKKNVDHLSQQDVLNLQKALRDVFD	46
HepoHaN	-----MSQ--LWF--LISLGLLVVWC-QATLLRKNVDHLSQQDVLNLQKALRDVFD	46
HeluHaD	-----MAP--TIV--WLA-FTFMLVCSNALLVRKDVHDLTPEEVLNLQKALREVSK	46
APCAH1	-----MVG--YLG--QALMALLLLALSNAALVRKSDQLTSEEIILNLQKSLREVND	47
LystH1	-----MAL--LWS--FLALALFVLSFGDAALVRKKNVDNLSEEDIINLQKTLRDVAD	47
PhacH1	-----MAR--LLP--LLALALYVLSFSEAVLVRKSVDSLSEEDILNLQKSLRAVAD	47
PocaH1	-----MLD--CRVFLFLFWLLHLQVVLTAQASLVRKDVATLSSEEIILALQKSLRLEV	49
MetuMH	-----MVTS--PRFALALLAFVCQVAFTTGSILRKNVENLSSQEIILSLQTSRLALAA	50
LisaH1	-----MTS--SALTLAFVALLAHVALTSGSLRRKNVASLSETEVLSLQTYLRLQLED	49
<b>CrfoH1</b>	----- <b>MTT--FCGPLALLVLWCQLL</b> ASGSLIRKKNVDSL <b>TESEVLT</b> Q <b>TVLRQLED</b>	<b>49</b>
RaveH1	-----MTS--PRVTGTLFLCLAYLCFATGSLRKNVDTLTQEILRFQNTLLELEK	49
OcvuHA	MAPTRSPWLLGATILCIISI-----FVPVITNGKFVRKKNVDSLTDQEVVSLQVALYSMQK	55
TopaH1	-----NLLRKNVDTLTPEEILNLQVSLRAMQD	27
EuscH1	MR-----RLNHLPLLC LAVILTLWMSGSNVTNGNLVRKKNVDTLSEHEIISLQVALRSMQD	55
SeofH1	MR-----HLNHLPLLYLTVILTFWMSGDLTVYGKILRKNVDTLSHDEIILNLQIALRNMQD	55
SemaH	MR-----HLNHLPLLYLTVILTFWMTGLDVTYGKILRKNVDTLSHDEIILNLQIALRNMQD	55
LijaH2	-----	0
NunuH1	-----MRLPLIFLALG-----IQVCIVHPLLRKDVDSLTTAEVLALQEALKEVQE	46
MecrH1	-----MLSVRLI--IV-----VLALANAENLVRKSVHEHLTQEETLDQAALRELQM	44
HacrH1	-----MGFVPLFL-VA-----LVALAGAENVIRKDASHLTADEIQALHGALHAITD	45
HarufH1	-----MGVVQLFL-VA-----LVALAGAENVIRKDASHLTADEIQALHGALHAVVN	45
HadishaH1	-----MGVVQLFL-VA-----LVALAGAENVIRKDASHLTADEIQALHGALHAVIG	45
HadivH1	-----	0
HaasH1	-----MWLVQFLL-VA-----LAAVAGADNVVRKDVSHLTDDEVQALHGALHDLTE	45
HarubH1	-----MWLVHFL-VA-----LTTGAGADNVVRKDVSHLTDDEVQALHGALHDLTE	45
CoasHaN	DSSSKGFDAI-AAHYGYPPQCKDG--DRDVACCVHGDNLNFPWHRLYTVQIEQALAEKGL-	103
HepoHaN	DNSSKGFDAI-AAHYGYPPQCKDG--DRAVACCVHGDNLNFPWHRLLTVQIEQALHEKGL-	103
HeluHaD	DTSSKGFDAI-AAHYGYPPQCKHG--SKDVACCVHGEPTFPQWHRLYAVQMEQALKEKGL-	103
APCAH1	DTSNLGYAAI-ASYHGYPTQCKDG--DRDIACCVHGMATFPQWHRLYVQMEQALKAKGL-	104
LystH1	DKSAGGYAAI-AAHYGYPAQCKDANRVPVACCVHGMATFPQWHRLYVQMEQALKEKGL-	105
PhacH1	DTSDKGYAAI-AAHYGYPTQCKHG--DKDVACCVHGMATFPQWHRLYVQMEQALKEKGL-	104
PocaH1	DKGVTGFQSL-AAIHGEPSCSNGDNTSVACCVHGMATFPQWHRLYVQMEQALVGEGLV	108
MetuMH	DESASGFKAV-AAHYGEPSCMEDNDGVAIACCVHGMATFPQWHRLYVQFEQALLVKNKGL-	108
LisaH1	DKSETGFQSLVAAHYGEPSCMEDAEGHPVACCVHGMATFPQWHRLYTVQFEQALVQRGLS	109
<b>CrfoH1</b>	<b>DKTATGFQAI-AAHYGEPSCQTKDGTAIACCVHGEATFPQWHRLYVQFEQALVSKGLA</b>	<b>108</b>
RaveH1	DNSDHGFQAL-GAFHGEPSGCQTKEGSAIACCVHGEATFPWHRAYTAQFEQLLVKGLA	108
OcvuHA	DEGPTGFQAI-SAYHGEPADCKAADGSTVVCCLHGMPTFPLWHRLYLVQFEQAMASHGS-	113
TopaH1	DEGASGYQAI-SAYHGEPADCKAADGSTVVCCLHGMPTFPMWHRLYLVQFEQALVAHGS-	85
EuscH1	DDGATGYQAI-SAYHGEPADCKAADGSSIVCCLHGMPTFPLWHRLYLVQFEQALAAHGS-	113
SeofH1	DEGATGYQAI-SAYHGEPADCKAADGSTVVCCLHGMPTFPMWHRLYMVQFEQAVAGHGS-	113
SemaH	DDGATGYQAI-SAYHGEPADCKAADGSTVVCCLHGMPTFPMWHRLYMVQFEQAVAGHGS-	113
LijaH2	-----MEDGTKIACCVHGMATFPWHRLYVVAQIEQSLVKHGS-	37
NunuH1	DTGPTGYQAI-AAHYGYPGDLCAHDGKMACCVHGMPTFPMWHRLLFTTQMEQALRQKGL-	104
MecrH1	DSSSIGFQAI-AAAHGAPA-SCVHKDTSIACCVHGMPTFPHWHRAYVVMERALQTKRR-	101
HacrH1	ASGPLSFREI-TSYHASPP-SCDHHGQKIACCVHGMPTFPWHRAYVVAERALLSQRK-	102

HarufH1	ASGPFGRFRI-SAYHASPP-SCDHNGRKIACCVHGMPTFPFWHRAYVVQAERALLAQRK-	102
HadishaH1	ASGPLSFRI-SAYHASPP-SCDHNGRKIACCVHGMPTFPFWHRAYVVQAERALLSQRK-	102
HadivH1	-----HRAVVVQAERALLSQRK-	17
HaasH1	ATGPMSPFRI-TAYHAAPA-SCDYKGRKIACCVHGMPSFPFWHRAYVVQAERALLSKRK-	102
HarubH1	ATGPMSPVRI-TTYHAAPA-SCEYKGRKIACCVHGMPSFPFWHRAYVVQAERALLSQRK-	102

\*\* .: \* : .

CoasHaN	SIGLPHYWDWTPQLYKLPDLVSNRVFLDSAGGKTRKNAWYQGNIVHG----	159
HepoHaN	SIGLPHYWDWTPQLYKLPDLVSNRVFLDSAGGKTRKNAWYQGYIIVHG----	159
HeluHaD	NIGIPYWEWTHQLDHLPELVSRVFIENDGGKARSNIWYQQIPTP----	159
ApcaH1	SIGLPHYWDWTRPLNHLPELVSRVFIENDGGKARNVVYQGDINID----	160
LystH1	SIGLPHYWDWTRPLTKLPALVSRVFIENDGGKARNVVYQGDIEVVENSKTIVIRHTARAL	165
PhacH1	SIGLPHYWDWTKPLSALPDLVSRVFIENDGGKARNVVYQGDITVD----	160
PocaH1	GLGVPHYWDWTPQLTELPRVQEPFKDPQGGGLGQRNAWFSGDIEVG----	164
MetuMH	NIGVPHYWDWTEPLTHLPSLVEEQIFRDPNGGKARNVVYQGEVFPV----	164
LisaH1	NIGVPHYWDWTEPLSELPLVQVQIFRDPNGGKARNVVYQGEVFPV----	165
<b>CrfoH1</b>	<b>NIGVPHYWDWTKPIKDLPALVKNQIFRDPKGLGQKNAWFSAEIDHD----</b>	<b>164</b>
RaveH1	NLGVPHYWDWTRPLTHLPELVSRVFIENDGGKARNVVYQGEVFPV----	164
OcvuHA	THGVPHYWDWTPQMSLPLVQHPFLIDPSGKAKKNVVFYSGAIAFE----	167
TopaH1	TLGPHYWDWTKPMTQLPELVQHPFLIDPDKKAKKNVVFYSGEIKFE----	139
EuscH1	ALGVPHYWDWTKPMDHLPELVQHPFLIDPDKKAKKNVVFYSGEIKFE----	167
SeofH1	TLGPHYWDWTKPMSHLPELVQHPFLIDPDKKAKKNVVFYSGEIKFE----	167
SemaH	TLGPHYWDWTKPMSHLPELVQHPFLIDPDKKAKKNVVFYSGEIKFE----	167
LijaH2	SIGVPHYWDWTKPIISDLPLVREPLFIPNABGKAQKNPFYEGHIDSE----	91
NunuH1	GIGVPHYWDWTPQMSLPLVQHPFLIDPDKKAKKNVVFYSGEIKFE----	157
MecrH1	TSGLPHYWDWTEPIITQLPSLAADPVYIDSGGKAHTNYWYRGNIDFL----	155
HacrH1	TVGMPYWDWTEPIITQLPALVTEPIFIDSLGGKQPNYWHRGEIAFA----	156
HarufH1	TVGMPYWDWTEPIITQLPALVTEPIFIDSLGGKQPNYWHRGEIAFA----	156
HadishaH1	TVGMPYWDWTEPIITQLPALVTEPIFIDSLGGKQPNYWHRGEIAFA----	156
HadivH1	TVGMPYWDWTEPIITQLPALVTEPIFIDSLGGKQPNYWHRGEIAFA----	71
HaasH1	TVGMPYWDWTPQLTHLPLVTEPIYIDSGGKAQTNFWYRGEIAFI----	156
HarubH1	TVGMPYWDWTPQLTHLPLVTEPIYIDSGGKAQTNFWYRGEIAFI----	156

\* : \* \* : : \* \* : . : : : \* : . : : \* \* \* :

CoasHaN	DKRLYQNVSGDEHTDLFEHVLHAFEYSSYQCFEVQFEVAHSTIHSLVGGRFYPYSVSSLDY	219
HepoHaN	DKRLYQNVSGDEHTDLFEHVLHAFEY-GYQCFEVQFEVAHNTIHSLVGGRYYPYSVSSLDY	218
HeluHaD	DPRLFQQVETGQYTDLFEHVLNALEYPNYQCFEVQFEVAHNTIHYLVGGRHTYSVSHLEY	219
ApcaH1	DDRLFQQVAPGENTQLFEMVLNALEYENYQCFEVQFEVAHNTIHYLVGGRHKYSMSHLEY	220
LystH1	DDRLFQKVEPGQNTDLFEHVLNALEYPNYQCFEVQFEVAHNTIHYLVGGRNKYSMSHLEY	225
PhacH1	DDRLFQKVDAGEHTDLFEHVIDALQYPNYQCFEVQFEVAHNTIHYLVGGRNKYSMSHLEY	220
PocaH1	DERLRFQTVAPGQNSRLDVLHHALEFDDYQCFEVHLEVAHNNIHYLVGGRHEFSMSTLEW	224
MetuMH	DDRLFQKVEPGQNTQLFNIILDALEQDDYQCFEVQFEVAHNNIHYLVGGRHQYSMATLEY	224
LisaH1	DDRLFQKVEPGQNSQLYNIILDALEQDDYQCFEVQFEVAHNNIHYLVGGRHYSMSSTLEY	225
<b>CrfoH1</b>	<b>DDRLFQKVEPGGHTLFLNMVLDALQDDYQCFEQAQFEVAHNNIHYLVGGRHYSMSSTLEY</b>	<b>224</b>
RaveH1	DERLFEHPQDGGKTRLFNMILDALEQDDYQCFEQAQFEVAHNNIHYLVGGRHYSMSSTLEY	224
OcvuHA	DDHLFDAS-HGKNSLLENILMAFEQEEYCRFEVQFEVAHNPVHFLVGGRYTYSMSSTLEY	226
TopaH1	DARLYQASQEQGNFLLEGVLNALEQEDFCHFEVQFEVAHNPVHFLVGGRFTHSMSSLEY	199
EuscH1	DARLYEASQKGAKNFLLEGVLNALEQEDYCHFEVQFEVAHNPVHFLVGGRFTHSMSSLEY	227
SeofH1	DARLYEASKEGKKNFLLEGVLNALEHEDYCHFEVQFEVAHNPVHFLVGGRFTHSMSSLEY	227
SemaH	DARLYEASKDSHKNFLLEGVLNALEHEDYCHFEVQFEVAHNPVHFLVGGRFTHSMSSLEY	227
LijaH2	DERLFEKPDAGKFTPLMEGILLALEQEDFCDFEVQFEVAHNAIHYVMGGPYNFGMSTLDY	151
NunuH1	DDRLFQKVEPGQNTGHYTKLMEVLDALQDDYQCFEQAQFEVAHNDIHYLVGGRYKYSMSSTLEY	217
MecrH1	DDRLFQKVEPGQNTGHYTKLMEVLDALQDDYQCFEQAQFEVAHNDIHYLVGGRYKYSMSSTLEY	215
HacrH1	DDRLFQKVEPGQNTGHYTKLMEVLDALQDDYQCFEQAQFEVAHNDIHYLVGGRYKYSMSSTLEY	216
HarufH1	DDRLFQKVEPGQNTGHYTKLMEVLDALQDDYQCFEQAQFEVAHNDIHYLVGGRYKYSMSSTLEY	216
HadishaH1	DDRLFQKVEPGQNTGHYTKLMEVLDALQDDYQCFEQAQFEVAHNDIHYLVGGRYKYSMSSTLEY	216
HadivH1	DDRLFQKVEPGQNTGHYTKLMEVLDALQDDYQCFEQAQFEVAHNDIHYLVGGRYKYSMSSTLEY	131
HaasH1	DDRLFQKVEPGQNTGHYTKLMEVLDALQDDYQCFEQAQFEVAHNDIHYLVGGRYKYSMSSTLEY	216
HarubH1	DDRLFQKVEPGQNTGHYTKLMEVLDALQDDYQCFEQAQFEVAHNDIHYLVGGRYKYSMSSTLEY	216

\* : \* \* : : \* \* : . : : : \* \* \* : \* \* \* : \* \* \* : \* \* \* : \* \* \* :

CoasHaN	TAYDPLFYLHHSNVDRLFAIYQEVQRIRGIESGSA---NSPICDVKTFYRPLEPFRSRTSN	276
HepoHaN	TGYDPLFYLHHSNVDRLYAIYQEVQKLRGVQSGSA---NSPLCDVKSYYRPLEPFRRKS	275
HeluHaD	TSYDPIFFLHHSNVDKIYIYETIQRSRYTPGPGCTGCELCDIVGFRTPLEPLNRDSN	279
ApcaH1	TSYDPIFFLHHSNVDRIYIYEAALQKQKGYTPGPGCTGKCELCDLKHFDEPLAPFSWETN	275
LystH1	TSYDPIFFLHHSNVDKIYIYEAALQKQKGYTPGPGCTGKCELCDLKHFDEPLAPFSWETN	285
PhacH1	TAYDPIFFLHHSNVDRIYIYEAALQKQKGYTPGPGCTGKCELCDLKHFDEPLAPFSWETN	280
PocaH1	TSYDPIFFLHHSNVDRIYIYEAALQKQKGYTPGPGCTGKCELCDLKHFDEPLAPFSWETN	278
MetuMH	SSYDPIFFLHHSNVDRIYIYEAALQKQKGYTPGPGCTGKCELCDLKHFDEPLAPFSWETN	278
LisaH1	SAYDPIFFLHHSNVDRIYIYEAALQKQKGYTPGPGCTGKCELCDLKHFDEPLAPFSWETN	279
<b>CrfoH1</b>	<b>TSFDPLFFLHHSNVDRIYIYEAALQKQKGYTPGPGCTGKCELCDLKHFDEPLAPFSWETN</b>	<b>278</b>
RaveH1	SSYDPIFFLHHSNVDRIYIYEAALQKQKGYTPGPGCTGKCELCDLKHFDEPLAPFSWETN	278
OcvuHA	TSYDPLFFLHHSNVDRIYIYEAALQKQKGYTPGPGCTGKCELCDLKHFDEPLAPFSWETN	279
TopaH1	TSYDPLFFLHHSNVDRIYIYEAALQKQKGYTPGPGCTGKCELCDLKHFDEPLAPFSWETN	252
EuscH1	TSYDPLFFLHHSNVDRIYIYEAALQKQKGYTPGPGCTGKCELCDLKHFDEPLAPFSWETN	280
SeofH1	TSYDPLFFLHHSNVDRIYIYEAALQKQKGYTPGPGCTGKCELCDLKHFDEPLAPFSWETN	280
SemaH	TSYDPLFFLHHSNVDRIYIYEAALQKQKGYTPGPGCTGKCELCDLKHFDEPLAPFSWETN	280



TopaH1	TDTVLSLGLALDSNYYVTAEVTAINGTLMPTQTI-PRPIVITYIPPOGFKDVMNMVNM----	417
EuschH1	TDKVISLGLPLSGDYVQAVVTAINGTLLPDGII-PSPIVSYVPGHGLDDEMKNV----	445
SeofH1	TDTVNSLGLPLSGNYYVQAIVTAINGTLLPDGII-PNPIVSYVPSVGLKDEEMKVV----	445
SemaH	TDTVTSGLPLSGNYYVQAIVTAINGTSLPDGIL-PNPIVSYVPSAGLKDEEMKVV----	445
LijaH2	TKAVDDANLPYDGNYYVSIKVSAVNGTSIPSSVL-YKQVYVQPAKQFVVDVTPPKG----	370
NunuH1	TDTFHKHGIDFHQNFYIKTELYAVNGSALPSDAL-VSGTVGHQPAGYLDPKPAPV----	435
MecrH1	SDTVHKMGKLDGHYTVKAEFLSVNGTALPDDLL-PHPVVVHHPEKGFDPVVKHH----	433
HacrH1	TKAVQRLGVVLSGHYYVKTLELFSVNGTSLSSDIL-PQPTVAYRPGKGHIDPPIHHR----	434
HarufH1	TKAVQRLGVVLSGHYYVKTLELFSVNGTSLSPDIL-PQPTVAYRPGKGHIDPPVHHR----	434
HadishaH1	TKAVQRLGVALS GHYYVKTLELFSVNGTSLSPDIL-PQPTVAYRPGKGHIDPPVHHR----	434
HadivH1	TKAVQRLGVALS GHYYVKTLELFSVNGTSLSPDIL-PQPTVAYRPGKGHIDPPVHHR----	349
HaasH1	TKAVKRLGVPLSGHYYVKTNLFSVNGTFLSPDLL-PQPTVAYRPGKGHIDPPVHHR----	434
HarubH1	TDAVHRLGVPLSGHYYVKANPFSVNGTSLSPDLV-PQPTVAYRPGKGHIDPPVHHR----	434
	:. . : : **: : *	
CoasHaN	EHEFHENVVRKNVDRLTIEEVVQLREALQEFQNDKSVVEGYQAIAEFHGDPGKCPNPTAK	505
HepoHaN	EHEFHENVAVRKNVDRLTREEVVQLREALQEFQNDKSVVEGYQAIAEFHGDPGKCPDPTAK	504
HeluHaD	-LEHKEEVSIRKDDVHLTREEILELREALEKQLSDHSVDGYQAIAEFHGDPGKCPPTAR	502
ApcH1	GHEVHDGVSIRKDDITLREEVNDLRQALTKFQNDASVDGYQAIAEFHGDPGKCPYPNAK	502
LystH1	EHEYHEGIVSRKDDIRLTNEEIEYELREALQKQFQNDKSVVDGYQAIAEFHGDPGKCPNPSK	513
PhacH1	EHEYHDDVAVRKNVDRLTREEVYELRQALARFQNDSSVDGYQAIAEFHGDPGKCPPTTSK	508
PocaH1	---RTDATVVRKNVKALTDDEETLALRQALSRFQNDTSVDGFQAVAEFHGLPARCPRPDAR	498
MetuMH	-RDVHRHAVRKNVVDLTDDEEVYELRLAMDRFQNDTSVDGYQAIAEFHGLPARCPRPDAA	514
LisaH1	-DGGKPTTVVRKDI AVLTDDEEVYALRQAMARFQNDTSVDGFQAVAEFHGLPAKCPHPDAA	503
<b>CrfoH1</b>	<b>-DSRKPSNVVRKDI ELLTDEEVYALRQAMHRFQNDTSIDGYQAVAEFHGLPAKCPHPDAA</b>	<b>497</b>
RaveH1	-PDLKKTTLVRKDVETLTDDEEVYLRQAMAKFQNDTSIDGFQAVAEFHGLPPKCPHPDAA	509
OcvuHA	---DKADVIRKDVSVLTKEEYGVRIAMERFMDLKSINGYQALAEFHGLPAKCPRPDAP	501
TopaH1	---DTSLLRFRKDISLSTTEEEYELRVAMERFMSDKSINGYQALAEFHGLPAKCPRPDAL	474
EuschH1	---DEMKLSTRKDLSSLTDEEEYELRQAMERFMSDKSINGYQALAEFHGLPAKCPRPDAQ	502
SeofH1	---DVTLLKTRKDVSSLTDEEKYDLRVAMERFMSDKSINGYQALAEFHGLPAKCPRPDAL	502
SemaH	---DATLKLTRKDVSSLTDEEKYDLRVAMERFMSDKSINGYQALAEFHGLPAKCPRPDAL	502
LijaH2	---DPAALAVRKNVDVSRLLTPEVYVNLREAMRRFQADTSVDGFQATVEYHGLPARCPRPDAK	427
NunuH1	---DRTGLAVRKNVDRLTDDEEVDSLRKAMARLQSDKSVGDYDAIAEYHGLPARCPRPDAA	492
MecrH1	---QSANLLVRKNINDLRTREEVNLREAFHKFQEDRSVDGYQATAEYHGLPARCPRPDAK	490
HacrH1	---QNEDYIVRKNINHLTRSETYELRKALERFQADTSVDGFQATVEYHGLPARCPEPDAK	491
HarufH1	---QNEDYIVRKNINHLTRSETYELRKALERFQADTSVDGFQATVEYHGLPARCPEPDAK	491
HadishaH1	---QNEDYIVRKNINHLTRSETYELRKAMERFQADTSVDGFQATVEYHGLPARCPEPDAK	491
HadivH1	---QNEDYIVRKNIDHLTRSETYELRKAMERFQADTSVDGFQATVEYHGLPARCPEPDAK	406
HaasH1	---HDEDLIVRKNIDHLTREEEYELRRALERFQADTSVDGYQATVEYHGLPARCQPDAK	491
HarubH1	---HDDDLIVRKNIDHLTREEEYALRRALERFQADTSVDGYQATVEYHGLPARCQPDAK	491
	**:: ** * : * * : . : * * : * : * . * : * * * : * * * :	
CoasHaN	DRHACCIHGLPTFPHWHRLIVTQVEDSLRRRGAPLGLPYWDFTRPDTHVPQLANDETYFN	565
HepoHaN	DRHACCIHGLPTFPHWHRLIVTQVEDSLRRRGAPLGVVYDFTRPDTHVPLLASEEIIYFN	564
HeluHaD	DRFACCIHGMPTFPHWHRLVQVEDALRRRGAHTGIPYWDWTKPNTHIPALAADETYVN	562
ApcH1	NRLACCIHGMPTFPHWHRLFVQVEDALRRRGAQIGIPYWDWTKPGTQIPALAAEETIYVD	562
LystH1	NRKACCIHGMPTFPHWHRLVQVEDALRRRGSNIGVYWDWTKPRTSVPKLADETYTD	573
PhacH1	VRQACCIHGMPTFPHWHRLVQVEDALRRRGSNIGVYWDWTKPGTQIPALAAADHDYLD	568
PocaH1	VRYACCLHGVPFPHWHRLFVQVEDALRTRGLDIGIPYWDWTKGMDTTVPLADDDVYLD	558
MetuMH	VRYACCIHGMATFPHWHRLFVQVEDELKARDLHFGMPYWDWTKPNAQIPALAAEATYQD	574
LisaH1	LYRACCIHGMATFPHWHRLFVVEVEDELRSRGLFEGIPYWDWTKPNAIPALAADITYTD	563
<b>CrfoH1</b>	<b>VRYACCVHGMATFPHWHRLFVVEVEDELRSRGLFEGIPYWDWTRPNAHIPALANDVSYED</b>	<b>557</b>
RaveH1	VRYACCIHGMATFPHWHRLFVTEVEDELKSRGLEFGIPYWDWTRPNTHPALAAEETIYED	569
OcvuHA	NRIACCEHGMATFPHWHRLVIMQLEDALDRGSTVGVYWDWTKPI THLPSLVAEETIYQD	561
TopaH1	NRVACCIHGMATFPHWHRLVVMQFENALFTRGSPIGVYWDWTKPFTALPSSLADETYVD	534
EuschH1	NRVACCVHGMATFPHWHRLVQVEDALINRGSPIGVYWDWTKPI TALPHLLGDETYVD	562
SeofH1	NRVACCVHGMATFPHWHRLVQVEDALINRGSPIGVYWDWTKPMPKALPDLLADETYVD	562
SemaH	NRAACCVHGMATFPHWHRLVQVEAALINRGSPIGVYWDWTKPMPKALPDLLAQETIYVD	562
LijaH2	ERFACCVHGMATFVHWHRLVQVEDAIRRGSGVGVYWDWTEPIEALPALAGEATYLD	487
NunuH1	ERVACCVHGMPTFPHWHRLVMVQLEDALVDRGSTIGLPHYWDWTKPLIGLPDLISNKMYKD	552
MecrH1	DRYACCVHGMPIFPHWHRLFVTQVEDALVGRGATIGIPYWDWTEPMTHIPGLAGNKTYVD	550
HacrH1	VRFACCMHGMAFPHWHRLFVTQVEDALVRRGSPIGVYWDWTKPMTLHPDLAASANYVD	551
HarufH1	VRFACCMHGMAFPHWHRLFVTQVEDALVRRGSPIGVYWDWTKPMTLHPDLAASENYVD	551
HadishaH1	VRFACCMHGMAFPHWHRLFVTQVEDALVRRGSPIGVYWDWTKPMTLHPDLAASEDYVD	551
HadivH1	VRFACCMHGMAFPHWHRLFVTQVEDALVRRGPPIGVYWDWTKPMTLHPDLAASEDYVD	466
HaasH1	VRFACCMHGMAFPHWHRLFVTQVEDALVRRGSPIGVYWDWTRPMTLHPDLASNETYID	551
HarubH1	VRFACCMHGMAFPHWHRLFVTQVEDALVRRGSPIGVYWDWTKPMTLHPDLASDATYID	551
	* ** * * : * * * * : . : * : * . * : * * * : * : * : . * :	
CoasHaN	PHI-GQNVHNPFFHDVEIAFLGNDVTRTERDI-SESLAQTPAWGEHTELFNAFLLALEQDNF	623
HepoHaN	PHT-GGNLHNPFFHDVQIAFLGNDVHTRTERDI-SPSLAQTPAWGEHTELFNAFLLALEQDYF	622
HeluHaD	PHD-NAEHVNPFFHHAFIGLGGDAKTSRDV-LPELTQTPKWDHTELFDFVFLLALEQDNF	620
ApcH1	PEN-GETVRNPFHAGADVAFNLNSGAKTSRDV-SDSLNDVPEWGDHTNLFQYLYALEQEDF	620
LystH1	PTT-NEVKRNPFFHDAPIAFLGDSTKTTRDV-SSLSSDSPKWDHTELFDAFLLALEQDNF	631
PhacH1	PKG-RVPSQPLPSRPHLSFLGADAKTSRDV-DNSLSQQPKLGDHTELFDAFLLALEQDNF	626
PocaH1	PIT-GEFQHNPFHDAVAFIGE--RTSRQV-QAELTQTPKFGDHTDLFDGMLLAFEQTDF	614
MetuMH	PHHPDVTLHNPFFYDAAVFLNT--RTTRDV-QNDLAEENPIYGDHNSKLYDGIILAFEQTDF	631

LisaH1 PHS-GVEVHNPFFHDAVAFLGE--RTTRDTMVLELSKNAEFGDHTPLFDGILLAFEQTFD 620  
**CrfoH1 ANT-HQQLHNPFHDAPIAFLGE--KTTRQV-QDDLSESPFEGDHTSLFDGILLAFEQTFD 613**  
 RaveH1 PHT-HHQVHNPFFHDALVAFLLK--KTTRDV-QADLTETPAFGDHTALFDGMLLAFEQTFD 625  
 OcvuHA PFT-LESKPNPFFRASIDFLHSGVYVTRDV-DPRLMQPAKGDHCLYDGLLLAYEQDEY 619  
 TopaH1 PYT-KETKPNPFFKAPIEFLLKAGVHTSRQI-DERLQKQPSKGDHGFYLDGLSLAFEQDDF 592  
 EuschH1 PYT-KETKANPFFSSAIEFLDAGVHTTRVI-DPRLQKQPSVGDHDSAMYDGLMLAFEQEDF 620  
 SeofH1 PYT-KETKPNPFFSAPIEFMHAGVSTKRVI-DARLFKEPTVGDHSYLHDGLMLAFEQHDF 620  
 SemaH PYT-KETKPNPFFSAPIEFMHAGISTKRVI-DARLFKEPTVGDHSYLHDGLMLAYEQHDF 620  
 LijaH2 TKT-NERLNPFFFKQIPMVG--SETSRDV-QDGLFEKPAFGEHTSLDEEVMYALEQDHF 543  
 NunuH1 PQT-GANTDNPFFHGSVAFEG--HVTDRDP-DSRLFEDPSYGDHTALFDGMLLALAEQEDF 608  
 MecrH1 SH--GASHTNPFHSSVIAFEENAPHTKRQI-DQRLFKPATFGHHTDLFNQILYAFEQEDY 607  
 HacrH1 PN--GHTRHNPFNNANISFEEGHHHTSRSI-DARLFAPAAYGDHTPLFDGILYAFEQEDF 608  
 HarufH1 PN--GHTRHNPFNNANISFEEGHHHTSRSI-DARLFAPAAYGDHTALFDGILYAFEQEDF 608  
 HadishaH1 PN--GHTRHNPFNNANISFEEGHHHTSRSI-DARLFAPAAYGDHTALFDGILYAFEQEDF 608  
 HadivH1 PN--GHTRHNPFNNANISFEEGHHHTSRSI-DARLFAPAAYGDHTALFDGILYAFEQEDF 523  
 HaasH1 PY--GHIHNPFFNANISFEEGHHHTSRSL-DPRLFAPAAYGDHTHLFDGILYAFEQEDF 608  
 HarubH1 PH--GHIHNPFFNANISFAEGHHHTSRSI-DSRLFAPAAYGDHTHLFDGILYAFEQEDF 608  
 : : \* \* \* \* \* : : \* \* \* :

CoasHaN CKFAVQFEVAHNLIHGLVGGNTSHGLSTLSYSADFPIFYIYHSNIDRIWAIWTALQQLRG 683  
 HepoHaN CKFAIQFEVAHNLIHGLVGGNTPHGLSTLSYSADFPIFYIYHSNIDRIWAIWTALQEHrg 682  
 HeluHaD CDFEVQFEIAHNLIHAYVGGNSKYGLSSLSYSADFPIFYLHHSNIDRIWAIWTALQQRG 680  
 ApcHa1 CNFEVQFEVAHNLIHALVGGNTQFGLSSLSYSADFPIFFLHHSNVDRIWAIWTALQQRG 680  
 LystH1 CDFEVQFEVAHNLIHALVGGKEQYSLATLSYSADFPIFYLHHSNVDRIWAIWTALQELRG 691  
 PhacH1 CDFEQFEVAHNLIHALVGGKPNHSLATLSYSADFPIFYLHHSNVDRIWAIWTALQELRG 686  
 PocaH1 CDFEQFEVHNAPHFLVGGFAPYSLATLHYSADFPIFYLHHSVDRMWAIWQALQVLRG 674  
 MetuMH CDFEQFEVIHNSPHFLVGGFGAYTLATLHYSADFPIFYLHHSNVDRMWAIWQELQIRRG 691  
 LisaH1 CDFEVQFEVHNAPHFLVGGFAPYSLTTLHYSADFPIFYLHHSNVDRLWAIWQKLIIRG 680  
**CrfoH1 CDFEVQFEVHNAPHFLVGGFGAYTLATLHYSADFPIFYLHHSNVDRLWAIWQQLIKRG 673**  
 RaveH1 CDFEVQFEVHNAPHFLVGGFDPYTMATLHYSADPIFYLHHSNVDRLWAIWQKLMRRG 685  
 OcvuHA CDFEVQFEVTHNFVHAWIGGKEAYSMSSLHYTSFDFLFWLHHAQVDRLWAIWQALQIHRR 679  
 TopaH1 CDFEVQFEVTHNSIHAWTGGSEPYSMSSLHYTSFDFPMFWLHHSQVDRLWAIWQALQIRG 652  
 EuschH1 CDFEVQFEVTHNAIHAWTGGAEPEYSMSSLHYTSFDFPMFWHSEVDRLWAIWQALQVHRK 680  
 SeofH1 CDFEVQFEVTHNAIHRWTGGSEPYSMSSLHYTSFDFPMFWLHSEVDRLWAIWQALQIHRG 680  
 SemaH CDFEVQFEVTHNAIHAWTGGSEPYSMSSLHYTSFDFPMFWLHSEVDRLWAIWQALQIHRG 680  
 LijaH2 CDFAVQFEIAHNRIHALVGGSQRYSMSSLHYTSYDPLFFLHHSNVDRLWAIWQQLILRG 603  
 NunuH1 CDFEVQFEVTHNHLHAWVGGSAKYSMSSLHYTAFDPVFFYLHHSNVDRLWAMWQALQQRG 668  
 MecrH1 CDFEVQFEITHNTIHAWTGGSEHFSMSSLHYTAFDPLFYFHHSNVDRLWAIWQALQMRH 667  
 HacrH1 CDFEQFEVLVHNSIHAWIGGSEDSMSTLHYTAFDPIFYLHHSNVDRLWAIWQALQIRRQ 668  
 HarufH1 CDFEQFEVLVHNSIHAWIGGSEDSMSTLHYTAFDPIFYLHHSNVDRLWAIWQALQIRRQ 668  
 HadishaH1 CDFEQFEVLVHNSIHAWIGGSEDSMSTLHYTAFDPIFYLHHSNVDRLWAIWQALQIRRQ 668  
 HadivH1 CDFEQFEVLVHNSIHAWIGGSEDSMSTLHYAADFPIFYLHHSNVDRLWAIWQALQIRRQ 583  
 HaasH1 CDFEQFEVLVHNSIHAWVGAEDYSMATLHYTAFDPIFYLHHSNVDRLWAIWQALQIRRH 668  
 HarubH1 CDFEQFEVLVHNSIHAWIGGSEDSMATLHYTAFDPIFYLHHSNVDRLWAIWQALQIRRH 668  
 \* . \* : \* \* \* : \* \* \* \* \* . : : \* : : : \* : : : \* : : : \* \* \*

CoasHaN KPYKAHCAQSYVHTPLKPFAPKPPYNNDAKTFAHSTATNIYDYEKELAYTFDSLEFGGMS 743  
 HepoHaN KPYKAHCAQSYVHTPLKPFAPKPPYNNDAKTYAHSTATNIYDYEKELAYTYDSLEFGGMS 742  
 HeluHaD KPYKAHCAQSYVHTPLKPFAPHTPYNNNEKTFAHSTPTDVYDYEKEFYQYDNLQFGLG 740  
 ApcHa1 KPYKAHCAQSYVHTPLKPFAPFSSPLNNNEKTFHSHVPTNVYDYAEELGYDYNLQFGLGM 740  
 LystH1 KPYKAHCAQSYTYEPLKPFAPFSSPYNNNDQKTFTHAVPRNVYDYTELNYAYDSLEFGGLT 751  
 PhacH1 KPYKAHCAQSYTYEPLKPFAPFSSPYNNNEKTSHAVPRNVYDYTELKYTYDQLEFGGMS 746  
 PocaH1 LPYQAHCALSRLTRDTLKPFGFPSPPLNNNQKTFSRARGDQVYEQKNFLYDYSLTFGGMT 734  
 MetuMH LPYKAHCAHTMTHEPMKPFAPFSSPLNNNEKTFHSHSEPSQVYDYERDLGYTYDSLTFGGMT 751  
 LisaH1 LPYKAHCASTQTQEVLPKPFPPPPNNDEKTFNNARPTNVYDYENVLDYTYDTLQFGGMT 740  
**CrfoH1 LPYKAHCASLSTQEVLPKPFPPPPNNDEKTFENARPTNIYDYEAVALDYTYDSLQFGGMT 733**  
 RaveH1 KLYKAHCAGSLTQELMPPFGFPFPPYHNDPKTHDNARPSQIYDYESVLDYTYDSLQFGGMT 745  
 OcvuHA KPYKAYCANSEVHRPLKPFAPFSSPLNNNEKTRHSHVPTNVYDYQAEALAYTYDTLFFGGMS 739  
 TopaH1 KPYKAYCANSEVYRPMKPFAPFSSPLNNNEKTRHSHVPTNVYDYQAEALAYTYDTLFFGGMS 712  
 EuschH1 LPYKHCASSEVYRPMKPFAPFSSPLNNNDHQTSHSVPTNIYDYQAEALKYAYDTLFFGGMS 740  
 SeofH1 LPYKAHCASSEVYQPMKPFAPFSSPLNNDELTHSHSVPTDIYDYQSELDYAFDTLFFGGMS 740  
 SemaH LPYKAHCASSEVYQPMKPFAPFSSPLNNDELTHSHSVPTNIYDYQSELDYAFDTLFFGGMS 740  
 LijaH2 KPFFKAYCANSDVHMSLKPFGFPAPLNSNPKTFEHAAPTNIYDYENELDYTYDTLQYHGMT 663  
 NunuH1 KPYKAHCATYTYEPLKPFAPKSPPLNNNEKTKHAPVPTNVYDYSELGYSFDSLEFGGMS 728  
 MecrH1 KPYRAHCAISLEHMLKPFAPFSSPLNNNEKTHANAMPNKIYDYENVLHYTYEDLTFGGIS 727  
 HacrH1 KPYRAHCAQSIEQLPMKPFAPFSSPLNNNEKTHSNVPTDIYDYEETLHYSYDDLTFGGMN 728  
 HarufH1 KPYRAHCAQSIEQLPMKPFAPFSSPLNNNEKTHSNVPTDIYDYEETLHYSYDDLTFGGMN 728  
 HadishaH1 KPYRAHCAQSIEQLPMKPFAPFSSPLNNNEKTHSNVPTDIYDYEETLHYSYDDLTFGGMN 728  
 HadivH1 KPYRAHCAQSIEQLPMKPFAPFSSPLNNNEKTHSNVPTDIYDYEETLHYSYDDLTFGGMN 643  
 HaasH1 KPYQAHCASVEQPMKPFAPFSSPLNNNEKTHSHSVPTDIYDYEELVHYSYDDLTFGGMN 728  
 HarubH1 KPYQAHCASVEQTPMKPFAPFSSPLNNNEKTHSHSIPTDIYDYEELHYSYDDLTFGGMN 728  
 : : \* \* \* : \* \* \* \* \* : : \* : : \* : : \* : : \*

CoasHaN VPELDNYINTNIKKNRNVFVGIQLHGIQTSGLAHYIEAPGKD-KFEAGRFAILGGPTEM 802  
 HepoHaN VPELDNYINTNIKKNRIFVGIQLHGIKTSGLAHIFIEAPGKE-KLEAGRFAILGGPTEM 801  
 HeluHaD IPQLENYINEHLKSKSRFTVGIHLHGKTSGLATHVSASGK--DYVAGHFAILGGPSEM 798  
 ApcHa1 IAELDQYIKSTQKSKDRVFGYILLHGKKAALATVYVKKPGVDTEYNAGQFALLGGPSEM 800

LystH1 IKQLNDYINNERKTKNRVAVGVLWGIKRSALATLYVN-----DVEAGRFBALLGGPSEM 805  
PhacH1 VQQLNDYINNNQTKDRTFAGILLHGIKKSATARIYVTPAGKD-KYEAGRFBVLLGGPSEM 805  
PocaH1 IPQLNQFLA-ERGNDRDTFVGVKHLNIGISAWVKLYISV-PGGERYLANSAFVLLGGEREM 792  
MetuMH IEQLENYLE-ERQSKERVVVGIELHNIGASAWVKVLIDSGSGGPEYLVGKIAVLGGEKEM 810  
LisaH1 VDQLEHYLE-ERQTRDRAFVGIALHNIGVSAWVDVKINA-PGGESYQVGHVAVLGGEKEM 798  
**CrfoH1** **VDQLEHYLH-ERKTQNRDFVGIALHNIGVSAWVTLSLQV-EGGEPYQVGVKIAVLGGEKEM** **791**  
RaveH1 IAQLDHYLE-ERKTHDRTFVGIIMLHNIGISAWATLAIEM-KNGEYTVGKVAVLGGEKEM 803  
OcvuHA IRELQKHID-EKKTHDRVFAGFLFGGIKTSANIDLYVVA--AGNEYLASPISILGGTKEM 796  
TopaH1 IRELQRYVE-EAKSKDRVFAGFLMGIQTSANVDLFFVVA--GGNEFFVGSIAVLGGSKEM 769  
EuschH1 VRELQGHID-NNKHKDRVFAGFLFMGIKTSAIVDFYVVA--AGNEFMAGSVAVLGGSKEM 797  
SeofH1 IRELQEHIE-ENELKDRVFAGFLFMGIKTSANIDFYVVA--GENEFMAGSIAVLGGSKEM 797  
SemaH IRELQEHIE-ENELKDRVFAGFLFMGIKTSANIDFYVVA--GENEFMAGSIAVLGGSKEM 797  
LijaH2 MEQLNIYLR-RRGQKDRVFAGCNLKGIRTSAMVDFQICKQDGTGCGESAGHFAVLGGPFEM 722  
NunuH1 IQELDHYIQ-ERKATDRVFVGFLEGFQKSANVDFFIER-SGQDKFKAGTIAVLGGSDEM 786  
MecrH1 LENIEKMIH-ENQQEDRIYAGFLLAGIRTSANVDFIKT-TDSVQHKAGTFAVLGGSKEM 785  
HacrH1 LDEIEETIH-HRQQHERVFAGFLGGIGKSALVNIFINK-PGSSPHKAGDIAILGGTKEM 786  
HarufH1 LDEIEDTIH-HRQQHERVFAGFLGGIGKSALVNIFINK-PGSSPHKAGDIAILGGTKEM 786  
HadishaH1 LDEIEETIH-HRQQHERVFAGFLGGIGKSALVNIFINK-PGSSPHKAGDIAILGGTKEM 786  
HativH1 LDEIEETIH-HRQQHERVFAGFLGGIGKSALVNIFINK-PGSSPHKAGDIAILGGTKEM 701  
HaasH1 LDQIEKAIH-LRQQHERVFAGFLLAGIGKSALVDIFIKK-PGSEPFKAGDIAILGGSKEM 786  
HarubH1 LDEIEEAIH-LRQQHERVFAGFLGGIGKSALVDIFIKK-TGSEPHKAGDIAILGGAKEM 786  
: : : : \* : \* : . : . : . . : \* \* \*

CoasHaN PWRYDRLYKHEITNALETQLHWAQPYNVNIEIARFDGTPIDAHQFGHVEVLYLPGKQTK 862  
HepoHaN PWRYDNLYKHEITNALEKLDLHWAQPYNVNIEIAQFDGTPIDAKQFGHLEILYLPKQTK 861  
HeluHaD DWYDRPYRHDI SHALEELGVNWAQPFVDVTIEMHTFDNKPIDVSQFPRINIIHKQEQA 858  
ApcH1 EWRFRDRNRYFEITKALEELGINFLSFFDVRLEMNEFDGTPIDVSQFPKSLIFKRRDILP 860  
LystH1 EWRFRDRLYRHEITDTLEELNLHWAQPYVTTLKLTDFEGHDIPIINFPKQIIFKAKEEPK 865  
PhacH1 QWRFRDRLYRQEITNALNELNLQWNDYTVSVEYEAIDNTPISDSELTHIQIYKAKEDVK 865  
PocaH1 TWTFRDLRFKVEITDTLLAIGRNYDEDFDVTLDITDVOQQKLSVDTFSHTTIFHQPKDYD 852  
MetuMH PWHFRDLRFKMEITDALSDDLGSYDDNFDVTLQITDVTGKEWSSDSFQHDTIFHEPGKTFE 870  
LisaH1 AWHFDRLFKLEITDALSALGLRYDDDFEVLHIVDVTGKQWPEDTFSHSTILHSPAQVHE 858  
**CrfoH1** **PWRFRDLRFKVEITSALQKGLSYDDDFNVHLDITDVTGKWAEDTFHHNTIIHVPGEHGE** **851**  
RaveH1 PWHFRDRTFKVEITEALHKLGYRYDDEYDVKLHKIDVTKGKWPEDTFSSHITIIHVPASTHL 863  
OcvuHA PWRFRDLRFQEIITDLSKALGVDPFGDFTLRVDIKDVNGTALPSTIISQPIVIFEPKQVET 856  
TopaH1 TWRFRDRPYKHEITDALSALGLTKEDVFSIHVDITDVTGKALNSNLVPSPIIFSPAAGEE 829  
EuschH1 SWRFDRVYKHEITGALAALGVDKYAEYTLRVDIKDVNGTALPATLTPQPTVIFVPGKADT 857  
SeofH1 SWRFDRVYKYEITAAALGVDKYAEYTLRVDIKDVNGTALSANILPSPPTVIFTPGKVE- 856  
SemaH SWRFDRVYKYEITAAALGVDKYAEYTLRVDIKDVNGTALSANILPSPPTVIFTPGKVE- 856  
LijaH2 DWRYDRPYKHEITKALSALGLTKEDVFSIHVDITDVTGKALNSNLVPSPIIFSPAAGEE 782  
NunuH1 KWRFRDRVYKHEITESLTLGLTKDSDFSISVTLTVDVGTELDASVLDHPTIIFEPSHAAE 846  
MecrH1 KWGFDRVFKFDITHVLKDLDLTADGDVEVTFDITDVTGKCLASSLIPHASVIREHARVKF 845  
HacrH1 PWAFRDLRFKVEITDALKALDVGDDYEVTFDIHDMHGKLSDDLIPHPAVVSEPAHPSF 846  
HarufH1 PWAFRDLRFKVEITDALKALDVGDDYEVTFDIHDMHGKLSDDLIPHPAVVSEPAHPTF 846  
HadishaH1 PWAFRDLRYKVEITDALKALDVGDDYEVTFDIHDMHGKLSDDLIPHPAVVSEPAHPSF 846  
HativH1 PWAFRDLRYKVEITDALKALDVGDDYEVTFDIHDMHGKLSDDLIPHPAVVSEPAHPSF 761  
HaasH1 PWAFRDLRYKVEITDALKNLSLEVDGDDYEVTFDIHDMHGKALDADLI PHAAVISEPAHPSF 846  
HarubH1 PWAFRDRVYKVEITDALKILSLEVDGDDYEVTFDIHDMHGNELDTDLI PHAAVISEPAHPSF 846  
\* : \* . : : \* : \* : . : . : . . : \* \* \*

CoasHaN KSA---DVEVHVRKEVSQLTGEEIIDLRLHALANLEVDKSIGGYQTLGRYHGASKWCPSPS 919  
HepoHaN KPA---DVEVHVRKEVSQLTGEEIIDLRLHALSNLEEDKSIGGYQTLGRYHGAYKWCPS 918  
HeluHaD DTS---PITKIRKNNVHLTEEEIIDLRLHALSLAFLEEDRSVGGYQTLGRFHGTPNWCPS 915  
ApcH1 EAQ---IDVTVRKNVDSLNAEEVLALRLALANLKEEDSISGGYQTLGRYHGTPPLWC 917  
LystH1 DVS---ADTIKIRKNNVATLTAEEVVDLRLQALANLQNDQSSAGGYQDLGRFHGT 922  
PhacH1 DET---PDSHRVRKNVNSLTREEVSLALREAMNLQGDNSAGGYQDLGRFHGT 922  
PocaH1 GDA---PEEVGRMRYNVDSLTRQAQSLRDLALQKLNQDSTVNGFGHIAAFHGQPSWCP 910  
MetuMH VRGV---SDKRTLRLDVLHNSLSEQIQLNRDAFKALEDSDSINGFNQIAAFHGQPSW 927  
LisaH1 HEGMTFGDREYLRSDVTSLSREQIQLNRDAMQSLSDSINGYNQIAAFHGQPNWCPS 918  
**CrfoH1** **D-P---GEDDDHVRNDIVTTLTAEQEQNRDAFQGLKNDKSVAGYNAIAAFHGQPNWCPS** **908**  
RaveH1 EED---HKEEDHVRNDVDTLTKEQIQNMREALATLQDHSPPGFDHLAAAFHGQPNW 921  
OcvuHA DVK---FDENFRTRKEVSTLTASQMAALREALEKEYEADKTANGYQQVAAFHGSTEW 914  
TopaH1 NVK---FDEQHRSRKNVDSMTVSEMNALRTAMAAFAADKEVTGYQQVAAFHGSTQ 887  
EuschH1 DVE---FDEQHRSRKNVDSMTKTEMDSIRSALTAFAADKSAANGFYQQVAAFHG 915  
SeofH1 DVH---FDEQHRSRKNVDSMTKSEMEDLRKAMTAFADKGVTYQQVAAFHGSTK 914  
SemaH EVH---FDEQHRSRKNVDSMTKSEMDDLKAMTTFADKGVTYQVAAFHGSTK 914  
LijaH2 E-F---ADDWTIVRKDVTRLSPEETALRVALQAVKDDSSVGGFQTIAGYHGIPTLC 839  
NunuH1 D-R---PTDFRVVRKNADRLTETETDLSLVHALKQEDTGAGGFQAFHGLPNW 902  
MecrH1 D-K---VPRSRLIRKNVDRLSPEEMNELRKALALLKEDKSAAGGFQQLGAFHGEPK 902  
HacrH1 E-D---VSTSLRIRKNVDYLTPEETNDLRHALDLLEKDPGAGFNQLGAFHGEPK 903  
HarufH1 E-D---VSTSLRIRKNVDYLTPEETNDLRHALDLLEKDPGAGFNQLGAFHGEPK 903  
HadishaH1 E-D---VSTSLRIRKNVDYLTPEETNDLRHALDLLEKDPGAGFNQLGAFHGEPK 903  
HativH1 E-D---VSTSLRIRKNVDYLTPEETNDLRHALDLLEKDPGAGFNQLGAFHGEPK 818  
HaasH1 E-D---ETHSFRIRKNVDSLTPPEETNELRKAMDLLQKDHAGGAGFNQLGAFHGEPK 903  
HarubH1 E-D---EAHSRLRIRKNVDSLTPPEETNELRKALDLLEKDPGAGFNQLGAFHGEPK 903  
\* : \* . : : \* : \* : . : . : . . : \* \* \*

CoasHaN	AEVKKVCCPHGMAIFPHWHRLLVVQFENALTNHGYPYWDWTPQLDPLPALVTAETY	979
HepoHaN	AEVKKVCCPHGMSIFPHWHRLLVVQFENALSSHGYHGAVPYWDWTPQLDPLPALVTAQTY	978
HeluHaD	AEKKVACCPHGPTFPWHHRLLTVAENALRRKHGKGYKALPYWDWTPATSLPAIVPETY	975
ApcaH1	AEKKVACCLHGMPVFPWHHRLLTVAENALRRHGFSGGLPYWDWTRPMTQLPDLVDSAKY	977
LystH1	AEVKKSCCLHGMPTFPHWHRLWTVQAEALRRHGVGAIYWDWTPQVKSLPDLVTSEKY	982
PhacH1	AEVKKSCCLHGMPTFPHWHRLWTVQAEALRRSGLKGAIPYWDWTPQIAALPELVQSEY	982
PocaH1	APVKYACCVHGMVFPWHHRLLTVMENALRRKGFKEGLPYWDWTPMSSLPDFVEEESY	970
MetuMH	AETKYACCLHGMPTFPHWHRLITVQLENALRDKGFESGLPYWDWTPMTELPESFTAETY	987
LisaH1	AEKKYACCAHGMAIFPHWHRLLTVAENALHANGLDGLPYWDWTPMETLPEIVRDETY	978
<b>CrfoH1</b>	<b>AERKYACCTHGMVFPWHHRLLTVAENGLIANGHLSGLPYWDWTPMSTLSPISVLNATY</b>	<b>968</b>
RaveH1	AEHKVACCPHGMPVFPWHHRLLTVAENALIAHGMHSGLPYWDWTPMTALPKFVADATY	981
OcvuHA	AEHKYACCHHGMAATFPWHHRLTLNFENGLRNGYDGGIYWDWLNIPISALPNLVAEEQY	974
TopaH1	AAQKYACCHHGMAATFPWHHRLIALNFENGLRNGWSSGGLPYWDWTRPIDALPALVLEAEY	947
Eusch1	AAQKYACCHHGMAATFPWHHRLVTLNENGLRNGYSSGGVYWDWTRPIDALPALVLEEYQY	975
SeofH1	AAQKYACCHHGMAATFPWHHRLTLNENGLRNGYSSGGIYWDWTRPIEALPALVLEEYQY	974
SemaH	AAQKYACCHHGMAATFPWHHRLTLNENGLRNGYSSGGIYWDWTRPIEALPALVLEEYQY	974
LijaH2	AASKFSCCVHGMAATFAHWHRLYTVQFEQALRKQFTGALPYWDWSKPISSLPALATDDTF	899
NunuH1	-DEKVACCAHGMAVFPWHHRLLVVQFEAGLHKGYPYWDWSSPMDSLPTLANAATY	961
MecrH1	ASKKFACCVHGMVFPWHHRLLTVAENALRRHGYDGALPYWDWTRPLNHLPELADHEKY	962
HacrH1	AEHKVACCVHGMVFPWHHRLLTVAENALRRHGYSGALPYWDWTRPLKLPDLVTPETY	963
HarufH1	AYHKVACCVHGMVFPWHHRLLTVAENALRRHGYSGALPYWDWTRPINKLPDLVTPETY	963
HadishaH1	AEHKVACCVHGMVFPWHHRLLTVAENALRRHGYSGALPYWDWTRPINKLPDLVTPETY	963
HadivH1	AEHKVACCVHGMVFPWHHRLLTVAENALRRHGYSGAPYWDWTRHPIKLPDLVTPETY	878
HaasH1	AVHKVACCVHGMVFPWHHRLLTVAENALRRKHGYPYWDWTRPLSHLNLVTPPEQF	963
HarubH1	AEHKVACCVHGMVFPWHHRLLTVAENALRRKHGYPYWDWTRPVSKLPDLVAHEHF	963

\* \* \* \* \* : : \* . \* \* . . \* \* \* \* \* \* \* \*

CoasHaN	TDPGSKNDVANPFYSAHI--DDANQDTERSFK-PELFLKPKYGDYTAIAKQVLLAFEQED	1036
HepoHaN	PDPASNKDVANPFYSAHI--DDVNQDTERSFR-PELFQKPKYGEYTSIAKQVLLAFEQED	1035
HeluHaD	IDPSNNQETHNPFHDAYI--DDVNQNTVRSVR-SDLYQQPAFGEYTDIAKQVLLALEQDN	1032
ApcaH1	SDPKSGQELDNPFYSGHI--DDANADTVRSVR-DDLFQQPFPGGYTDIAKQVLLALEQDD	1034
LystH1	RDPSSNNEIKNPFYSAHI--TDADQDTRSVR-DDLFEQKPFQNTAIAKQVLLALEQDN	1039
PhacH1	EDGN-GKSVKNPFYSAHI--TDVDKDTTRSVR-DDLFQKPAFGEYTDVAKQVLLALEQED	1038
PocaH1	QNPRTEDIEFPNPFHGTV--EG--QDTRSVR-EDLFKVPYEQMTPVAEQVLLALEQTD	1025
MetuMH	DNPVSGANEANPLHHAHV--EG--HKTTRSVR-GELEFQKPAFGEYTRIGEQLVLAPEQTD	1042
LisaH1	IHPKTGQSTENPLHHAHV--DG--HKTTRVTLREELFEAPAFGKLTIAEKVMLLAPEQTD	1034
<b>CrfoH1</b>	<b>VHPKTGESVANPLYSGQV--DG--HDTTRSVR-QELFEQKPFQGHMTKIAEKVMLLAPEQDN</b>	<b>1023</b>
RaveH1	ENPKTHHTTEPNPWFSGEV--DG--HNTSRMVR-DDLFEQPEFDKMTIAEKVMLLAPEQDN	1036
OcvuHA	TDM-NGETHVNPFFSGSI--DHIGKTTSRAPS-ADLFQKPFGEFTSLTNEVIYALEQED	1030
TopaH1	TDA-NGEAKPNPFFSGAI--DSIGASTSRAPT-EALYKPKDFGKYTHLANEIIYALEQED	1003
Eusch1	TDV-NGEAHPNPFFSGAI--DEAGAVTSRAPS-ANLFKPEFGKYTHLANEIIYALEQED	1031
SeofH1	TDS-NGESHNPFFSGAI--DEAGAVTSRAPS-EHLYKPKDFGKYTELANEVIYALEKED	1030
SemaH	TDS-NGESHNPFFSGAI--DEAGAVTSRAPS-EHLYKPKDFGKYTELANEVIYALEKED	1030
LijaH2	KDPTTGEDKPNPWSATIAA-AGDVATERAPR-DELFDVQPGFQNTNLGKQLLYAFAEQED	957
NunuH1	THPEDGSSIPNPFYSFDVVEGGVSHSTTSRAPR-QELFENPGFGKHTKVAKQVMLLAPEQED	1020
MecrH1	VDPEGDGVEKHNPFWDGHI--DTVDKTTTRSVQ-NKLFQKPEFGHYTSIAKQVLLALEQDN	1019
HacrH1	TDPSDHDQKHNPNWYNGHI--DTVDEYTTTRNVR-EDLYQQPEFGHFTDIAQVLLALEQDD	1020
HarufH1	TDPSDNQDKHNPNWYNGHI--ETVDQYTTTRNVR-EDLYQQPEFGHFTDIAQVLLALEQDD	1020
HadishaH1	TDPSDNQDKHNPNWYNGHI--ETVDQYTTTRNVR-EDLYQQPEFGHFTDIAQVLLALEQDD	1020
HadivH1	TDPSDNQDKHNPNWYNGHI--ETVDQYTTTRNVR-GDLYQQPEFGHSTDIAQVLLALEQDD	935
HaasH1	EDPSDHHVKHNPNWYNGHI--DSVNQTTTRSVR-ADLYQKPEFGHYTDIAQVLLALEQDD	1020
HarubH1	TDPSDHDHEKHNPNWYNGHI--DTVNHNTTRSVR-SDLYQQPEFGHFTDIAQVLLALEQDD	1020

. \* \* \* : \* \* \* : \* : \* : \* : \* : \* : \*

CoasHaN	FCDFEVQFEIAHNFIAHNLVGGSKPYSMASLAYTAYDPLFLHHSNTDRLWAIWQALQKNR	1096
HepoHaN	FCDFEVQFEIAHNFIAHNLVGGSKPYSMASLAYTAYDPIFYLLHHSNTDRLWAIWQALQKNR	1095
HeluHaD	YCDFEVQYEIAHNFIAHALVGGSEVYSMASLLYTAFDPLFLYLLHHSNTDRIWAIWQALQAYR	1092
ApcaH1	FCTFEVQYEIAHNFIAHALVGGSEQFSLASLQYTAFDPIFYLLHHSNTDRIWAIWQALQKYYR	1094
LystH1	FCDFEVQFEIAHNFIAHALVGGNEQFMSASLLYTAFDPLFLYLLHHSNTDRLWAIWQALQKYYR	1099
PhacH1	FCTFEVQFEIAHNFIAHALVGGQEPYSLASLMYTAFDPLFLYLLHHSNTDRIWAIWQALQKYYR	1098
PocaH1	FCSFEVQFEIAHNFIAHALVGDKEEYSMASLRYTAYDPLFLYLLHHSNTDRLWAMWQVLQGYR	1085
MetuMH	YCSFEVQFEVAHNFIAHALVGGNELYSMASLRYTAYDPLFLALHHSNTDRIWAIWQALQKYYR	1102
LisaH1	FCDFEIQFEIAHNFIAHALVGGNEEYSMASLRYTAYDPIFYLLHHSNTDRIWAIWQQLQYER	1094
<b>CrfoH1</b>	<b>FCDFEIQFEIAHNFIAHALVGGNETYSMASLRYTAYDPIFYLLHHSNTDRIWAIWQALQKYYR</b>	<b>1083</b>
RaveH1	FCDFEIQFEIAHNFIAHALVGGNKLYSMASLRYTAYDPLFLYLLHHSNTDRIWAIWQTLQKMR	1096
OcvuHA	FCSFEVQFEIAHNFIAHALVGGTEPYSMSSLEFTAAYDPIFYLLHHSNTDRIWAIWQALQKHR	1090
TopaH1	FCDFEVQYEIAHNFIAHALVGGTEAVSMASLEYSAFDPIFYLLHHSNTDRIWAIWQALQKFR	1063
Eusch1	FCDFEVQYEIAHNFIAHALVGGTEPFSMASLEYSAFDPIFYLLHHSNTDRIWAIWQALQKFR	1091
SeofH1	FCDFDVFQFEIAHNFIAHALVGGTEAFSMSSLEYSAFDPIFYLLHHSNTDRIWAIWQALQKFR	1090
SemaH	FCDFGVQFEIAHNFIAHALVGGTEAFSMSSLEYSAFDPIFYLLHHSNTDRIWAIWQALQKFR	1090
LijaH2	FCDFSVQFEIATHNFIAHALVGGKPPSMASLRYTTFDPIFYLLHSSVDRYWAIWQALQKFR	1017
NunuH1	FCSFETQFEIAHNFIAHALVGGKDEYSMASLRYTAYDPIFYLLHHSNTDRIWAIWQALQKYYR	1080
MecrH1	FCDFEIQFEIAHNFIAHALVGGQPYGMSALRYTAYDPLFLYLLHHSNTDRIWAIWQALQKYYR	1079
HacrH1	FCSFEVQYEISHNFIAHALVGGTTPYGMSALRYTAYDPIFYLLHHSNTDRIWAIWQALQKYYR	1080
HarufH1	FCSFEVQYEISHNFIAHALVGGTTPYGMSALRYTAYDPIFYLLHHSNTDRIWAIWQALQKYYR	1080
HadishaH1	FCSFEVQYEISHNFIAHALVGGTTPYGMSALRYTAYDPIFYLLHHSNTDRIWAIWQALQKYYR	1080
HadivH1	FCSFEVQYEISHNFIAHALVGGTTPYGMSALRYTAYDPIFYLLHHSNTDRIWAIWQALQKYYR	995

HaasH1	FCSFEVQYEISHNFIHALVGGKDPYGMASLRYTAYDPIFFLHHSNTDRIWAIWQSLQKYR	1080
HarubH1	FCSFEVQYEISHNFIHALVGGTDPYGMASLRYTAYDPLFFLHHSNTDRIWAIWQSLQKYR : * * * : * : * * * * * . . : * * : * : * * * * * : * * * * *	1080
CoasHaN	GKSHNSANCATGLLRKPLAPFSLDSVNPDPVTKAHSQPHQVFDYKQSFHYEYDNLEFNG	1156
HepoHaN	GKPHNSANCATGLLRKPLAPFSLDSVNPDPVTKAHSQPYQVFDYKDSFHYDYDNLEFNG	1155
HeluHaD	GKPENANCAIGKLRIPLPFFSLTDPVNPDPVTRHSSPLRVFNKYKDSFQYEDTLDGFSG	1152
ApcHa1	GKPYNTANCAIGQLRNPLSPFSLTSDINPDPVTRHSSVFPQVFDYQTNFHYEYDALEFNG	1154
LystH1	NKPYNTANCAIGQLRKPLSPFSLTSSVNPDPVTRHSLPFQVFDYRNNFHYEYDNLEFNG	1159
PhacH1	GKPYNTANCAIGQLRKPLSPFSLTSDINPDPVTRHSAVFPQVFDYRNNFHYEYDNLEFNG	1158
PocaH1	GLPMFTANCALTSFRQPLQPFPAQPSVNPDPITRENSIPFRVFDYTRNFHYRDNIDFNG	1145
MetuMH	GLPYNTANCAITQLRKPLQPFPAQTSVNPDPVTRDHSVFPQVFDYSSSFHYDYDNLEFNG	1162
LisaH1	GLPYNSANCAIASLRKPLQPFPAQPSVNPDPVTRDHAVPFDVFNVEDSFHYHYDNQFNG	1154
<b>CrfoH1</b>	<b>GKPYNSANCAIAEMRKPLQPFPAQTSVNPDPVTRDHSVFPQVFDYRNNFHYEYDNLEFNG</b>	<b>1143</b>
RaveH1	GKPYNSANCAIAELRKPLQPFPAQTSVNPDPVTRDHSVFPQVFDYKYNFHYRFDNMQFNG	1156
OcvuHA	GNAYNSANCAIEILLRQPMSPFSLSDINPDSQVTRHSSVFPQVFDYKSNFHYEYDSDLVSG	1150
TopaH1	GKAYNSANCAIEILLRQPMSPFSLSDINPDSQVTRHSSVFPQVFDYKSNFHYEYDSDLVSG	1123
Eusch1	GKPYNTANCAIEMLRKPMSPFSLSDINPDSMTREHSSVFPQVFDYKKNFHYEYDSDLVSG	1151
SeofH1	GKPYNTANCAIEMLRKPMSPFSLSDINADAMTREHSSVFPQVFDYKKAHFHYEYDSDLVSG	1150
SemaH	GKPYNTANCAIEMLRKPMSPFSLSDINADAMTREHSSVFPQVFDYKKAHFHYEYDSDLVSG	1150
LijaH2	GKPYNKANCALEQMRGSLMPFNLDGADVNPDLTRDNSAPPFRVFDYENVFYNYDNLEFNG	1077
NunuH1	GKPYNRANCALDSMRAHLKPFAMPSSVNPYPITRDNIPFGAFDYKSNFHYSDNLEFNG	1140
MecrH1	GKPYNVANCAVTSMREPLQPFGLSANINTDHTVKEHSSVFPQVFDYKTNFHYEYDSDLVSG	1139
HacrH1	GKPYNTANCAIASMRKPLQPFGLSDVNPDIITREHAIPFDVFNKYDNFHYKYDSDLVSG	1140
HarufH1	GKPYNTANCAIASMRKPLQPFGLSDVNPDIITREHAIPFDVFNKYDNFHYKYDSDLVSG	1140
HadishaH1	GKPYNTANCAIASMRKPLQPFGLSDVNPDIITREHAIPFDVFNKYDNFHYKYDSDLVSG	1140
HadivH1	GKPYNTANCAIASMRKPLQPFGLSDVNPDIITREHAIPFDVFNKYDNFHYKYDSDLVSG	1055
HaasH1	GKPYNTANCAIASMRKPLQPFGLSDVNPDIITREHAIPFDVFNKYDNFHYGYDSDLVSG	1140
HarubH1	GKPYNTANCAIASMRKPLQPFGLSDVNPDIITREHAIPFDVFNKYDNFHYVYDSDLVSG . * * * * : * : * * * * * : * : * * * * * : * : * * * * *	1140
CoasHaN	LTIPQLERLIQHNGEDRVFAGFLLHGIQHSALLRFNICKTEDHCEDEGGEFYILGDPHE	1216
HepoHaN	LTIPQLERVIQHNAEDRVFAGFLLHGIHESALLRFNICKTEDHCEPQGGEFYILGDPHE	1215
HeluHaD	LGIPLQAKLLEENKADDRVAGFLLHGVGHSALVTFICRNDTDCNHNGGEFYILGDPNE	1212
ApcHa1	LSPVQLARVLEQNKAGDRVAGFMLHGIQSSALVKFYICKSDDCCNNYAEFFYVLDGVNE	1214
LystH1	LSIPQLAKELEKNKGDVRFVGFMLHGIKRSALVKFYICKSDDCCNNYAEFFYVLDGVNE	1218
PhacH1	LSIPQLAKQLEEIKGKDRVAGFMLHGLKQSALVTFISICKSENSCKK-AGEFYILGDINE	1217
PocaH1	LSIPQLQREIVKQRSRERVFVGFMLHGIQSSALVVLNLCQV--TCQK-AGEFYILGDANE	1202
MetuMH	LSIAQLQHEVVVRKAHERVFAGFMLHGIKQSALVRFEICSSGGACGQ-AGEFYILGDDYE	1221
LisaH1	LSIPQIQREVVRRQAKERI FAGFMLHGIKQSALVVFIEICKPNDGCKE-AGEFYLLGDENE	1213
<b>CrfoH1</b>	<b>MSVAQLQREVIRRRGLERAFAGFMLHGVKKSCLLVFDICKPDGTCTK-SGEFYLLGDENE</b>	<b>1202</b>
RaveH1	MSPQLLRETVRRKGLDRVAGFMLHGIKQSALVVFIEICKPDGTCKE-AGEFYLLGDEFE	1215
OcvuHA	MSIPQLQTEINKRRAKDRIFVTFLELGHKQSLVVEYIRLHGSTDNRNKAGEFYVLSGENE	1210
TopaH1	LSIXQLSREINRRKAKNRVXVTFMLEGLKKSLLVEYFIAADGTDQKMKAGEFYVLSGENE	1183
Eusch1	LSIPQLSREINRRKAKNRVXVTFMLEGLKKSLLVEYFIAADGTDQKMKAGEFYVLSGENE	1211
SeofH1	MSIPQLSREINRRKSKNRVITFMLEGLKKSLLVEYFIAADGTDQKMKAGEFYVLSGENE	1210
SemaH	MSIPQLSREINRRKSKNRVITFMLEGLKKSLLVEYFIAADGTDQKMKAGEFYVLSGENE	1210
LijaH2	LSISDLSHEIENQKSHDRVFAGFMLHGIKRSALVKFYICKSDDCCNNYAEFFYVLDGVNE	1137
NunuH1	LSIPQLSRELESRSKSLYRVFAGFMLHGIKDTVLVKFYICLSNGDCSNYAGEFLLGDPAE	1200
MecrH1	LSISQLNKKLEAIKSDRFFAGFLLSGFKKSLVKFNICTDSSNCH-PAGEFYLLGDENE	1198
HacrH1	LSISQLNRELDRIKSHDRVFAGFLLSGIKKSVLVKFNVCAPPDCH-PAGEFYLLGDENE	1199
HarufH1	LSISQLNRELDRIKSHDRVFAGFLLSGIKKSVLVKFNVCAPPDCH-PAGEFYLLGDENE	1199
HadishaH1	LSISQLNRELDRIKSHDRVFAGFLLSGIKKSVLVKFNVCAPPDCH-PAGEFYLLGDENE	1199
HadivH1	LSISQLNRELDRIKSHDRVFAGFLLSGIKKSVLVKFNVCAPPDCH-PAGEFYLLGDENE	1114
HaasH1	LSISQLNRELEKIKSHDRVFAGFLLSGIKKSVLVKFEVCTPPDQCH-KAGEFYLLGDENE	1199
HarubH1	LSISQLNRELDRIKSHDRVFAGFLLSGIKKSVLVKFEVCTPPDCH-KAGEFYLLGDENE : : : * * * * * : * : : * * * * *	1199
CoasHaN	IPWSYNRLFYKYEITEQLKSLGLHYDDSYTIHYTLLDVGADLG--QRYSQPTVIHEVGTS	1274
HepoHaN	IPWSYNRLFYKYEITEQLKSLGLHYDDNYNIHYTLLDVGADLG--QRYSQPTVIHQVGTGTS	1273
HeluHaD	MEWSYDRLYKYEITEELKHLRLRYNDRYFVRYEIHDLTGQDLG--QPFPPTPTVIRQIGTS	1270
ApcHa1	MAWSYDRLYKYEITDALAALGLRYNDRYSVRYEVLNLQGEDIG--QPFPPTPTVVKEDGTS	1272
LystH1	MEWSYDRLYKYEITQLEDFGLRYNDRYYIKYQVLDLGDGSIY--STDTFSTPTVLFELGTS	1277
PhacH1	MEWNYDRLYKYEITHQLEEQNIHNDRYDITVKVQALDGSDI TEAANI PPTVLFELGTS	1277
PocaH1	MPWKYDRLFKYEITRLLAELKLRADDPYRVSYTVYDLNGQELE--DNLFGNVTIVRDTGTG	1261
MetuMH	MPWEYDRLFKYEITDQLKEQGLEPLDRYDIKYTVFALDGSVVG--TDTFGQVTVVHTYGAG	1280
LisaH1	LPWSYDRLFKYEITDQLHEFDLQHGDNIDHYVYVYDLGSSLG--DDLFGSSTTVYSAGHQ	1272
<b>CrfoH1</b>	<b>LPWEYDRLYKYEITHELEDMGLEPQDRFVQYHVYDLDNTDLG--DDVFGKAVI IYSAGQG</b>	<b>1261</b>
RaveH1	MPWEYDRLFKYDITEQLEKFDLEPMDRYDIQYTVSDLNGQNLG--DDLFGNATIVYTPGLG	1274
OcvuHA	MPWKFDREVYKADITRQMEHLHLHYNDLYHIEYTLKDMSGAEVT-----SLG	1256
TopaH1	MPWKFDREVYKSDITYVMDAMKLYHTDKYHVELRITDMTGAEVT--DLKLVTSVIYEPGIG	1241
Eusch1	MPWKFDREVYKADITHTMDEMMLHYTDKYHVEYKISDMTGAEVT--GVSLSSSVVYEPGLG	1269
SeofH1	MPWKFDREVYKSDITHVMDEMMLHYTDKYHVEYKISDMTGAEVT--DIKLESSVYEPGIG	1268
SemaH	MPWKFDREVYKSDITHVMDEMMLHYTDKYHVEYKISDMTGAEVT--DIKLESSVYEPGIG	1268
LijaH2	MPWIYDRAYKYEITSQKALNLRFGDSFSIRSVILDLNGNEIP--PDTFPPSATVDFVPAQG	1196
NunuH1	MPWYDRAYKYEITDVLDSLHLHEDRYSIKYEVYDLNQNQNTG--TQPYPEPTILHEPSSG	1259
MecrH1	MPWAYDRVFKYDITEKLDLKLHAEDHFYIDYEVFDLKPASLG--KDLFKQPSVIHEPRIG	1257

HacrH1 MAWAYDRVFKYDITNVLDENNLHFYDHI F I S Y E V F D L H G T S L G - T D I F H K A N V I H D P G T G 1258  
HarufH1 MAWAYDRVFKYDITNVLDENNLHFYDHI F I S Y E V F D L H G T S L G - T D I F H K A N V I H D S G T G 1258  
HadishaH1 MAWAYDRVFKYDITNVLDENNLHFYDHI F I T Y E V F D L H G T S L G - T D I F H K A N V I H D S G T G 1258  
HativH1 MAWAYDRVFKYDITNVLDENNLHFYDHI F I T Y E V F D L H G T S L G - T D I F H K A N V I H D S G T G 1173  
HaasH1 MAWAYDRVFKYDITNVLDENNLHFYDNI F I K Y E V L D L K G A S L G - T D L F H T A N V I H Y T G T G 1258  
HarubH1 MAWAYDQLFKYDITNVLDENNLHFYDNI F I H Y E V F D L K G T S L G - T D L F H T A N V I H D K G T G 1258

: \* : : : \* : \* : : . \* : : :

CoasHaN HLY--GQEYRPVVAASQVRRNLESLSGEIESLRAAFLAIQND--HSYEGIASYHGKPG 1330  
HepoHaN HIY--GQEYRPVVAASQVRRNLESLSGEIESLRAAFLAIQND--HSYEAIASFHGKPG 1329  
HeluHaD HLY--GREYRDAVTAVSHVRKDLDTLGTAGEIESLRS AFLDIQQD--HTYENIASFHGKPG 1326  
ApcHa1 QLY--GQEYREAVTAASYVRRDLSTLNEGEVESLRAAFLSIQKD--DTYANIAAFHGKPG 1328  
LystH1 NLY--GKEYRFAVSTAASQIRRNLDLSQGEIESLRAAFLSIKED--GTYEKIASFHGKPG 1333  
PhacH1 NLY--GKEYRPSVTTISNIRRNLDLSEGEVESLRS AFLSIQQD--GTYGDIAAFHGKPG 1333  
PocaH1 VRR--GRVFNVIETASHVRKNELEDLSPGETESLRS ALLQMOKD--ESFQKIASFHGFPG 1317  
MetuMH HVE--RKS YG EELKASSHVRNVDLDTGGEIESLKAALQKMEED--GSFENIAEFHGAPG 1336  
LisaH1 D----GNAYRQDVQASSHVRNIDDLTEGEAESLRAALLHMEED--GSFEAIARFHGFPG 1326  
**CrfoH1 HQKGHEEDYLEEVQASSHVRNLEDLTTGAEESLRSALRMEED--GSFDAIARFHGYPG 1319**  
RaveH1 HMKGHEEDYRAEVRASSHVRNKLKSLTTGECESLRSALHMEED--GSFEAIARFHGKPG 1332  
OcvuHA H--YGEDRAWIEPVTASNRIRKNELEDLDTGGEIESLRAAFLAIQND--GTYERIAAFHGLPA 1313  
TopaH1 N--FGEGRRWISPIITSASRIRKNELEDLDTGGEIESLRAAFLAIQND--GRYEEIASFHGVPA 1298  
Eusch1 K--FGEGRAWIEPVTASRIRKNELEDLDTGGEIESLRAAFLAIQND--GRYQEIASFHGLPA 1326  
SeofH1 K--YGEGRAWIEPVTASRIRKNELEDLDTGGEIESLRAAFLAIQND--GSYQIASFHGLPA 1324  
SemaH K--YGEGRAWIEPVTASRIRKNELEDLDTGGEIESLRAAFLAIQND--GRYQIASFHGLPA 1325  
LijaH2 T--YGTAEAWVEPI TDASRIRKNELEDLDTGGEIESLRAAFLAIQND--GRYQIASFHGLPT 1255  
NunuH1 A--GVDHEYEEVVTASHIRKNELEDLDTGGEIESLRAAFLAIQND--GTYAKIAQAHGKPG 1315  
MecrH1 H--HEGEVYQAEVTSANRIRKNELEDLDTGGEIESLRAAFLAIQND--GTYESIAKFAHGSPG 1313  
HacrH1 S--REHDKYVEEVTGASHIRKNELEDLDTGGEIESLRAAFLAIQND--GTYESIAQYHGKPG 1314  
HarufH1 S--REHDKYVEEVTGASHIRKNELEDLDTGGEIESLRAAFLAIQND--GTYESIAQYHGKPG 1314  
HadishaH1 S--REHDKYVEEVTGASHIRKNELEDLDTGGEIESLRAAFLAIQND--GTYESIAQYHGKPG 1314  
HativH1 S--REHDKYVEEVTGASHIRKNELEDLDTGGEIESLRAAFLAIQND--GTYESIAQYHGKPG 1229  
HaasH1 T--RR--ENYVEEVTGASHIRKNELEDLDTGGEIESLRAAFLAIQND--GTYESIAQYHGKPG 1313  
HarubH1 T--REHHTYVEEVTGASHIRKNELEDLDTGGEIESLRAAFLAIQND--GTYESIAQYHGKPG 1314

: : : : : \* \* : : : : : \* \* \* \*

CoasHaN LCEH-QGR-KVACCVHGNPTFFSWHRLYVELVEHALLSHGSSVAVPYWDWISPIKLPKL 1388  
HepoHaN LCEH-EGR-KVACCVHGDPTFFSWHRLYVELVEHALLSHGSSVAVPYWDWISPIKLPKL 1387  
HeluHaD LCQH-EGH-KVACCVHGMPTFFLWHRLYVEQVEEALLDHGSSVAVPYFDWISPIKLPDL 1384  
ApcHa1 LCEL-NGR-KVACCVHGMATFFPAWHRLYVEQVEEALLGRGSSVAVPYWDWTQPI TELPKL 1386  
LystH1 KCKDSAGR-TVACCVHGMPTFFPAWHRLYVEQVEEALLGRGSSVAVPYWDWTAFTELPLD 1392  
PhacH1 LCEH-DGH-KIACCVHGMPTFFPAWHRLYVEQVEEALLSRGSSVAVPYWDWTSPFQKLP 1391  
PocaH1 MCKH-NNR-SVACCVHGMPTFFLWHRLYVEQVENCLLARGSAVSVPYWDWTQPIRQLP 1375  
MetuMH LCEL-NGK-KVACCVHGMATFFPHWHRLYVEQVENALLSHGSSVAVPYWDWTQPIRQLP 1394  
LisaH1 LCEL-NGQ-KKACCVHGMPTFFPHWHRLYVEQVENALLSHGSSVAVPYWDWTQPIRQLP 1384  
**CrfoH1 LCEH-DGH-KAACCVHGMPTFFPHWHRLYVEQVENALLSHGSSVAVPYWDWTQPIKLPAL 1377**  
RaveH1 LCDH-KGK-QVACCVHGMPTFFPHWHRLYVEQVENALLDHGSSVAVPYWDWTDFDHLPLK 1390  
OcvuHA KCPNEDGSKVYTCCLHGMPTFFLWHRLYVEQVENALLDHGSSVAVPYWDWTDFDHLPLK 1373  
TopaH1 QCPSEDGTMVHTCCLHGMPTFFPHWHRLYVEQVENALLDHGSSVAVPYWDWTDFDHLPL 1358  
Eusch1 QCPNADGTVVHTCCLHGMPTFFPHWHRLYVEQVENALLDHGSSVAVPYWDWTDFDHLPL 1386  
SeofH1 QCPSEDGTVVHTCCLHGMPTFFPHWHRLYVEQVENALLDHGSSVAVPYWDWTDFDHLPL 1384  
SemaH QCPSEDGTVVHTCCLHGMPTFFPHWHRLYVEQVENALLDHGSSVAVPYWDWTDFDHLPL 1385  
LijaH2 KCPSPDNP-KYACCVHGMPTFFPHWHRLYVEQVENALLDHGSSVAVPYWDWTDFDHLPL 1314  
NunuH1 LCNG-----KGCCVHGMATFFPHWHRLYVEQVENALLDHGSSVAVPYWDWTDFDHLPL 1369  
MecrH1 LCQLNGNP--IACCVHGMPTFFPHWHRLYVEQVENALLDHGSSVAVPYWDWTDFDHLPL 1371  
HacrH1 KCELNHHS--IACCVHGMPTFFPHWHRLYVEQVENALLDHGSSVAVPYWDWTDFDHLPL 1372  
HarufH1 KCELNHHS--IACCVHGMPTFFPHWHRLYVEQVENALLDHGSSVAVPYWDWTDFDHLPL 1372  
HadishaH1 KCELNHRS--IACCVHGMPTFFPHWHRLYVEQVENALLDHGSSVAVPYWDWTDFDHLPL 1372  
HativH1 KCELNHRS--IACCVHGMPTFFPHWHRLYVEQVENALLDHGSSVAVPYWDWTDFDHLPL 1287  
HaasH1 KCQLNGRN--IACCVHGMPTFFPHWHRLYVEQVENALLDHGSSVAVPYWDWTDFDHLPL 1371  
HarubH1 KCELNHRS--IACCVHGMPTFFPHWHRLYVEQVENALLDHGSSVAVPYWDWTDFDHLPL 1372

\* : : : \* \* \* \* : \* \* \* \* : . \* : : \* : \* :

CoasHaN ISKSTYYNSRQRFDPNPFSSGRIA--GENAVTTRDPQPELFNNDYFLEQALFALEQDHY 1446  
HepoHaN ISKSTYYNSRQRFDPNPFSSGRIA--GENAVTTRDPQPELFNNDYFLEQALFALEQDHY 1445  
HeluHaD ISKATYYNSRQRFDPNPFSSGKVA--GEDAVTTRDPQPELFNNDYFYEQALYALEQDNF 1442  
ApcHa1 INDATYFNSRQRFDPNPFSSGKVL--GEDAVTTRDPQPELFNNDYFYQQLFALEQDNF 1444  
LystH1 INQATYFNSRQRFDPNPFSSGKVA--GEDAVTTRDPQPELFNNDYFYQQLFALEQDNF 1450  
PhacH1 ISKATYYNSRQRFDPNPFSSGKIK--GEDAVTTRDPQPELFNNDYFYEQALYALEQDNF 1449  
PocaH1 ISSPSFFNSRQRFDPNPFSSGKIP--DTDAFTTRDPQPELFNNDYFLDNVLLALEQTSF 1433  
MetuMH IAQETYFNSRQRFDPNPFSSGKIP--DTDAFTTRDPQPELFNNDYFLDNVLLALEQTSF 1454  
LisaH1 IDAPTYFNSRQRFDPNPFSSGKIS--DTDKYTTTRDPQPELFNNDYFLDNVLLALEQTSF 1442  
**CrfoH1 INDATYYNSRQRFDPNPFSSGKIP--DTDKYTTTRDPQPELFNNDYFLDNVLLALEQTSY 1435**  
RaveH1 LAMPTYNSRMSKDVNPFSSGKIS--GTDEYTTTRDPQPELFNNDYFLDNVLLALEQTSY 1448  
OcvuHA INEATYFNSRQRFDPNPFSSGKIS--FENTETTRDPQPELFNNDYFYEHALFALEQDNF 1431  
TopaH1 INEATYFNSRQRFDPNPFSSGKIS--FENAETTRDPQPELFNNDYFLDNVLLALEQTSF 1416  
Eusch1 FNDATYYNSRQRFDPNPFSSGKID--GDAVTTTRDPQPELFNNDYFLDNVLLALEQTSF 1444  
SeofH1 FNDATYYNSRQRFDPNPFSSGKIS--FANTLTTRDPQPELFNNDYFLDNVLLALEQTSF 1442

SemaH	FNDATYYNSRTHLIESNPFRRGSDV--FASTLTDRDAQDVIYNNHEFYDKALLVLEQTFD	1443
LijaH2	VSDSTFFYNSRTRLRLDPNPFRRGEIS--FENAVTTRDPQPELYNNDYFYNQVLLAFEEEDF	1372
NunuH1	LSDDPTFFNSRSHFEDVNPFFSGDIP--GENSDTSRDPQPRLFNNDYFYDNMLLAFQTSF	1427
MecrH1	ISDATYYNSRQHYYETNPFFHGGKIT--HENEITTRDPKDSLHSDYFYEQVLYALAEQDNF	1429
HacrH1	IDDATYFNSRQQRYPNPFRRGKIS--FENAVTTRDPQPEELFNSEYMHNVLALAEQDNY	1430
HarufH1	IDDATYFNSRQQRYPNPFRRGKIS--FENAVTTRDPQPEELFNSEYMHNVLALAEQDDY	1430
HadishaH1	IDDATYFNSRQQRYPNPFRRGKIS--FENAVTTRDPQPEELFNSEYMHNVLALAEQDNY	1430
HadivH1	IDDATYFNSRQQRYPNPFRRGKIS--FENAVTTRDPQPEELFNSEYMHNVLALAEQDNY	1345
HaasH1	IDDATYFNSRQQRYPNPFRRGKIT--FENAVTTRDPQPEELFNSEYMYENVLLALAEQENY	1429
HarubH1	IDDATYFNSRQQRYPNPFRRGRIT--FENAVTTRDPQPEELFNSEYMYENVLLALAEQENY	1430
	. : : : * * * * . : . * * * : : : : * * . : * :	
CoasHaN	CDFEIQFEILHNALHLSWLGGHAFYSMASLDYAAFDPVFFLHHANTDRIWAVWQELQRYRG	1506
HepoHaN	CDFEIQFEILHNALHLSWLGGHAFYSMASLDYAAFDPVFFLHHANTDRIWAVWQELQRYRG	1505
HeluHaD	CDFEIQFEILHNALHLSWLGGHAKYSFSSLDYAFDPVFFLHHANTDRLWAIWQELQRYRG	1502
ApcHa1	CDFEIQFEILHNALHLSWLGGRAEYSFSSLDYAFDPVFFLHHANTDRIWAIWQELQRYRG	1504
LystH1	CDFEIQFEILHNALHLSWLGGRAYSLSSLDYAFDPVFFLHHANTDRIWAIWQELQRYRG	1510
PhacH1	CDFEIQFELVHNALHLSWLGGRAYSLSSLDYAFDPVFFLHHANTDRIWAIWQELQRYRG	1509
PocaH1	CDFEIQFELVTHNALHLSWLGGRDKHSMSSLDYAFDPAFFLHANTDRLWAIWQALQQYRG	1493
MetuMH	CDFEQAQFEVTHNAFHSFLGGRGKYSLSLTDYAFDPVFFLYHANTDRIFAIWQALQSYRG	1514
LisaH1	CDFEVQFEIVHNALHLSWLGGRGKFSMSGLDYAFDPVFFLHHANTDRIWAIWQALQRHRG	1502
<b>CrfoH1</b>	<b>CDFEVQFEILHNALHLSWLGGRGKYSLSLTDYAFDPVFFLHHANTDRIWAIWQALQKIRG</b>	<b>1495</b>
RaveH1	CDFEVQFEIVSHNAIHSFLGGRGKVMSTLDYAFDPIFFLHHANTDRIWAIWQALQKHRD	1508
OcvuHA	CEFATHLEVLHNSLHLSWLGGRDPHSMSSLDYAAAYDPAFFLHNSNDRLWAIWQELQRYRK	1491
TopaH1	CEFEVHYEVLHNTIHSWLGGRDVHSMSSLDYAAAYDPVFFLHNSNDRLWAIWQELQRYRK	1476
Eusch1	CEFEVQLEVLHNRHLSWLGGRVYSMASLDFAAAYDPVFFLHNSNDRLWAIWQELQRYRK	1504
SeofH1	CEFEVQLEVLHNRHLSWLGGRVYSMASLDFAAAYDPVFFLHNSNDRLWAIWQELQRYRK	1502
SemaH	CEFEVQLEVLHNRHLSWLGGRVYSMASLDFAAAYDPVFFLHNSNDRLWAIWQELQRYRK	1503
LijaH2	CNFEVQFEIAHNALHAWIGGRDPYMSLTDYATAYDPIFFLHNTDRLWAIWQELQRYRK	1432
NunuH1	CDFEIQMELVHNALHLSWIGGRAHYSLSLTDYAFDPAFFLHANTDRLWAIWQELQRYRK	1487
MecrH1	CDFEIQLEILHNALHLSWLGGRGKYSMSNLDYAAFDPVFFLHHANTDRIWAIWQDLQRFRK	1489
HacrH1	CDFEIQFELHNALHLSWLGGRGQYSMSLTDYAFDPAFFLHHANTDRLWAIWQELQRYRG	1490
HarufH1	CDFEIQFELVHNALHLSWLGGRGQYSMSLTDYAFDPAFFLHHANTDRLWAIWQELQRYRG	1490
HadishaH1	CDFEIQFELVHNALHLSWLGGRGQYSMSLEYSAFDPAFFLHHANTDRLWAIWQELQRYRG	1490
HadivH1	CDFEIQFELVHNALHLSWLGGRGQYSMSLTDYAFDPAFFLHHANTDRLWAIWQELQRYRG	1405
HaasH1	CDFEIQFELVHNALHLSWLGGRGQYSLSLTDYAFDPAFFLHHANTDRLWAIWQELQRYRE	1489
HarubH1	CDFEIQFELVHNALHLSWLGGRGQYSMSLTDYAFDPAFFLHHANTDRLWAIWQELQRYRD	1490
	* : * : * * : * : * : * : * : * : * : * : * : * : * : * : * : * : * : * : *	
CoasHaN	LQYNEADCAINLMRKPPEFNRV--TNTDEVTRKNSRPVDTFDYRNNLHYEYDTLEFNHL	1564
HepoHaN	LQYNEADCAINLMKPKLPFNRT--TNTDEVTRKNSRPVDTFDYRNNLHYEYDTLEFNHL	1563
HeluHaD	LPYNEADCAINLMRKPPLQPFQDKK--LNPINI TNYSRPADTFDCRNHFHYEYDTLELNLHQ	1561
ApcHa1	LPYNEADCAINLMRKPPLQPFNDD--FNHG DATNRSRPADTFDYRNHFQY EYDTLTFNHM	1563
LystH1	LKYNADCAINQMRTPLRPFNDQ--LNHGDLTHRSRPADTFDYRNHFHYEYDTLEFNHM	1569
PhacH1	LLYNEADCAINLMRKPPLQPFNDGG--LNHG DYTHRSRPADTFDYRNHFHYEYDNLEFNHM	1568
PocaH1	LPSHESDCSLHLMRRPLQPFDDSTS--QNPVEITNQFSRPADVFDTNHFSYRYDNLLFNSW	1552
MetuMH	LHWDEADCALNHMRDGLHPFDVKA--QNEFDITHKYSRPVDVWDYSDHLQYHYDNLLNDW	1573
LisaH1	LSFEESDCALNLMRKLHPFDKKT--QNEFDLTHYSRPADLFDYSGHLDYHYDNLFNSW	1561
<b>CrfoH1</b>	<b>LPFDESDCALNIMGTPLHPFDKKE--ENQFDLTKNYSRPIDAFDYSNHFDYHYDTLKFNSW</b>	<b>1554</b>
RaveH1	LPFDETDCAVNLMSKPLHPFDNKD--INHFG LTYKHSRPADVWDYSRHLDYHYDNLQLHGW	1567
OcvuHA	LPYNSANCALGLLNQPMRPFNSKT--ANSNLLTFTHSRANDVFQYQNVLHYKYDLSLAFNGL	1550
TopaH1	LSYNEANCALPLMNQPMRPFNSST--ANNDRLTFTNSRPNDVFQYQNVLHYKYDTLNFAGL	1535
Eusch1	LPYNEANCALPLLNQPMRPFNSST--ANQDRLTFTNSRPNDVFQYQNVLHYKYDTLTFEGL	1563
SeofH1	LSYNEANCALPLLNQPMRPFNSKT--ANHDRLTFTNSRPNDVFQYQNVLHYKYDTLKFEG	1561
SemaH	LSYNEANCALPLNQPMRPFNSKT--ANHDRLTFTNSRPNDVFQYQNVLHYKYDTLEFEG	1562
LijaH2	KPNTAFKALPYMQTPIKPFSFDRSINYDWMTRVSRPSEAFDYQNNFGYKYDTLEFNMG	1492
NunuH1	LPYNKADCSINLMRKPPLPFAGD--ENKDTLTKKNSHPQDVFDYRNNLHYKYDNLEFHMM	1545
MecrH1	RPYREANCAIQLMHTPLQPFDKSD--NNDEATKTHATPHDGFYQNSFGYAYDNLELNHY	1547
HacrH1	LPYEEANCAINLMHQPLKPFSDPN--ENHDNITLHESKQPQCFDYRNNFGYKYDNLEFHHL	1549
HarufH1	LPYEEANCAINLMHQPLMPFSDPH--ENHDNVTLKHKSKPQCFDYRNNFGYKYDNLEFHHL	1549
HadishaH1	LPYEEANCAINLMHQPLMPFSDPN--ENHDNVTLKHKSKPQCFDYRNNFGYKYDNLEFHHL	1549
HadivH1	LPYEEANCAINLMHQPLMPFSDPN--ENHGNVTLKHKSKPQCFDYRNNFGYKYDNLEFHHL	1464
HaasH1	LPYEEANCAINLMHQPLKPFCDPH--ENHDNITLKYKSKPQCFDYQNHFGYKYDNLEFHHL	1548
HarubH1	LPYEEANCAINLMHQPLKPFSDPH--ENHDNITLKYKSKPQCFDYQNHFGYKYDNLEFHHL	1549
	: * : : : * * * * : * * * : * * * : * * * : * * * : * * * : * * * :	
CoasHaN	SIPQLEELIQS--RKRNGRVFAGFLIHNIGLSADVDVYVCSVSGKYGEQDCDHKAATFSVL	1623
HepoHaN	SIPQLEELIQS--RKRNGRVFAGFLIHNIGLSADVDVYVCVTVGQFGEQDCNKHAGTFSVL	1622
HeluHaD	TVPQLENLLKR--RQYGRV FAGFLIHNIGLSADVTYVVCVPSGPKGKNDCNKHAGVFSVL	1620
ApcHa1	TIPQLENLLHQ--RQESGRV FAGFLHNI GASADVEIYICVPTGPRGKKNCGTRAGVFSVL	1622
LystH1	TVPQLEALLKK--RKESGRV FAGFLHNI GASAVANIYVVCVPSGPRGKKNCHNPAGSFSIL	1628
PhacH1	TIPQLEALLKK--RKQNGRV FAGFLHNI GASAEVDIYVVCV PAGPRGKKNCHNAGSFSIL	1627
PocaH1	TLPQLEAVLRN--QRSRDR LFAGFLLSNIGTSADVLIFICVATGN--GGRSCNHPAGEFSIL	1610
MetuMH	TIPQLEEV LKT--QRSRDR MFAGFLHNI GTSADVELHICVATGN--GDRNCNHPAGKFSIL	1631
LisaH1	TVPQLEEV LKQ--QRSRDR LFAGFLHNI GTSADVEIDVCVATGN--GAQSCNHPAGKFSIL	1619
<b>CrfoH1</b>	<b>TIPQLEQVLEK--QRSRDR VFAGFLHNI GTSADVEIDVCVATGN--GAQSCNHPAGKFSIL</b>	<b>1612</b>
RaveH1	SIAQLEEV LKK--QRSRDR MFAGFLHNI GTSADVEIDVCVDTGD--GGRSCHHPAGKFSIL	1625

OcvuHA	TVIQLENLLQHNHRHQDRVFAAFLHLHGKASADVRIYICAPTGR-NQENCANYAGVFSIL	1609
TopaH1	SIPQLERILQK-NQGRDRIFAGFLLHGIKASADVRIYICVPTGI-GEENCGNYAGIFSVL	1593
Eusch1	SIPLEALLNK-RQDHDRVFAGFLLHGIKTSAEVHIYICVPTGV-NEENCGNFVGTFSVL	1621
SeofH1	SIPLEALLNK-RKSHDRVFAGFLLHGIKTSADISIYICVPTGV-NEENCGNYVGTFSAL	1619
SemaH	SIPLEALLNK-RKSHDRVFAGFLLHGIKTSADISIYICVPTGV-NAENCGNYVGTFSAL	1620
LijaH2	SIPKLSLLEV-RKSQDRVFAGFLLSGIQTSANRIFICLPR-T-GYTDCSNYAGIFSVL	1549
NunuH1	SIPLEATLDA-RKSRDRVFAGFLLHNIIGTSADVEIYICVPGKD-GYKDCNNYAGVFSVL	1603
MecrH1	SIPQLDHMLQE-RKRHDRVAGFLLHNIIGTSADGHVFCVCLPTGE-HTKDCSHEAGMFSIL	1605
HacrH1	SIPSLDATLKA-RKEHDRVAGFLLHNIIGTSADISIYICQPDGS-GGNDCSHKAGTFYVL	1607
HarufH1	SIPSLDATLKA-RKEHDRVAGFLLHNIIGTSADISIYICQPDGS-GGNDCSHKAGTFYVL	1607
HadishaH1	SIPSLDATLKA-RKEHDRVAGFLLHNIIGTSADISIYICLPGDS-GGNDCSHKAGTFYVL	1607
HadivH1	SIPSLDATLKA-RKEHDRVAGFLLHNIIGTSADISIYICLPGDS-GGNDCSHKAGTFYVL	1522
HaasH1	SIPSLDATLKQ-RKQHDRVAGFLLHNIIGTSADITIYICLPNEH-HGHDCGHEAGTFYIL	1606
HarubH1	SIPSLDVTLKQ-RKHDRVAGFLLHNIIGTSADITIYICLPDGR-RGHDCSHEAGTFYLL	1607

:: .\*: : : .\*:\*\*.\*: : : \*\* : : \* . \* . \* \*

CoasHaN	GGET-EMPFEFNRLYKQDITRTVRELGLKLDYANFHLLEVEIKAPNGSQLDPHILPDPSSI	1682
HepoHaN	GGET-EMPFEFNRLYKQDITRTVRELGLKLDNAANFHLLEVEIRAPNGSQLDPHILPDPSSI	1681
HeluHaD	GGEL-EMPTFDRLYMLQITDITIKQLGLKLNNAASYQLKVEIKAVNGTLLDPHILPDPSSI	1679
ApcaH1	GGQLEEMPFVFDRLYKYDITRAVRDLGLRPNDSGAKFDLKNIHAVNGSYLNPSLLPAPT	1682
LystH1	GGEL-EMPFVFDRLYKYDITKTVRDLGLKLDNAANFELRVDIKAVNGSLLDSDILPKPTI	1687
PhacH1	GGEL-EMPFVFDRLYKHDITSTIKALGLKLDNAANFELKIDIKAVNGSLLSADILPTPTI	1686
PocaH1	GGET-EMPTFDRLYKYDISEPIRQLGLKLDAAEFEELEIKIRAFNGSYLEPDLVDKPSI	1669
MetuMH	GGEY-EMPTFDRLYKYDSDSIRKLGKLDGANFDLKVQIHAYNGSYLDPSSLKEPTI	1690
LisaH1	GGEY-EMPTFDRLYKFDISDTVRKLGRLDAAFDLEIKIKSYNGSYIDSNLLGRPSI	1678
<b>CrfoH1</b>	<b>GGEY-EMPTFDRLYKYDSDTVRKLGLRLDAAFDVQIKIFAYNGSYIDASLLHRPTI</b>	<b>1671</b>
RaveH1	GGEY-EMPTFDRLYKHDIIDAIRKLGRLDAAFDLEVKIHSYNGSYIDASLLHPPSI	1684
OcvuHA	GGES-EMPWRFTHLFRYEITDLSMLKGLNHNH--HFRISTEVRAVNGSKIEKKIFDEPTI	1666
TopaH1	GGET-EMPWQFDRLFRYEITDELKGLNQN--HFRVEMELTAVNGSKITQKIFPNPTI	1650
Eusch1	GGES-EMSWNFDRLYRYEITDMLLGLNHNH--HFRIAVEIITANRSMADSNIFPTPTV	1678
SeofH1	GGES-EMSWNFDRLYRYEITDELMLLGLNHNH--HFRISIEIMTMNRSIASSKIFPNPTI	1676
SemaH	GGET-EMSWNFDRLYRYEITDELMLLGLNHNH--HFRISIEIMTMNRSIASSKIFPNPTI	1677
LijaH2	GGET-EMPWSFDRLYKYDITPALSDELGLTADS--NFDVSEIIEIARNGSSLSPLFRPTI	1606
NunuH1	GGEV-EMPFVFDRLYKYDITPTIHSGLNPDASDFDLKVAIHGNGAILSSDALPRPTI	1661
MecrH1	GGQT-EMSFVFDRLYKLDITKALKKNGVHLQ--DFDLEIEITAVNGSHLDSHVIHSPTI	1662
HacrH1	GGET-EMPFVFDRLYKFEITEALHKLGVKLHG-GVFDLELDIVAYNGSHLDSHIF-DPTI	1664
HarufH1	GGET-EMPFVFDRLYKFEITEALHKLGVKLHG-GVFDLELDIVAYNGSHLDSHIF-DPTI	1664
HadishaH1	GGET-EMPFVFDRLYKFEITEALHKLGVKLHG-GVFDLELDIVAYNGSHLDSHIF-DPTI	1664
HadivH1	GGET-EMPFVFDRLYKFEITEALHKLGVKLHG-GVFDLELDIVAYNGSHLDSHIF-DPTI	1579
HaasH1	GGET-EMPFIFDRLYKFEITKALQKLGVLHG-GVFDLELEIRAYNGSYLDPHTF-DPTI	1663
HarubH1	GGET-EMPFIFDRLYKFEITKPLQKLGVLHG-GVFELELEIRAYNGSYLDSHTF-DPTI	1664

\*\* : \*\* : \* : \* : : \* : . : : : \* : . \* : :

CoasHaN	IYIPGTDEEQD-----	1693
HepoHaN	IYIPGTDEVED-----	1692
HeluHaD	IFEPGTKERQA-----	1690
ApcaH1	LYVPGSNERQE-----	1693
LystH1	LFEPGTGERQK-----	1698
PhacH1	QFVPGTTEVQK-----	1697
PocaH1	IFLPGTGKRQT-----	1680
MetuMH	IFEPGENTFQT-----	1701
LisaH1	IFMPGEGKHQD-----	1689
<b>CrfoH1</b>	<b>IFEPGQGQTQD-----</b>	<b>1682</b>
RaveH1	IFQPGEGKTQD-----	1695
OcvuHA	IFVPKHGA-----	1674
TopaH1	IFVPSDVEFEEDTWRDVVTSANRIRRNKDLKSKEDMFSLRAAFKRMTDDGRYEEIAAFHG	1710
Eusch1	IFVPTADSEHKESWGGVITSANRIRRNKDLKKEEMICFADAFRLMAADGRYEEIAAFHG	1738
SeofH1	IFVPGADSGGKDAWGGVITSANRIRRNKDLNKKQMICFAEAFRMAESGRYEEIAAFHG	1736
SemaH	IFVPGADSEGKDAWGGVITSANRIRRNKDLNKKQMICFAEAFRMAESGRYEEIAAFHG	1737
LijaH2	VYVPGTGHEMQS-----	1618
NunuH1	LLQPGRGAAAP-----	1672
MecrH1	LFEAGTDSAHT-----	1673
HacrH1	IFEPGKDTHTVL-----	1675
HarufH1	IFEPGIDHTVL-----	1675
HadishaH1	IFEPGIDHTVL-----	1675
HadivH1	IFEPGIDHTVL-----	1590
HaasH1	IFEPGTDHTIL-----	1674
HarubH1	IFEPGTDHTIL-----	1675

CoasHaN	-----	1693
HepoHaN	-----	1692
HeluHaD	-----	1690
ApcaH1	-----	1693
LystH1	-----	1698
PhacH1	-----	1697
PocaH1	-----	1680

MetuMH	-----	1701
LisaH1	-----	1689
<b>CrfoH1</b>	-----	<b>1682</b>
RaveH1	-----	1695
OcvuHA	-----	1674
TopaH1	LPAQCPNADGSNIHTCCLHGMPVFPWHRLYLALVENELLARGSDVAVPYWDWIEPFDSL	1770
Eusch1	LPAQCPDETGDNVYTCCLHGMPVFPWHRLYLALVENELLARGSCIAPVYWDWIEPFDEL	1798
SeofH1	LPAQCPDESGDKVFTCCLHGMPVFPWHRLYLALVENELMSRGSIAVYWDWIEPFDVL	1796
SemaH	LPAQCPDESGDKVFTCCLHGMPVFPWHRLYLALVENELMARGSCIAPVYWDWIEPFDVL	1797
LijaH2	-----	1618
NunuH1	-----	1672
MecrH1	-----	1673
HacrH1	-----	1675
HarufH1	-----	1675
HadishaH1	-----	1675
HadivH1	-----	1590
HaasH1	-----	1674
HarubH1	-----	1675

CoasHaN	-----	1693
HepoHaN	-----	1692
HeluHaD	-----	1690
ApcaH1	-----	1693
LystH1	-----	1698
PhacH1	-----	1697
PocaH1	-----	1680
MetuMH	-----	1701
LisaH1	-----	1689
<b>CrfoH1</b>	-----	<b>1682</b>
RaveH1	-----	1695
OcvuHA	-----	1674
TopaH1	PGLISDETYKHPKTNEDIENPFHHGKISFADAVTVRKPRDQLFNRRYLYEHALFAFEHTD	1830
Eusch1	PPLINDLTYYNPKTDKTLNPFLLKGSVSFENTHTQRTLSSDIFGNMNYDHALFALEQTD	1858
SeofH1	PVLINDLTYYNPQTNKVHPNPFLLKGNISFASSETHRSPKELYGNRYLYDHALFALEQTD	1856
SemaH	PVLINDLTYYNPKTNKVHPNPFLLKGNISFASTETHRSPTKELFGNRYLYDHALFALEQTD	1857
LijaH2	-----	1618
NunuH1	-----	1672
MecrH1	-----	1673
HacrH1	-----	1675
HarufH1	-----	1675
HadishaH1	-----	1675
HadivH1	-----	1590
HaasH1	-----	1674
HarubH1	-----	1675

CoasHaN	-----	1693
HepoHaN	-----	1692
HeluHaD	-----	1690
ApcaH1	-----	1693
LystH1	-----	1698
PhacH1	-----	1697
PocaH1	-----	1680
MetuMH	-----	1701
LisaH1	-----	1689
<b>CrfoH1</b>	-----	<b>1682</b>
RaveH1	-----	1695
OcvuHA	-----	1674
TopaH1	FCDFEVHFEVLHNSIHSWIGGPNPHSMSSLDFAAYDPIFFLHHSVDRDLWAIWQDLQRYR	1890
Eusch1	FCEFEIQYEVHLNTHALIGGASKYSMSSLDYAAAYDPIFFLHHSNVDRLWAVWQELQRYR	1918
SeofH1	FCEFEVHFEVLHNTIHSWIGGPNPHSMSSLDFAAYDPVFFLHHSNVDRLWAIWQELQRYR	1916
SemaH	FCEFEVHYEVHLNTHIHSWIGGPKHSMSSLDFTAYDPVFFLHHSNVDRLWAIWQELQRYR	1917
LijaH2	-----	1618
NunuH1	-----	1672
MecrH1	-----	1673
HacrH1	-----	1675
HarufH1	-----	1675
HadishaH1	-----	1675
HadivH1	-----	1590
HaasH1	-----	1674
HarubH1	-----	1675

CoasHaN	-----	1693
HepoHaN	-----	1692
HeluHaD	-----	1690

ApcHa1	-----	1693
LystH1	-----	1698
PhacH1	-----	1697
PocaH1	-----	1680
MetuMH	-----	1701
LisaH1	-----	1689
<b>CrfoH1</b>	-----	<b>1682</b>
RaveH1	-----	1695
OcvuHA	-----	1674
TopaH1	KLDYNVANCALNLLNDPMRPFNNTANQDHLTFTNSRPNDVFDYQNSLNYKFDLSLFSGL	1950
Eusch1	KLEYNTATCAKNSLNKPMRPFNSNSTANHDRLTYVNSKPNDFDYQNVLHYKYETIEFSGL	1978
SeofH1	KLDYNTATCAKNYLNKPMRPFNSNSTANHDRLTYVNSKPNDFDYQNVLHYKYDSISFSGL	1976
SemaH	KLDYNTATCAKNYLNKPMRPFNGNSTANHDRLTYVNSKPNDFDYQNVLHYKYDSISFSGL	1977
LijaH2	-----	1618
NunuH1	-----	1672
MecrH1	-----	1673
HacrH1	-----	1675
HarufH1	-----	1675
HadishaH1	-----	1675
HativH1	-----	1590
HaasH1	-----	1674
HarubH1	-----	1675

CoasHaN	-----	1693
HepoHaN	-----	1692
HeluHaD	-----	1690
ApcHa1	-----	1693
LystH1	-----	1698
PhacH1	-----	1697
PocaH1	-----	1680
MetuMH	-----	1701
LisaH1	-----	1689
<b>CrfoH1</b>	-----	<b>1682</b>
RaveH1	-----	1695
OcvuHA	-----	1674
TopaH1	SIPRLDDLLESRQSHDRVFAGFWLSGIKASADVNIHICVPIGVEHEDCDNYAGTFAVLGG	2010
Eusch1	NIPQLENVLKNNKAHDRVFGVSLHGIKASAFFRFYICVPVGA-EKNCDHFGGTFVSVLGG	2037
SeofH1	NIPQLENVLTNNKAHDRVFGVSLHGIKASADVRIYICVPVQEEKNCDHYAGIFSVLGG	2036
SemaH	NIPQLENVLTNNKAHDRVFGVSLHGIKASADVRIYICVPVQEEKNCDHYAGIFSVLGG	2037
LijaH2	-----	1618
NunuH1	-----	1672
MecrH1	-----	1673
HacrH1	-----	1675
HarufH1	-----	1675
HadishaH1	-----	1675
HativH1	-----	1590
HaasH1	-----	1674
HarubH1	-----	1675

CoasHaN	-----	1693
HepoHaN	-----	1692
HeluHaD	-----	1690
ApcHa1	-----	1693
LystH1	-----	1698
PhacH1	-----	1697
PocaH1	-----	1680
MetuMH	-----	1701
LisaH1	-----	1689
<b>CrfoH1</b>	-----	<b>1682</b>
RaveH1	-----	1695
OcvuHA	-----	1674
TopaH1	ETEMPWAFDRLFRYEISDEMKKLQLTEDSKFRLTNI IASNGSKVSNDFPTPTVIFVVK	2070
Eusch1	ETEMPWQFDRLFRYDITPILNELNLNYKSDFTIKTEVIGVNGTKLNEKLFHDPSIIFVVK	2097
SeofH1	ETEMPWQFDRLFRYEISHALNALELTHKSDFTIKVEVTAANGTRINEKIFPEPTIIFAPR	2096
SemaH	ETEMPWQFDRLFRYEISHALNELELTHKSDFTIKVEVIAANGTRINEKIFPEPTIIFAPR	2097
LijaH2	-----	1618
NunuH1	-----	1672
MecrH1	-----	1673
HacrH1	-----	1675
HarufH1	-----	1675
HadishaH1	-----	1675
HativH1	-----	1590
HaasH1	-----	1674
HarubH1	-----	1675

CoasHaN	-----QLGATTTYLVKRNVESLNPLEGYHLADALIALKKDTSADGFQSIATFHAIPP	1745
HepoHaN	-----QHGSTTYLVKRNVESLNPLEGYLLAEALIALKKDTSADGYQSIATFHAIPP	1744
HeluHaD	-----EDGDVKDIIVVRKNVDALSPRETLISLHAEALQADSSADGYQSIAAFHAVPP	1742
ApcaH1	-----ADGQVGNLYLVKRNKIASLSPQEBLSLLKAMAALQADSSADGYQSIASFHAIPP	1745
LystH1	-----SSGEVNTYLVKRIEINSLSPREVQSLYSAMEALQADSSADGWQSIASFHAIPP	1750
PhacH1	-----PSGEVSSYLVRKEINSLSPRETFNSLYKAMESLQADSSADGWQSIAAFHAIPP	1749
PocaH1	-----ANGRTENRLVRKNAWALSPNELRSMWAMRLLQEDSSPNGFQALASFHALPP	1732
MetuMH	-----QDGKTQRNMIRRNVLTLSLAERRSLVLAMRRLQEDHSADGFQALASFHALPE	1753
LisaH1	-----EHGHSERRLVKRDWAISPAERRSLVIALRNLQADSSADGFQSLAAFHAVPP	1741
<b>CrfoH1</b>	<b>-----DVGHVERRLVKRSVLSLSPAERRSLVLAMRSLQEDSSADGFQSIASFHALPP</b>	<b>1734</b>
RaveH1	-----DEGHPISDLVRKSVWTLSPAERRSLVLAMKSLQADSSADGFQSLASFHAQPP	1747
OcvuHA	---HSESEKKLVRANLIRKKNINRSLLEEDSLMHALKRMQKDKSSDGFESIATFHALPP	1731
TopaH1	KERTEQVSTTKSVRGNLVRKNVDRLSLQEIINSLIHALKRMQKDRSSDGFETIASFHALPP	2130
Eusch1	EERSFKSATNKNLYGNLVRKNVDRLSLQEIINSLVHALKRMQKDKQADGFETIASFHAIPP	2157
SeofH1	EEHTVQSATHSNLYGNLVRKNVDRLSLQEVNSLVHALTRMOKDKSADGFETIASFHALPP	2156
SemaH	EEHTVHSATHSNLYGNLVRKNVDRLSLQEVNSLVHALKRMQKDKSADGFETIASFHALLS	2157
LijaH2	-----DSQHSRELERKKNINSLSPGDISSLMTAFRSMADTSADGYQAIASFHAIPA	1670
NunuH1	-----HHG-GSDVTVRKEINTLTPLEVDLSMKAMRRLQEDSSPNGYQAIASFHAVPP	1723
MecrH1	-----DDGHTPEVPMIRKIDITQLDKRQQLSLVKALESMKADHSSDGFQAIASFHALPP	1725
HacrH1	-----DHDNTEEILIRKNINDLSPREVRSLVKALKGMQHDRSADGFQAIASFHALPP	1727
HarufH1	-----DHDNTEEILVRKNINDLSPREVRSLVKALKGMQNDRSADGFQAIASFHALPP	1727
HadishaH1	-----DHDNTEEILVRKNIIDLSPREVRSLVKALKGMQNDRSADGYQAIASFHALPP	1642
HadivH1	-----DHDNTEEILVRKNIIDLSPREVRSLVKALKGMQNDRSADGYQAIASFHALPP	1642
HaasH1	-----DHDHKEEILVRKNINDLSPREKVS LKALEGMKNDRSADGYQAIASFHALPP	1726
HarubH1	-----DHDHKEEILVRKNINDLSPREVRSLVRALEGMKNDRSSDGYQAIASFHALPP	1727

\*.: : : \* : \* \* :\*: :\*:\*\*\*

CoasHaN	LCPSPTASLRYACCIHGGSFQLQWHRLYTVQFEDALKRHGSPIGVPYWDWTRFSTELPRT	1805
HepoHaN	LCPSPTASRRYACCIHGGSFQLQWHRLYTVQFEDALKRHGSPIGVPYWDWTRFSKELPRT	1804
HeluHaD	LCPSPSASTRYACCLHGMSTFFPQWHRLYTVQVEDALRRHGSVVGIPYWDWTRASQSLPHF	1802
ApcaH1	LCPSPSASKRYACCLHGMATFFPQWHRLYTVQVEDALRRHGSVVGIPYWDWTRQTDHLPGL	1805
LystH1	LCPSPSAENRYACCLHGMATFFPQWHRLYTVQVEDALRRHGALVGIPIYWDWSRVSDSLPHF	1810
PhacH1	LCPHPASAKRYACCLHGMATFFPQWHRLYTVQVEDALRRHGSVVGIPYWDWSRLSSSLPHL	1809
PocaH1	LCPYPEAPVRYACCVHGMPTFFPQWHRLYTVQFEDALRRHSALVGIPIYWDWSVEPSGFHPYM	1792
MetuMH	LCPYPEAAKRYACCVHGMATFFPQWHRLYTVQFEEALRRHGSVVGIPYWDWSRQKALPAF	1813
LisaH1	LCPSPEASERFACCVHGQATFFPQWHRLYTVQFEDALRRHGALVGIPIYLNLSLEPVSMPLVL	1801
<b>CrfoH1</b>	<b>LCPYPEATRKFACCIHGGMATFFPQWHRLYTVQFEDALRRHGALVGIPIYWDWTVVNSRLPDF</b>	<b>1794</b>
RaveH1	LCPYPEATRKFACCVHGMATFFPAWHRLYTVQFEDALRRHGALVGIPIYWDWTVVQSELPPF	1807
OcvuHA	MCPNPTAKHRYACCLHGMATFFPQWHRLYTVQFEQALQRHGAKVGVPIYWDWTVPMSEIPSF	1791
TopaH1	LCPNPTAKHRHACCLHGMATFFPQWHRLYTVQFEHSLNRHGAIVGVPIYWDWTVPMTEVPGL	2190
Eusch1	LCPNPTAKHRHACCLHGMATFFPQWHRLYTVQFEQALHRHGATVGVPIYWDWTVPMKEVPNL	2217
SeofH1	LCPNPTAKHRYACCLHGMATFFPQWHRLYTVQFEQALHRHGATVGVPIYWDWTVPMKEVPHL	2216
SemaH	LCPNPTAKHRYACCLHGMATFFPQWHRLYTVQFEQALHRHGATVGVPIYWDWTVPMKEVPHL	2217
LijaH2	LCPNPTAAQRYACCVHGMATFFPQWHRLYTVQVEDALRRQGSTVGLPYWDWLEPFTALPSI	1730
NunuH1	LCPSFAAANRFACCVHGMATFFPQWHRLYTVQFEQALKDHGATVGLPYWDWTRPMTSLPDF	1783
MecrH1	LCPSFAASKRFACCVHGMATFFPQWHRLYTVQFQDSLRRKHGAVVGLPYWDWTLPRSELPEL	1785
PhacH1	LCPNPTAAHRFACCVHGMATFFPQWHRLYTVQFQDALRRHGSVVGIPYWDWTKPVSELPEF	1787
HarufH1	LCPNPTAAHRFACCVHGMATFFPQWHRLYTVQFQDALRRHGSVVGIPYWDWTKPVSELPEF	1787
HadishaH1	LCPNPTAAANRFACCVHGMATFFPQWHRLYTVQFQDALRRHGSVVGIPYWDWTKPVSELPKL	1787
HadivH1	LCPNPTAAANRFACCVHGMATFFPQWHRLYTVQFQDALRRHGSVVGIPYWDWTKPVSELPKL	1702
HaasH1	LCPSFASAKRYACCVHGMATFFPQWHRLYTVQFQDALRRHGSVVGIPYWDWTKPVTELPKL	1786
HarubH1	LCPNPTAAAHRYACCVHGMATFFPQWHRLYTVQFQDALRRHGSFVGVPIYWDWTKPATLPELKL	1787

:\*\* \* \* \*.\*\*\*:\*\* \* :\* \*\*\*: .\*.:.\*. :. : :\*:\*\* :

CoasHaN	FTFTNYSDFPTNVTNLNPFYKGRVEFEHV--DTERDVQEDKLFKSG-PHGWDTWLYTQVL	1862
HepoHaN	FTYANYSDFPTNLWTLNPFYSGRVEFEHV--DTERDVQEDKLFKRG-PHGWDTWLYTQVL	1861
HeluHaD	LSDNNTDPYTKEVDNPNWGHASIDFEHS--HTERDIQSALFKLG-PHGWDTWLFEQAL	1859
ApcaH1	LANPTYTDVYTGQTIIDNPWKAKIEFENS--VTERDVQGDYLFKQG-PHGFDTWLNFQAL	1862
LystH1	IDDENFVDVPTGDQKANPWWKARIEFENS--ETEREVVSDRLFKRG-PHGWDTWLNFQAL	1867
PhacH1	IDDESVIDPNSETVANPWWKAKIAFENS--ETERDVVSDKLFKRG-PHGWDTWLNFQAL	1866
PocaH1	FTNKTYQVPVHKFQWNNPWEAATFAGK--RTARDFQNDRLADS--DGLGGWQWKQFV	1848
MetuMH	ATDEKFTDPVLNVEFDNPNWLGADIEFENS--HTEREPNLARGEEG-EHGYDTWLYEQYL	1870
LisaH1	MSDETWDPLFNSDIPNPWAGADIEFKGA--KVARVDVQSDRLAKKG-PHGYDTWSWKQYL	1858
<b>CrfoH1</b>	<b>IAESVWDDPLFHANFSPNPWAGADIEFDNS--AVVRDVNMDRISQKG-PKGYDTWSWKQYL</b>	<b>1851</b>
RaveH1	FNDEVDWDDPLFHANFNPWQGAIEFNH--KVARDFDMDQLAKKG-PKGYDTWSWKQYI	1864
OcvuHA	LALKEYVNPYTGIEFMFPFNHGHISFISIPETMTTRDISP-HMFE-KPELGKQTWLFNNII	1849
TopaH1	LTSEKYTDPTGIEFTFNPNHGHISFISIPETMTTREVSE-HLFE-QPALGKQTWLFNNII	2248
Eusch1	LTSEKYVDPTGIEFTFNPNHGHISFISIPETMTARDVSD-HLFE-QPGLGKQTWLFNNII	2275
SeofH1	LTSEKYTDPTGIEFTFNPNHGHISFISIPETMTTRDVSE-HLFE-QPALGKQTWLFNNII	2274
SemaH	LTSEKYTDPTGIEFTFNPNHGHISFISIPETMTTRDVNE-HLFE-QPALGKQTWLFNNII	2275
LijaH2	FTDDEFTDSVSLKTFKNPLKNVHIAFEDT--HVERDVIAEKLIE-QPKNGKQTWLYEQVM	1787
NunuH1	FNDGQYTDARTGVTFDNPNFHHADINFEAGVHTRSPNEQRLFE-DPSLYNTWLFENVI	1842
MecrH1	LTVSTIHDPEPGRDIPNPFIGSKIEFEGENVHTKRDINRDLFQ-GSTKTHHWNFFIEQAL	1844
HacrH1	LSAETFHDPHNLNVSNPFAKADIEFEGDVHHTERDIQSDFLFHSQDHEGYHWNFFETVL	1847
HarufH1	LSAETFHDPHNLNVSNPFFKADIEFEGDVHHTERDIQSDFLFHSQDHEGYHWNFFETVL	1847
HadishaH1	LSAETFHDPHNLNVSNPFFKADIEFEGDVHHTERDIQSDFLFHSQDHEGYHWNFFETVL	1847



MecrH1	DTLEFAGWSIRGIDHIVNRQEHRSRVFAGFLLEGFGTSATVDFQVCRTA--GCEDAGYF	2021
HacrH1	DSLELEGRSVAHIDELIKERQEHDRTFAGFLKGFSGTSATVTLKICRDD--HTCQNAGYF	2024
HarufH1	DNLELEGRSVAHIDELIKERQEHDRTFAGFLKGFSGTSATVTLKVCRRDD--HTCQNAGYF	2024
HadishaH1	DNLELEGRSVAHIDELIKERQEHDRTFAGFLKGFSGTSATVTLKICRDD--YTCEAGYF	2024
HadivH1	DNLELEGRSVAHIDELIKERQEHDRTFAGFLKGFSGTSATVTLKICRDD--YTCEAGYF	1939
HaasH1	DNLELEGRSISHIDELIKERQEQDRTFAGFLKGFSGTSATVDFDICTVN--HTCKEAGYF	2023
HarubH1	DNLELEGRSIAHLDELIKERQEQDRTFAGFLKGFSGTSATVDFEICRVD--HTCKEAGYF	2024
	: ** . * . : : . : : . * . * * * . : * . : : . . . *	
CoasHaN	SILGGS AEMPWFQFNRLYKYEITDQLESHHVHYDDEYHLTVHLKS-INGTELD SHLVHEPT	2100
HepoHaN	SILGGS AEMPWFQFNRLYKYEITDQLESHHVHYDDEYHLTVHLKS-INGTELD SHLIEPT	2099
HeluHaD	TVLGG E L E K P W Q F D R L Y R Y P I T D V L K E N H I H Y D D D Y H F H I H I K A - L N G T E L D S K L I P E P S	2096
Apc aH1	TVLGG S A E M P W F Q F D R L F K Y E I T D Q L I A A N R F D D N Y S F N I S V R L - P D G T I L D S S L I P T P S	2098
LystH1	TVLGG S A E M P W F Q F D R L F K Y E I T D Q L N E H K L R F D D E Y T F Q I K I R A - P D G T E L S S D L I G T P S	2103
PhacH1	TILGGS A E M P W F Q F D R L F K Y E I T E Q L E E H G L R F D D E Y T F E I H I K A - P N G S E L A S S L I S A P S	2103
PocaH1	NVLGGS A E M P W R F D R L Y R Y E I T D V L K T K G L K V D D M F Q I K V V I T A - Q N G T V L D S N S L P Q P T	2087
MetuMH	DILGGS A E M P W S F D R A Y K Y D I T N V L E K K G L E A T D A F K V K V A I T A - Q D G T A L D S S L P E P S	2107
LisaH1	NVLGGS A E M P W R F D R L Y K Y E I T D V L E S K G L D V R D T F H I E V M V T A - A N G T V L D S H I L P D A S	2095
<b>CrfoH1</b>	<b>DILGGS A E M P W R F D R L Y K Y E I T D V L E K K G L D V H D S F N I T V S L T A - L D G S A L S S S L I P T P S</b>	<b>2088</b>
RaveH1	DVLGGALETPWRFDRLYKYEISDVLKEKGLEVRDAFHIELTITA-SNGTALPSSLIPEPS	2101
OcvuHA	SILGGS A E M S W A F D R L Y V M E I T D I L H D M G L K F D S H F H I H H H I K T - A N G T D L P S D L I P Q A T	2086
TopaH1	SVLGG S T E M P W A F D R L Y K M E I T D I L Q A M B L K F D S H F T I K T K I V A - H N G T E L P E S L L P E A T	2485
EuscH1	SVLGG S T E M P W S F D R L Y R I E I T D I L K D M G L E F D S H F T I K V Q L A A - H N G T E L N H - V L S E P T	2511
SeofH1	SVLGG S T E M P W A F D R L Y R I E I T D I L K D M G L Q F D S H F T I K V N L L A - H N G T E L T G - V L T E P T	2510
SemaH	SVLGG S T E M P W A F D R L Y R T E I T D I L K D M G L Q F D S H F I I K V N L L A - H N G T E L T G - V L T E P T	2511
LijaH2	HVLGGS M E M P W A F D R Q Y R Y D I T S A V E A L G A R K D D A L S I S V S I V A - T N G T E V P S S I L K T P N	2024
NunuH1	EVLGG T L E M P W R F D R L Y K Y E I T D V L D K M D I R Y D D P F H F E W T V T P T M P G D T I D S N P I S E P S	2080
MecrH1	TVLGG S E K E M P W A F D R L Y K Y D I T E T L D K M N L R H D E I F Q I E V T I T S - Y D G T V L D S G L I P T P S	2080
HacrH1	TVLGG S A E M P W A F D R L Y K Y D I T K T L D H M N L R H E D T F Y I N R T V T A - Y D G T V L P G D L I P P A S	2083
HarufH1	TVLGG S A E M P W A F D R L Y K Y D I T K T L D H M N L R H E D T F Y I N G T V T A - Y D G T V L P G D L I P P A S	2083
HadishaH1	TVLGG S A E M P W A F D R L Y K Y D I T K T L D H M N L R H E D T F Y I N G T V T A - Y D G T V L P G D L I P P A S	2083
HadivH1	TVLGG S A E M P W A F D R L Y K Y D I T K T L D H M N L R H E D T F Y I N G T V T A - Y D G T V L P G D L I P P A S	1998
HaasH1	TILGGS A E M P W F D R L Y K Y D I T K T L N D M H L R H E D T F S V R V N V T S - Y D G T V L S G D L I Q T P S	2082
HarubH1	TVLGG S A E M P W F D R L Y K Y D I T K T L H D M H L R H E D T F S V K V T V T S - Y N G T V L S G D L I Q T P S	2083
	: * * * * * : : : : : : : : : : : : : : : : : .	
CoasHaN	VIYVPAKHVDVTLKKVTVNHIRRNLDDVEDRDTQS IQTALRKLADTGPDGWASLASFHGA	2160
HepoHaN	VIYVPAKHVDVTLKKVTVNHIRRNLDEVEERDTQS IQRALRDLADTGPDGWASLASFHGA	2159
HeluHaD	VLFAAHEDVHHVEATPNNIRRNLMLLEERDIQSIQAALRDLQRDTTNDGWANLASFHGA	2156
Apc aH1	VLFKSAQRDFKREHVAPNHRVRRNLDTLEERDLQSLKAALRDLQHDNSNDGWASLASFHGS	2158
LystH1	VILEPFFKRDYDLEKVTPNFVRHDLSTLTERDLQGLKSALRDLQLDESADGWASLASFHGA	2163
PhacH1	VILDPAHKVTDLEKVTHNYVRRDLSLTERDVQSLKAALRDLQLDPTTNDGWASLASMHGA	2163
PocaH1	VIFMPKIRERDLQDAPPSRVRRDLRDLKEVDIQNLKAAMASFQRDKGGNGWEAITAFHGV	2147
MetuMH	IILQHATRDPQGYELAPNVRVRRSLDLTERDVMSLSKALHDLQEDDSATGWQSLAAYHGV	2167
LisaH1	VIYDPKSRDSQLHEVAPNHRVHDLSHLSELDVMSLKAAMLDMQRDKGTSQYQNIAAFHGA	2155
<b>CrfoH1</b>	<b>VIIFPKSRTELHEVAPNRRHDLTLHLSELDIRSLKSSIRDLQLDSDNDGYQNIASFHGA</b>	<b>2148</b>
RaveH1	VIYAPKSHDTELHEVAPNRRVHDLADLTDRIKLNKSAIRDQNDISKDGYQHIAATFHGA	2161
OcvuHA	IIRIPAMRLEKQRFIPHNRIKRNINSLEERDVQNIIMSALTRMKEDESDFGFQTVASYHGS	2146
TopaH1	IVRIPPSAQNLVAIPLNRIRRNLNSLESDVQNLMSALKRLKEDESDFGFQTIAGYHGS	2545
EuscH1	IVRIPPEKEHIEVAIPLNKIRRQIESLEERDIQNLMSSALRRLKEDESDFGFQTIAGYHGS	2571
SeofH1	VVRIPPEREHIEVAIPLNKIRRNIQTLEERDIQNLMSSLRRLKEDESDFGFQTIASYHGS	2570
SemaH	VVRIPPEREHIEVAIPLNKIRRNIQTLEERDIQNLMSSLRRLKEDESDFGFQTIASYHGS	2571
LijaH2	VFFVPGKDTVRNAQTPPNVRQNIINNVDNERDIQS IMAALAEKKEDESADGFAAIAAFHGS	2084
NunuH1	VIFVPAIVDHNEKFAHNQVRRNINELDEADVRSLSALTDLMADEDKDGQAIARFHGM	2140
MecrH1	I I Y D P A H H D I S S H H L S L N K V R H D L S T L S E R D I G S L K Y A L S S L Q A D T S A D G F A A I A S F H G L	2140
HacrH1	I I F V P G R H T L N S R K H A P N K I R H E L S S L S S R D V A S L K A A L T S L Q K D D G P D G Y Q A I A A F H G L	2143
HarufH1	I I F V P G R H L L N S R K H A P N K I R H E L S S L S S R D V A S L K A A L T S L Q K D D G P D G Y Q A I A A F H G L	2143
HadishaH1	I I F V P G R H N L N S R K H T P N K I R H E L T S L S S R D V A S L K A A L T S L Q K D D G P N G Y Q A I A A F H G L	2143
HadivH1	I I F V P G R H N L N S R K H T P N K I R H E L T S L S S R D V A S L K A A L T S L Q K D D G P N G Y Q A I A A F H G L	2058
HaasH1	I I F V P G R H K L N S R K H A P N K I R H E I S S L S S R D V A S L K A A L T S L Q H D D G P D G Y Q S I A A F H G L	2142
HarubH1	I I F V P G R H K L N S R K H A P N K I R H E L S S L S S R D I A S L K A A L T S L Q H D S G T D G Y Q A T A A F H G V	2143
	: . : * : : : : . * . : : : * * : : * *	
CoasHaN	PARCPDPDHPTVACQHGMPFFLHWHRLF TLQVEQAIQKHGSSIAIPIYWDWTQPLKLPD	2220
HepoHaN	PARCPDPDHPTVACQHGMPFFLHWHRLF TLQVEQAIQKHGSSIAIPIYWDWTQPIKLPD	2219
HeluHaD	PARCPDPKHPTVACQVHGMPFPPHWHRLFALQIEQALHRRHGSSIALPYWDWTLAIDDLPS	2216
Apc aH1	PAKCTPANESVACCIHGMPFPPHWHRLF TLQVEQALRRHGSSIAIPIYWDWTLPITDLPE	2218
LystH1	PSMCENESGAKVACCIHGMPFPPHWHRLF TVQVERALQRHGSSAVIPIYWDWTKPI TALPE	2223
PhacH1	PARCKDDAGKDVACCIHGMPFPPHWHRLF TLQVEQALQRHGSSIAIPIYWDWTKPIKALPE	2223
PocaH1	PARCPSPOKPEKACCIHGMPFPPHWHRLF TLQVDMSSVVRKGSVALPYWDWTLPITDLPS	2207
MetuMH	PALCPSPEEAKYACCIHGMPFPPHWHRLFVLAVEHSLIKHGSSVAIPIYWDWTNPLDKLPE	2227
LisaH1	PAMCPTPEAAEYACCVHGMPFPPHWHRLFVMEDEALIKHGSSVALPYWDWTLP I E H L P D	2215
<b>CrfoH1</b>	<b>PALCPSPEAAEYACCVHGMPFPPHWHRLFVMEDEAMVRHGSSVALPYWDWTMPITQLPD</b>	<b>2208</b>
RaveH1	PALCPDPNAPTHACCLHGNPTFPPHWHRLFVMEDEALIRHGSSVALPYWDWTKPI TELPE	2221
OcvuHA	T-MCPDPENAKYACQHGMPFPPHWHRYALQFEEALRRHGSSVAIPIYWDWTTHSEKLP	2205
TopaH1	L-MCPTPEAPEYACCLHGMPTFMHWHRYLLHFEESSMRRHGSSVAIPIYWDWTMPSDNLP	2604
EuscH1	K-NCPSPDDAQYACCLHGMPTFPPHWHRYLLHFEDAMRRHGSSVAIPIYWDWTLP I S S L P H	2630



RaveH1	ELQKRRGLTYDRADCAVNFQHKMHPFDW--DDLNPVTRTREHAHPQTVFDYEDLGYHYDD	2398
OcvuHA	ELQEYRKMPFDKAYCALDQMSLAMKPF--WESNPNLATRALATPSMLFDYKRFYKYDND	2366
TopaH1	ELQKHRKPLPHDKAYCALDQMAFPMKPF--WESNPNPTTRAVTSPKSLFDYKSLGYDYD	2780
Eusch1	ELQHFRKLPDNKAFKALDQMAFAMKPF--WESNPNPHTRSVAVPAKLFYKSLGYIYDD	2806
SeofH1	ELQHFRKLPDKKAFKALDQMSFQMKPF--WDSNPNNAHTRSSTPAKLFYKGLGYIYDD	2805
SemaH	ELQHFRKLPDKKAFKALDQMSFQMKPF--WDSNPNPHTRSSTPAKLFYKGLGYIYDD	2806
LijaH2	ALQKHRKRPYLKAYCAAQMTMPMKPFKFNETHFNMSVTRSHARPDSVFNENLGYTYDD	2320
NunuH1	ELQKARGMPHNSANCALNLMSEPMHPFD--MGININKVTKRHATPSTVDFDFEDFGYTYDS	2373
MecrH1	ALQEKRGLPSDRADCAVSLMTQNMRFPH--YEINHNQFTKKHAVPNDVFKYELLGYRYDN	2376
HacrH1	ELQSRRLQFNGADCAVSLMGKAMRPF--KDFNHNPFTRKHAVPNTLFDYEDLGYNYDN	2379
HarufH1	ELQSRRLQFNGADCAVSLMGKAMRPF--KDFNHNPFTRKHAVPNTLFDYEDLGYNYDN	2379
HadishaH1	ELQSRRLQFNAADCAVSLMGKAMRPF--KDFNHNPFTRKHAVPNTLFDYEDLGYNYDN	2379
HadivH1	ELQSRRLQFNGADCAVSLMGKAMRPF--KDFNHNPFTRKHAVPNTLFDYEDLGYNYDN	2294
HaasH1	ELQSRRLQFKAADCAVGLMSKAMRPF--KDFNHNPFTRKHAVPNTVFDYEDLGYNYDN	2378
HarubH1	ELQSRRLQYKAADCAVGLMSKAMRPF--KDFNHNPFTRKHAVPNTVFDYEDLGYNYDN	2379
	: * * * * * : : * * : : * : : * * *	

CoasHaN	LNVGGYSLDQLEELIHDNQQHARVFAGFLLHGKITSGSVKIKLQCG---DKCTSAGQFNL	2454
HepoHaN	LKVGGYDLQLEELIHDNQQHARVFAGFLLHGKITSGSVKIKLQCG---DKCTSAGQFNL	2453
HeluHaD	FNIGGYDLDELEKLIHDRQIKPRIFAGFLLKGKITSGSVNLKVKCF---EECTPAGFFNL	2450
ApcaH1	LSIGGYDLQLEQVIADRQSKARVFAGFLLKGVKTSGSVVINICLRN---NVCNYAGRFNV	2453
LystH1	LNIGGYDLQLEQLIAHQSRPRVFAGFLLKGKITSGSVVLNVCKG---SQCTYAGRFLN	2456
PhacH1	LEIGGYDLQLEALIAQRQSRPRVFAGFLLKGKITSGSVVLQICNSR---KQCSYAGRFLN	2458
PocaH1	LNIGDKTPELEKLIHQSRPRVFAGFLLHGKITSADVVISVCVH---SNCVRAGAFFI	2441
MetuMH	YQIGGKNLDELEALIKDNQAHPRVFAGFHLKIGSSADVVKICKDKSGGQCTRAGSIFI	2464
LisaH1	LELGGKTLLELEELIHDQSHPRVFAGFHLHAIGTSADVIFEVCKTD--DECYRAGGFFI	2450
<b>CrfoH1</b>	<b>YQIGGKSIELEELIKKQSHPRVFAGFHLHGLGTSADVELSVCKSQ--NSCVSAGVIFI</b>	<b>2443</b>
RaveH1	FKLLGGNTLELEEMIHEKSHPRVFAGFHLHNIGTSADVVFHICTE--SRCTRAGLFI	2456
OcvuHA	LEFHGMDTEHLEIEIQKQKQKDRVFATFLLHGKITSADVHLKLCDD---KCSAAGVFI	2423
TopaH1	LNPHGMSIQLEALIQKQKADRVFAGFLLHGKITSADVHLKICIE---ADCQEAGVIFV	2837
Eusch1	LTFFHGMSIAQLDAAIQNQKAKDRVFAGFLLHGKITSADVHLKVCNE---DICKDAGVFI	2863
SeofH1	LTFFHGMSIAQLEAAIQRQKEDDRVFAGFLLHGKITSADMHLKVCNE---ADCKEAGTIFI	2862
SemaH	LTFFHGMSIAQLEAAIQRQKEDDRVFAGFLLHGKITSADMHLKVCNE---ADCKEAGTIFI	2863
LijaH2	LKFDGHSIEELSEIVERQTSQSDRVFANFLLHGIGTSADVFKSVCKTD--NTCERAGLFFI	2378
NunuH1	FKIGGLSLSEIQEEDNRRNHDRVFAGFLLSGIKTSALVHFHLCKSD--DDCIKAGEFGV	2431
MecrH1	LEIGGMLHEIEKEIKDKQHVRVFAGFLLHGIGTSADVQFQICKTS--EDCHHGQIFV	2434
HacrH1	LEISGLNLKEIEALIAKRKSHARVFAGFLLFGIGTSADVHLDICKTS--DDCHHAGVFI	2437
HarufH1	LEISGLNLKEIEALIAKRKSHARVFAGFLLFGIGTSADVHLDICKTS--DDCHHAGVFI	2437
HadishaH1	LEISGLNLKEIEALIAKRKSHARVFAGFLLFGIGTSADIHLDICKTS--DDCHHAGVFI	2437
HadivH1	LEISGLNLKEIEALIAKRKSHARVFAGFLLFGIGTSADIHLDICKTS--DGCHHAGVFI	2352
HaasH1	LEISGLNLNEIEALITKRKSHARVFAGFLLFGIGTSADVHLDICKTS--ENCHDAGVIFI	2436
HarubH1	LEISGLNLNEIEALISKRKSHARVFAGFLLFGIGTSADIHLDICKTS--ENCHHAGVIFI	2437
	. . . : : * : * * * : * . : : * * * . * . . :	

CoasHaN	LGGPLESPWAYNRLYKRDITQYLAALNLHPEDVDFDPTLQVHLEVEVRDQDQVHLEAQNVL	2514
HepoHaN	LGGPLESPWAYNRLYKRDITQYLEDLNLHPEDVDFDPTLQVHLEVEVRDQDQVHLEAQNVL	2513
HeluHaD	LGGPLEMPWAFDRLFKKDIWTWIARIGLNPDIDHKTDSGFKLEVQAFNVEGTALPLSQAI	2510
ApcaH1	LGGPIEMAWAFDRLFTYDITSALEQSGINPEDVDFDAEALFTLDIKVDFVEGHALPVSSVL	2513
LystH1	LGGPSEMPWAFDRLYKDDITWALEAADIKPESVFDADSLFTLDVKVDFVEGHAINVKSVM	2516
PhacH1	LGGPTEMEWAFDRLYKDDITWALEEAGISPEVFDADSLFTLDVKVYNVDGESINVKSVM	2518
PocaH1	LGGALEMQWSDRLRYRHDITQALLAVSVRQDLFTEHTPFTLDYSITAVNGTQLPRS-LM	2500
MetuMH	LGGEKEMWAFDRLYKYDITDALKDNGIQPEDVDFDAEAPFYLYKVEYAVNGTQLPRS-VI	2523
LisaH1	LGGHLEMPWTFDRLYKYDITDLDADISPEVDFNAQTPFTIKYSIHAVNGSALPLS-SV	2509
<b>CrfoH1</b>	<b>LGGKLEMPWAFDRLFKLDITDTHLDMGIEPEDVDFDQAPFFLSYEVHAVNGTTLPLS-TI</b>	<b>2502</b>
RaveH1	LGGALEMPWAFDRLYKYDITDLDLHDLGIEPEDVFNVAQPFHLKYEIHAVNGSTLAPS-TI	2515
OcvuHA	LGGETEMPWFDRTRYKMDITMVMKEKGIAMESLFHHDSEKVFHLEMEIITIDGTVLQDN-SL	2482
TopaH1	LGGETEMPWFDRTRYKMDITDVLKRNIPPEALFEHDSKIRLEVEIKSVGAVLDPN-SL	2896
Eusch1	LGGETEMPWFDRTRYKMDITDVLKRNIPPEALFEHDSKIRLEVEIKSVGAVLDPN-SL	2922
SeofH1	LGGETEMAWHFDRLYRFEITSVLEEMKIPFDKLFHEHESKIHWEVEIKVVDGTPVDAG-VI	2921
SemaH	LGGETEMPWFDRTRYRFEITSVLEEMKIPFDKLFHEHESKIHWEVEIKVVDGTPVDAG-VI	2922
LijaH2	LGSDLEMPWAFDRTFKYDITKALQKLGVSLS-----DDLYHLKVTIVAVNGTTLSDN-VI	2432
NunuH1	LGGEFEMPWAFDRLYKYDITSAVKEAGLNPNDFNAEAPFHLKLEITKVDGTQIPSS-EL	2490
MecrH1	LGGETKEMAWAYNRLFKYDITHALHDAHITPEDVHFHSEPFKIKVSVTAVNGTVLPAS-IL	2493
HacrH1	LGGSDEMHWAYNRLYKYDITTEALHEFGINPEDVHFADEAFFLKVSVAVAVNGTVLPAS-LL	2496
HarufH1	LGGSDEMHWAYNRLYKYDITTEALHEFGINPEDVHFADEAFFLKVSVAVAVNGTVLPAS-LL	2496
HadishaH1	LGGSDEMHWAYNRLYKYDITTEALHEFGINPEDVHFADEAFFLKVSVAVAVNGTVLPAS-LL	2496
HadivH1	LGGSDEMHWAYNRLYKYDITTEALHEFGINPEDVHFADEAFFLKVSVAVAVNGTVLPAS-LL	2411
HaasH1	LGGSDEMHWAYNRLYKYDITTEALHEFGINPEDVHFADEAFFLKVSVAVAVNGTVLPAS-LL	2495
HarubH1	LGGSDEMHWAYNRLYKYDITTEALHEFGINPEDVHFADEAFFLKVSVAVAVNGTVLPAS-LL	2496
	** . * * * : : * : * * : : . . : : . :	

CoasHaN	PQPTIIFDPPHEGAED---V-SSTSIAIGVGRKDVSTLSTSEIYINRSALRQVKDDAGAN	2570
HepoHaN	PQPTIIFDPPHEGAED---V-SSTSIAIGVGRKDVSTLSTSEIYINRSALRQVEDDGTGN	2569
HeluHaD	PKPSVTTYKALGVEKD---I-HTTAVAGVGRKDVTRTLTSETENLREAPRRRIKADNGSN	2566
ApcaH1	PEPTIYKAAVGAASEE---VSSSSSLAGVGRKDVSTLSTSEIYINRSALRQVADAGSN	2570
LystH1	PTPSIYDPAKGVANV---D-ITSVAGVGRKDVSTLSTSEIYINRSALRQVQEDGPN	2572
PhacH1	PTPSLIFDPAHGAED---V-SSTSIAIGVGRKDVSTLSTSEIYINRSALRQVADNGPN	2574

PocaH1	PPPTIIYEPASGGFRG---M-PSYSLAGIGVRKDINTLSEAETENLREALSSVMDTRI I	2556
MetuMH	NAPTLIFVPATGASVD---A-KSYAVAGMSVRKDINTLSSAETQSLKNSLQKVMDEGEPGL	2579
LisaH1	SPPTIIFEPALHAGASQD---H-TSYSVAGVGVKRDINTLTAETESLRSLEMRQEGSGSL	2565
<b>CrfoH1</b>	<b>SPPTLVFQPAEGAAAE---H-SSYSIAGVGVRKDINTLTAEMENLRDALGRVQAGTGRL</b>	<b>2558</b>
RaveH1	SAPTLIFKPAEGASVD---Q-TSYSIAGVGVRKDINTLTAEMENLRDALRRVQAGTGRL	2571
OcvuHA	PKPSLVYSPAKGVSQHVVH-HKEYAEGTIVRKNVNSLTPSEIDNLRHALSDVMADKSEN	2541
TopaH1	PKPSLIYAPAKGLIIQ---Q-VGEYDAGSGVRKNVNSLTPSEIENLRNALAAVQADKTD	2952
EuscH1	PKPSLIYLPKAKHHA---K-EQPEMAGTLIRKNVNSLTPYEVGNLRDALAAVQADITKT	2978
SeofH1	PKPSLIYLPKAPHLIE---K-EKPHSAGAAVRKNVNSLTPYETNNLRMALAAVQADITDT	2977
SemaH	PKPSLIYLPKAPHHIE---K-EKPHAAGAAVRKNVNSLTPYETDNLRDALAAVQADITDT	2978
LijaH2	PAPTLTFVFPASRSGRQ---A-EADHAAAPGIRKNVNDLSSSEMENLRAMRQVQDPSSR	2488
NunuH1	HKPTIIYEPAHGHES---V-GVASHAGRVRKNVNTLTPSEIDNLKDMARAVQADKGVN	2546
MecrH1	HAPTIIYEPGLDHHED---H-HSSSMAGHGVRKEINTLTTAEVDNLKDMARAVMADHGP	2549
HacrH1	HEPTILYEPGVGHED---H-ESGSLAGSGVRKDVNTLTTAETDNLRRALRGVKEDHGH	2552
HarufH1	HEPTILYEPGVGHED---H-ESGNLAGSGVRKDVNTLTTAETDNLKALRGVKEDHGH	2552
HadishaH1	HEPTILYEPGVGHED---H-ESGSLAGSGVRKDVNTLTTAETENLRKALRGVKEDHGH	2552
HadivH1	HEPTILYEPGVGHED---H-ESGSLAGSGVRKDVNTLTTAETENLRKALRGVKEDHGY	2467
HaasH1	HQPTIIYEPGEDHHED---H-ESGSIAGSGVRKNVNTLTTAETDNLREALRGVMDHGP	2551
HarubH1	HQPTIIYEPGEHDD---H-ESGSIAGSGVRKDVNTLTTAETDNLREALRGVMDHGP	2552
	* : : : . : * * * : * : * : : : .	
CoasHaN	GFLNIASFHGKPARCEH-DHHPVACCVHSGPKFPHWHRLYVKQWEDALTAHGAKIGIPYW	2629
HepoHaN	GFQNIASFHGSPPARCEH-DHHPVACCVHSGPNFPQWHRLYVKQWEDALTAHGAKIGIPYW	2628
HeluHaD	GFQNIASFHGSPPCVH-ENHSVACRIHGMANFPQWHRLYVKQWEDALTAQGAKIGIPYW	2625
ApcaH1	GFASIAAFHGEPAJCEL-NGRRIACQHGTMNFPQWHRLYVKQWEDALTAQGTMIPIYW	2629
LystH1	GFQNIASFHGSALACEY-QGHPIACCLHGMANFPQWHRLYVKQWEDALTSQGSKVGIPYW	2631
PhacH1	GFQNIASFHGSAPARCEH-NGHSIACCLHGMANFPQWHRLYVKQWEDALTAQGAMIGIPYW	2633
PocaH1	SYQRLAGWHGFPGLCSM-NGQQVACCVHSGSFPWHRLYVRSLEIAMTLEGARIGIPYW	2615
MetuMH	SYQKIAEMHGWPQDCEQ-NGKKVACCHGMASFPQWHRVYTRLLEMAMTWQGANVGIPIYW	2638
LisaH1	PFQQFASWHGWPASCEH-DGHPVACCHGMASFPQWHRLYVRILEMVMTFRGARVGIPIYW	2624
<b>CrfoH1</b>	<b>TYDNVVSAGHYPPQCTH-DGHKVVACCVHSGSFPWHRLYVTRQMEVALSWEGAKVGIPIYW</b>	<b>2617</b>
RaveH1	TYDFIAGAHGYPAAKCM-GEYDVACCVHSGSFPWHRVYTRQMEVALSWEGAKVGIPIYW	2630
OcvuHA	GYQKIAAYHGMPKSCYKPNGTAYACCQHGVMVTFPHWHRLYVKQMEDALKKGAKTGIPYW	2601
TopaH1	GYQKIASFHGMPLSCQYPDGTAFACCQHGVMVTFPHWHRLYMKQMEDALKKGAKTGIPYW	3012
EuscH1	GYQKIASYHGMPLSCYKDNGTALACCQHGVMVTFPHWHRLYMKQMEDAMRARGARVGIPIYW	3038
SeofH1	GFQKIASFHGMPLSCYKDNGSAYACCQHGVMVTFPHWHRLYMKQMEDALRARGAKVAIPIYW	3037
SemaH	GFQKIASFHGMPLSCYKDNGSAYACCQHGVMVTFPHWHRLYMKQMEDALRARGAKVGIPIYW	3038
LijaH2	GYQIAAAYHGLPAKCKNGQKMACCLHGMANFPWHRLYTKQMEDALAEKGARLGMPIYW	2548
NunuH1	GYQSIAYHGLPAQCG----DYACCLHGMVTFPHWHRLYVKQMEDALAAHGAEVGMPY	2601
MecrH1	GYQIAAFHGNPMPMPDGNKNSCCTHGMATFPHWHRLYTKQMEDALTAHGARGVGLPYW	2609
HacrH1	GFQIAAFHKGKPMCPMDGHNLSCTHGMATFPHWHRLYTKQMEDALKAHGSHVGLPYW	2612
HarufH1	GFQIAAFHKGKPMCPMDGHNLSCTHGMATFPHWHRLYTKQMEDALKAHGSHVGLPYW	2612
HadishaH1	GFQIAAFHKGKPMCPMDGHNLSCTHGMATFPHWHRLYTKQMEDALKAHGSHVGLPYW	2612
HadivH1	GFQIAAFHKGKPMCPMDGHNLSCTHGMATFPHWHRLYTKQMEDALKAHGSHVGLPYW	2527
HaasH1	GFQIAAFHKGKPMCPMDGHNYSCTHGMATFPHWHRLYTKQMEDALRAHGSVLVGLPYW	2611
HarubH1	GFQIAAFHKGKPALCPMDGHNYSCTHGMATFPHWHRLYTKQMEDALKSHGSLVGLPYW	2612
	: . . * * * : * * * . * * * : : * : . : . : * * *	
CoasHaN	DWTYAFKKLPSLVT-AEVDNPFHHGVTHDG---HVTTRAPRSLLFNDPEFGDESFFYRQV	2685
HepoHaN	DWTYAFKKLPSLVT-AGDNNPFHHGVTHDG---HITTRAPRSLLFNDPEFGDESFFYRQV	2684
HeluHaD	DWTTAFTALPALVT-EEVDNPFHHGTIYNG---EITTRAPRDKLFNDPEFGKESFFYRQV	2681
ApcaH1	DWTTAFTALPALVT-EQENNPFYFNKIFNG---EVTSRAPREQLFNDPEFGSESFFYRQA	2685
LystH1	DWTTAFTALPTLVT-EETNPNFHHGKIYNG---EITTRAPRDLFNDPEFGSESFFYRQA	2687
PhacH1	DWTTAFTALPTLVT-EEENNPFHHGKIYNG---EITTRAPREQLFNDPEFGSESFFYRQT	2689
PocaH1	DWTTTFTNLPSLLTADSSNPFKLGHIKALN--QSTSRSPRQLFNDPERGEESFFYRQI	2673
MetuMH	DWTEAFTALPSLVT-DDGDNPFNHGHIEALG--KVTSRPRPQLFKDPEHGEESFFYRQI	2695
LisaH1	DWTQAFTELPAALVT-EEENNPFHHGVIEAL--NMTSRAPRQLFNDPEHGEESFFYRQV	2681
<b>CrfoH1</b>	<b>DWTEAFTALPALVR-EEENNPFHHGRIPGTD--TVTTRAPRQLFRDPEHGEESFFYRQV</b>	<b>2674</b>
RaveH1	DWTEAFTALPTLVS-EDHDNPFHHGVPGKAENITTTTRAPRQLFKDPEHGEESFFYRQA	2689
OcvuHA	DWTFPFTDLPTLIT-EETDNPFFHHAHIDSVD--EITTRAPRHLFEG----RHFFYDQM	2653
TopaH1	DWTTAFHSLPILVT-EPKNNPFHHGYIDVAD--TKTTRDPRPQLFDDPEQDQSFYRQI	3069
EuscH1	DWTIAFSSLPDLVT-DEKNNPFHHAHIDVAN--TITTRNPRPQLFDDPEQDQSFYRQI	3095
SeofH1	DWTTAFSSLPALVT-DEKNNPFHHSYIDVAG--TITTRSPRQLFDDPEHGEESFFYRQI	3094
SemaH	DWTTAFSSLPALVT-DEKNNPFHHSYIDVAG--TITTRSPRQLFDDPEHGEESFFYRQI	3095
LijaH2	DWTMPFTSLPHLVT-ETENNPFNKVIVFKN--KYTTRDPVNLFRDPEYGEKSFFYRQV	2605
NunuH1	DWTTAFTALPALVT-DPNDNPFVHHAHIEYLN--EVTTRAPRQLFKDPEHNEGSFFYRQM	2658
MecrH1	DGTTAFTALPTFVT-DEEDNPFHHGHIYDLG--VDTTRSPRDKLFNDPERGESFFYRQV	2666
HacrH1	DWTTAFSHLPTLVT-DSDDNPFHHDGTIEYLN--VTTTRSPRDLFNDPEHGESFFYRQV	2669
HarufH1	DWTTAFSHLPTLVT-DNDNPFHHDGTIEYLN--VTTTRSPRDLFNDPEHGESFFYRQV	2669
HadishaH1	DWTTAFSHLPTLVT-DNDNPFHHDGTIEYLN--VTTTRSPRDLFNDPEHGESFFYRQV	2669
HadivH1	DWTTAFSHLPTLVT-DNDNPFHHDGTIEYLN--VTTTRSPRDLFNDPEHGESFFYRQV	2584
HaasH1	DWTTAFTALPTLVT-DTENNPFNHGHIEYLD--VDTTRSPRDLFNDPEHGESFFYRQV	2668
HarubH1	DWTTAFTALPTLVI-DTKNNPFNHGHIEYLD--VDTTRSPRDLFNDPEHGESFFYRQV	2669
	* * * * * : : . * * * . * * * * * . * * * *	
CoasHaN	LLAFEQTDYCDFEVQFEITHNAIHSWVGQSPYGLSTLEYTAYDPLFLHHSNVDRQFAI	2745
HepoHaN	LLAFEQTDYCDFEVQFEITHNAIHSWVGQSPYGLSTLEYTAYDPLFLHHSNVDRQFSI	2744



\* \* . : . \* : : . . : : \* \* : : \* \* \* : . \* . \* . : \* \*

CoasHaN	GDLEMDWEFDRLFKYDVTNVFNKHLHQPDSAYHFVDHITAVNGTELDNSLIRPPSVHFVFP	2925
HepoHaN	GYLEMDWEFDRLFRYDVTNVFNKHLHQPDSYHFVDHITAVNGTELDSQLIRPPSVHFVFP	2924
HeluHaD	GPLEHPWAFDRLFKYDVTNVFVSKLHLRDPSEYHFNIHIVAVNGTELDLSLIRSPVQFVFP	2921
ApcaH1	GAIEMPWAFDRLFKYDVTNVFRKLNHLPPDDVYHFVVKIMAVNGTELYPGLIRPPSVQFVFP	2924
LystH1	GLLEMPWAFDRLFKYDVTNVFNKLNLRDHDSDYHFVVEIKAINGTVLDSNLIRSPSVQFVFP	2927
PhacH1	GPLEMPWAFDRLFKYDVTNVFNKLNLRPDSNYHFVEVEIKAINGTILDSNLIRSPSVQFVFP	2929
PocaH1	GPLEMPWAFDRLFKYDVTNVFVKQLHLRDPSEYRFKMRRLTAVNGTELDPHLLHAPSVSFLP	2913
MetuH1	GPLEMPWAFDRLFKYDVTNVFRQMLRDPDSYSVKVIHIVSVNGTELDSDLLLEAPSVSFVP	2935
LisaH1	GPLEMAWRFDRLFKYDVTNVFRQMLRDPDSYVYHFRVHLTAVNGTELD AHLIEDPSVSFLP	2921
<b>CrfoH1</b>	<b>GPLEMPWAFDRLFKYDVTNVFVKQMLRDPDSSEYFRVSLTAVNGTQLDSRLIEAPSVSFVP</b>	<b>2914</b>
RaveH1	GPLEMPWAFDRLFKYDVTNVFRQMLRDPDSNYTIPIRIRAVNGMQLDPDLLEPPSVTFAP	2929
OcvuHA	GKHEMFWAFDRLFKYDITSTLEQLRLHAHQDSITVSIKATDGS TLAPSLIPPPS IMYHP	2893
TopaH1	GDYEMPWAFDRLFLYDISKSLVHLRDLAHDSDYHFVVEIKAINGTIMGIDGKSLPNNLPSPTILFKP	3309
Eusch1	GEHEMAWAFDRLFLYDITKALTKLSIDAYDDFKVQVGTITSITGEKLPQSLPPPTILFKP	3334
SeofH1	GQHEMAWAFDRLFLYDITKALNKLHLDAYDDFLINVSIVNIEGVKLPSSLLPRPTIMFKP	3334
SemaH	GQHEMAWAFDRLFLYDITKALNKLHLDAYDDFVINVSIVNIEGVKLPSSLLPKPTIMFKP	3335
LijaH2	GTKEMWAFDRLFKYEITEALSALRADHSDYHFVVEITTIATNKTKLASNLIRPRTVIYEP	2845
NunuH1	SEYEMHWYDRAFRYDITDMEELHLKPESEFTIKLHLVQADGTELDSDLVDSPEVIFVP	2898
MecrH1	GEHEMPWAFDRLFRYDITQVLKQMHLEYSDFTFHMRIIDTSGKQLPSDLIKMPTVEHSP	2906
HacrH1	GELEMPWAFDRLFRYDITKVMQQHLRHDSDFSFKVKVGTDDKELPPGILKEPTIEYEP	2909
HarufH1	GELEMPWAFDRLFRYDITKVMQQHLRHDSDFSFKVKVGTDDKELPHGILKEPTIEYEP	2909
HadishaH1	GELEMPWAFDRLFRYDITKVMQQHLRHDSDFSFKVKVGTDDIELPPGILKEPTIEFEP	2909
HadivH1	GELEMPWAFDRLFRYDITKVMQQHLRHDSDFSFKVKVGTDDIELPPGILKEPTIEFEP	2824
HaasH1	GDQEMPWAFDRLFRYDITKVMKQLHLRHDADFTFVSVKIVSTDGQELPSDSVKPTIEFEP	2908
HarubH1	GELEMPWAFDRLFRYDITKVMKQLHLRHDSDFTFRVKIVGIDDHLELPSDSFKAPTIEFEP	2909

CoasHaN	GVKKPHG-VAAAS---GPASGVLRKNIQNSQDEAVSLRDALYQLQQDQGLGGFEAIAG	2981
HepoHaN	GVKKPLG-VAAAS---GPASGVLRKNIQNSQDEAVSLRDALYQLQQDQGLGGFEAIAG	2980
HeluHaD	GVKDYELTAQKT---EAHEDVLLRKNINLSLEESANLRSALNKLQQDQGPNGFEAIAG	2978
ApcaH1	GVKGYERVAAKT---AKSASVLRKDVNDLTLAEASNLRDALYQLQQDQGPNGFEAIAG	2981
LystH1	GVKRFYKVAEKT---VVRDDKLRKNIQNSQDEAVSLRDALYQLQQDQGPNGFEAIAG	2984
PhacH1	GTKKFYKIAEKT---AKRDKLRKNIQNSQDEAVSLRDALYQLQQDQGPNGFEAIAG	2986
PocaH1	GRGEQRA-RAAREDPVTTVSSVVTRYDVTTLTLEQASSLRLNHLKQNDHGLTYSYEAIAS	2972
MetuMH	GKRGR----KAREDYRPTVSEDLIRKEVSSLSLEEVSNLKNALYKQNDHGPNGFEAIAS	2991
LisaH1	GRDHGVSHAQAHEDPIPLASEDVTRYEVSSLSLEQSSNLRNALYKQNDHGPNGFEAIAS	2981
<b>CrfoH1</b>	<b>GNKPGSKTSAAHEDPVPLDTGDVTRYEVSSLSLAQVTLNLRDALYKQNDHGPNGFEAIAS</b>	<b>2974</b>
RaveH1	GKKSQGRAVVYEEPTPSETQDLTRHDLSLSMAQISNLEDALYKQNDHGPNGFEAIAS	2989
OcvuHA	AAA-----	2896
TopaH1	GTGKI-----	3314
Eusch1	GKGSQQHH-----	3342
SeofH1	GKGTQHHH-----	3342
SemaH	GKGTQHHH-----	3343
LijaH2	GKAST---STKRV--TQPKRTALLRKGNTLTLDEAKNLRAMFMMNDEGPNGFIAIAT	2900
NunuH1	GSAT---HEHHTV--HEDHRDVLRLKNIQNSLSEARSRLDALYKQNDQTENGFEHIAG	2953
MecrH1	----GKHEKHH--EDHEDILVRKNIHLSLHHEAEELRDALYKQNDHSHGGYEHIAIAG	2960
HacrH1	ALGDGLKHEDEGH--DDRLSHVLRKEVDFLSLKEANA IKDALYKQNDHSHGGYEHIAIAG	2967
HarufH1	ALGDGLKHEDEGH--DDRLSHVLRKEVDFLSLKEANA IKDALYKQNDHSHGGYEHIAIAG	2967
HadishaH1	ALGDGLKHEDEGH--DDRLSHVLRKEVDFLSLKEANA IKDALYKQNDHSHGGYEHIAIAG	2967
HadivH1	ALGDGLKHEDEGH--DDRLSHVLRKEVDFLSLKEANA IKDALYKQNDHSHGGYEHIAIAG	2882
HaasH1	GVQDGRKLDGEHH--DDRHSVLRKEVDFLSLQEANA IKDALYKQNDHSHGGYEHIAIAG	2966
HarubH1	GVHDGKHKDDGHH--DDRHSVLRKEVDFLSLQEANA IKDALYKQNDHSHGGYEHIAIAG	2967

CoasHaN	FHGAPFLCPEHGAE-KYACCVHGMPSFPFHRLFTVQFEQSLKQHGSTTGIPYWDWTSPG	3040
HepoHaN	FHGAPFLCPENGDK-KYACCVHGMPTFPFHRLFTVQFEQALKLHGSTTGIPYWDWTSPG	3039
HeluHaD	FHGAPFKCPENGTD-KYACCVHGMSPFPFHRLFTVQFEQALKAHGAKGVPYWDWTAPI	3037
ApcaH1	YHGAPFKCPANGED-KYACCAHGMSPFPFHRLFTVQFEQALKEHGALVGVYWDWTAPI	3040
LystH1	FHGAPFRCPATGDD-KYACCVHGMSPFPFHRLFTVQIEQALKSNGALIGVYWDWTTPV	3043
PhacH1	FHGAPFKCPADGDD-KFACCVHGMSPFPFHRLFTVQVEQSLKKNALVGVYWDWTAPV	3045
PocaH1	FHAGSGLCPENASV-TFACVPHGFANLPHFNRLLLVQMEMALLEKATGIPYWDWTRTI	3031
MetuMH	FHGDPLCPEGSGS-KYACCVHGMSPFPFHRLFTVQFERALKKAGSVGVYWDWTRPS	3050
LisaH1	FHGAPGLCPENGGKLRACCRHGMPFPFHRLFTVQFERALRDKGAVGVYWDWTRPV	3041
<b>CrfoH1</b>	<b>FHGAPGLCPENATD-HYACCRHGMPFPFHRLFTVQFERALKKAGAVGVYWDWTRPA</b>	<b>3033</b>
RaveH1	YHGAPGLCPENATE-HYSCCVHGMSPFPFHRLFTVQFEHALKEKAGAVGVYWDWTRPA	3048
OcvuHA	-----	2896
TopaH1	-----	3314
Eusch1	-----	3342
SeofH1	-----	3342
SemaH	-----	3343
LijaH2	FHGAPGKCPGATENTFSCCIHGMPFPFHRLFTVQAEKSLQDHGATVGIYWDWTRSM	2960
NunuH1	FHGPNVCPEDSE-SKYACCAHGMPIFPFHRLFTVQFERGLVDNGALIGLPIYWDWTRPS	3012
MecrH1	FHGYPNLCPKEDG-EKYPCCVHGMSPFPFHRLFTVQFERALKKAGSHLGIYWDWTRTI	3019
HacrH1	YHGYPNKCPKEDG-DKYPCCVHGMPIFPFHRLFTVQMERALKKAGSHLGIYWDWTRSM	3026
HarufH1	YHGYPNKCPKEDG-DKYPCCVHGMPIFPFHRLFTVQMERALKKAGSHLGIYWDWTRSM	3026

HadishaH1	YHGYPNKCPEKGD-DKYPCCVHGMPIFPPHWRHLHTIQMERALKNHGSQIGIPYWDWTKRM	3026
HadivH1	YHGYPNKCPEKGD-DKYPCCVHGMPIFPPHWRHLHTIQMERALKNHGSQIGIPYWNWTKRM	2941
HaasH1	YHGYPNMCPERGS-EKYPCCVHGMPIFPPHWRHLHTIQMERALKNHGSPMGIPYWDWTKKM	3025
HarubH1	YHGYPNMCPHGT-DKYPCCVHGMPIFPPHWRHLHTIQMERALKNHGSPMGIPYWDWTKKM	3026

CoasHaN	NELPLFLADTSDNPFSGYTI SFAGQRTSRNPLEALFSTN--TSAGTSLLYQLTLDALEE	3098
HepoHaN	NELPLFLADTDNDNPFSSYTI SFVQRTSRNPLGALFSTN--TSAGTSLLYQLTLDALEE	3097
HeluHaD	GKIPSLFGDSADYNPFYSYTI SFNNQRTTRDIQSELYNPH--QINGYNYLYYLLALSTLEE	3095
ApcHa1	NALPSLIGDSSNNHNPFFYKYHISFVNQDTRDIQDLSLFNPR--TINGYNYLYYLLALSTLEE	3098
LystH1	KSLPSFFGDDADHNPFAFRIAAANEKTRQVQSELYSER--KVHGFYLYYLLALSTLEE	3101
PhacH1	QSLPSLFGDSSDFNPFYSFQIASANHKTRDQVQDLYSNR--KVNQYHYLYYLLALNTLEE	3103
PocaH1	RALPSLVAES-GDNSSFFSYHIRQANKDTRDQDLYVAS--RGTRNLIYLYYLLALALEE	3087
MetuMH	RGIPQLFSNTYGNPNPFLTYEIQAAASAFVERDQVDALEHTTGHEGFEDSLFHQALETLEE	3110
LisaH1	KALPSLLTDFDNNPFYSYRISFVEQYTRDQVQALFKHP--SEGDLSELYYLLALALEE	3099
<b>CrfoH1</b>	<b>KAMPSLFTDSYDNNPFQTYRMTFNDQYIQRDVSELEFNHP--SEGDSLELYYLLALALEE</b>	<b>3091</b>
RaveH1	KGMPTLFTDNSAINPFYTYLMSFNNHHIERKVHEDLFSHS--AEGDPESLYYLLALALEE	3106
OcvuHA	-----	2896
TopaH1	-----	3314
EuscH1	-----	3342
SeofH1	-----	3342
SemaH	-----	3343
LijaH2	KALPNFVVSQAE-NNPWHRGYIRFEGKYTRTRVDPRDFDP-QVS-GRNFLHNNVLLSLEE	3017
NunuH1	KALPSLFAEES-NNPFYKYHIGFANTDTEGDVQDFLNFQP-SFYGKYNYLYYLLALTTLEE	3070
MecrH1	SSLPTFFADSGNPNPFYKYHIRSINQDTRDQVNEAIFQQT-K-FGEFSSIFYLALQALEE	3077
HacrH1	SSIPAFFGDDSNPNPFYKYHIRAVNQYTRDQVDFLNFQT-K-FGEYDLYYLLALQVLEE	3084
HarufH1	SSLPAFFGDDSNPNPFYKYHIRAVNQYTRDQVDFLNFQT-K-FGEYDLYYLLALQVLEE	3084
HadishaH1	SSIPAFFGDDSNPNPFYKYHIRAVNQYTRDQVDFLNFQT-K-FGEYDLYYLLALQVLEE	3084
HadivH1	SSIPAFFGDDSNPNPFYKYHIRAVNQYTRDQVDFLNFQT-K-FGEYDLYYLLALQVLEE	2999
HaasH1	SSLPAFFGDDSGNPNPFYKYIRGVHETTREINQMQMNFQT-K-FGEYDLYYLLALQVLEE	3083
HarubH1	SSLPAFFDTAGGNPFYKYIRGVQHETTREINPIIFNFQT-K-FGEYDLYYLLALQVLEE	3084

CoasHaN	DDYCFEIMLEFFHNRIHFLIGGSETYSMSTLDYSADFPIFMIVHSGMDRLWVWLQELQK	3158
HepoHaN	DDYCFEIMLEFLHNRIHFLIGGSETYSMSTLDYSADFPIFMIVHSGMDRLWVWLQELQK	3157
HeluHaD	DNFCDFEVQYEVILHNEIHALIGGNGTYSMATLDFSAFEPFMIHSSLDRIWIWQELQK	3155
ApcHa1	DSFCDFEIQYEILHNEIHGLIGGHTYSMSTLDYSADFPLFMIHSSIDRIWAIWQELQK	3158
LystH1	DNYCDFEVQYEVILHNEIHADIGGSGTYSMATLDYSADFPIFMIVHSSIDRIWIWQELQK	3161
PhacH1	DNYCDFEVQYEVILHNEIHATIGGSGQYSMATLDYSADFPIFMIVHSSIDRIWIWQELQK	3163
PocaH1	VNFCDFLVQMDLLHVRVHALVGGKEAFSMATMEHAADFPLFWLHAANVDRLWQAWQELQK	3147
MetuMH	TSYCFEVEVQYFEMLHNAVHALVGGKTKTHSMATLEWSADFPIFMIVHSSIDRIWRIWQELQR	3170
LisaH1	NNYCDFEVQYEMLHNAVHELVGGRKYGSMATLEYSADFPIFMIVHSSIDRIWQIWIQTLQK	3159
<b>CrfoH1</b>	<b>TNYCDFEVQYEMLHNAVHELIGGNGTYSMATLEYSADFPIFMIVHSSIDRIWQIWIQTLQK</b>	<b>3151</b>
RaveH1	TNYCDFEVQYEMLHNAVHELVGGRKYGSMATLEYSADFPIFMIVHSSIDRIWQIWIQTLQK	3166
OcvuHA	-----	2896
TopaH1	-----	3314
EuscH1	-----	3342
SeofH1	-----	3342
SemaH	-----	3343
LijaH2	DDYCDFEVQYEIFHNTIHAWVGGADPRSLSTLEYSADFPLFIHHSNMDRLWIWQELQK	3077
NunuH1	DNYCDFEVQYEVILHNEVHALIGGAEKYSMATLDFSAFDPFMIHSSIDRIWIWQELQK	3130
MecrH1	DNYCDFEVQYEVILHNEVHALIGGAEKYSMATLDFSAFDPFMIHSSIDRIWIWQELQK	3137
HacrH1	NSYCDFEVQYEVILHNAVHAYLGGAGKYSMATLEYSAYDPVFMIVHSSLDRIWIWQELQK	3144
HarufH1	NSYCDFEVQYEVILHNAVHAYLGGAGKYSMATLEYSAYDPVFMIVHSSLDRIWIWQELQK	3144
HadishaH1	NSYCDFEVQYEVILHNAVHAYLGGAGKYSMATLEYSAYDPVFMIVHSSLDRIWIWQELQK	3144
HadivH1	NSYCDFEVQYEVILHNAVHAYLGGAGKYSMATLEYSAYDPVFMIVHSSLDRIWIWQELQK	3059
HaasH1	NSYCDFEVQYEVILHNAVHAYLGGAGKYSMATLEYSAYDPVFMIVHSSLDRIWIWQELQK	3143
HarubH1	NSYCDFEVQYEVILHNAVHAYLGGAGKYSMATLEYSAYDPVFMIVHSSLDRIWIWQELQK	3144

CoasHaN	LRRKPFVAIECGEK-SLHEPLHPFDYD-INTISLTRENAVPSIFDHHHLQYDYDTTEIS	3216
HepoHaN	LRRKPFVAIECGEK-SLHEPLHPFDYD-INTIALTREHAVPTLFDHHHLGYEYDTTEIS	3215
HeluHaD	LRHKPFNYAHCGGH-VLDDPLHPFSGEINKNDLTRLNSQPSVFDYTHFGYEFDKLELN	3214
ApcHa1	LRRKPFNSARCAGG-IMEEPLQPFYSQINTNDFTRMNSQPSKVFYAHLYGFEFNDLELN	3217
LystH1	LRHKPFNGASCAGH-IMERPLQPFYSPEVNKNEFTRLNSVFNVDSDERLGYKYDKLELN	3220
PhacH1	LRHKPFNSARCAGH-IMERPLQPFYSPEVNHNEFTRLNSVFNVDSDERLGYQYDNLELN	3222
PocaH1	LRRRSYHSGSCARV-TEDTPLPFSSETLNPNPVTHANARPVQSVQVEVDKFRYSYDHLDFN	3206
MetuMH	LRKPFNVARCAMR-YLRKPLEPFYSASVNTDEVARTNSRPIDIFDTAKFHYDFDNLDLG	3229
LisaH1	LRHKPFNFARCAGR-SLFKPEPFYASVNTDPITRANAEVHVFDTEKPHYHYDSLELN	3218
<b>CrfoH1</b>	<b>LRHKPFNYASCASR-SLYKPLEPFYSYSLNADPLTLNNAQPVHIFDTAKFHYHYDSLDLN</b>	<b>3210</b>
RaveH1	LRHKPFNYAVCAAR-SMYKPLEPFYSINPDPLTRENSKPAQIFDTHKPHYHYDNLNLN	3225
OcvuHA	-----	2896
TopaH1	-----	3314
EuscH1	-----	3342
SeofH1	-----	3342
SemaH	-----	3343
LijaH2	LRGKPYNSAPCAGR-KMSLPMQPFHFDLSLHNTLTKENSRAINAFDHHAFGYEYDSLDLN	3136

NunuH1	LRHKSFSYAECKH-HLDRPLHPFNYASVNHNELTRTHSMFNQAADQSAFGYWDNLDMN	3189
MecrH1	RRVKPAHAGSCAGD-IMHVPLHPFNYESVNNDDFTRENSLPNAVVDShRFNYKYDNLNLH	3196
HacrH1	RRMKPYAAADCAGD-LMKVPMHPFSYKSENEDEFTRVNSVFNIVFDHYKFNIDYDNMRIR	3203
HarufH1	RRMKPYAAADCAGD-LMKFPMHPFSYKSENEDEFTRVNSVFNIVFDHYKFNIDYDNMRIR	3203
HadishaH1	RRMKPYAAADCAGD-LMKFPMHPFSYKSENEDEFTRVNSVFNIVFDHYKFNIDYDNMRIR	3203
HativH1	RRMKLYAAADCAGD-LMKFPMHPFSYKSENEDEFTRVNSVFNIVFDHYKFNIDYDNMRIR	3118
HaasH1	RRMKPYAALDCAGDRLMKAPLHPFNYESVNEDEFTRANSYPNSVFDHYRNFYEDNLRIR	3203
HarubH1	RRMKPYAALDCAGDRLMKAPLHPFNYETVNEDEFTRTNSYPNIVFDHYRNFYEDNLRVR	3204
CoasHaN	GHDAAEVLEIIRRRHSETRVYLGGAAGQYGESYRTEWVNLDA--GEKFSVGTNFFLLGSA	3274
HepoHaN	EHDAAEVLEIIRRRHSETRVYLGGAAYGHGQSFRTAAWVNLDA--GEEFDAGTNFFLLGSS	3273
HeluHaD	GHDVQGIIDIIHNLRHENRVYLGFLVFGQSSLEYKIDLIDDA--GQAHTAGSFHLLGGE	3272
ApcaH1	GHDVNDLNNIINLRDQDRVYLAFFNCCGKQGSFAFDVHFETAE--NNVPLGRFVYVLGGE	3275
LystH1	GNSVEEINNIINLHHQERIFVGLAFGHQKSLTIHISLINDN--DEAFDGGNIHILGGE	3278
PhacH1	GKNVQEIHEIINKLNQDQDRVIFGVIGGLQQLSLTLDISLLND--NGVAAAGAVHVLGGE	3280
PocaH1	RRSVSELLEATQSLRVKDRLFAAFLLSGVHTSARLHVTLQTS--DEGAVEVGSIIYLLGGL	3265
MetuMH	GHSIGEVMKINDMRSKTRLFVGFVLSGIETSAVKIEMDDSQ--GNSHEVGTFFVVLGGH	3287
LisaH1	GHSVKQLQEMVDSMRNSPRVFGFLLHGSRTSASVHVDVVAAD--GSKVNAGNFVVLGGE	3276
<b>CrfoH1</b>	<b>GHGVEHLHDMIEDMQATSRI FAGFVLSGIST SARVHVDVTRGE---DTVSVGNFVVLGGS</b>	<b>3267</b>
RaveH1	GHSVAELNMIHAMQAQSRTFAGFVLSGIST SARVHVDIYKGD---EVASVGNFVVLGGP	3282
OcvuHA	-----	2896
TopaH1	-----	3314
EuscH1	-----	3342
SeofH1	-----	3342
SemaH	-----	3343
LijaH2	DLGVNQLLAYIKEIKQHDFVFGFNTHGFSASANVDFVICRRNDNECNNYAGSFSVLGGP	3196
NunuH1	HHTVAELAEIINSLRNSEVRFAGFVHLHGASATRKVFN-----DHYAGTFNVLGDR	3242
MecrH1	GHNIEELEEVLRLSLRLKSRVRFAGFVLSGIRTTAVVKVYIKSG-TDSDEYAGS FVILGGA	3255
HacrH1	GHDINELEAIINLNRNKDRIFAGFVLSGIRITATVKVFIHETGVTDHEEFAGKFAILGGE	3263
HarufH1	GHDINELEAIINLNRNKDRIFAGFVLSGIRTTATVKVFIHGTGATDHEEFAGKFAILGGE	3263
HadishaH1	GHDINELEAIINLNRNKDRIFAGFVLSGIRITATVKVFIHGTGATDHEEFAGKFAILGGE	3263
HativH1	GHDINELEAIINLNRNKDRIFAGFVLSGIRITATVKVFIHGTGATDHEEFAGKFAILGGE	3178
HaasH1	GHDIQEVEEILRELNRDRVRFAGFVLSGIRITAMVKVFIHSKNETRHEEYAGEFAILGGE	3263
HarubH1	GHDIHLEEVLKELRNKDRIFAGFVLSGLQISAPVKVYIHSKNETIHEEYAGEFAVLGGE	3264
CoasHaN	KEMPWSNEIILWKFDITNAVRQAGVST-DKNIKFHFI AETYNGLT-HHD-QVAYGIFVIRE	3331
HepoHaN	KEMPWANEIILWKFDITTAHVHQAGVST-DKNIKFHFKIETYNGLT-FAS-ESAYAIFLVRE	3330
HeluHaD	REMPWAYERLFKYDITDVAKKYDITD-DHPIKVKVTSTYNGEP-HQEYT-DEIVAVERH	3329
ApcaH1	REMPWCFERVFVKYDVAEVLQANNVDI-HKPVKIGGQLFRYDGEH-VQNFTTYATFIERP	3333
LystH1	KEMPWAYERIMKLDVTEAIRKGIIST-DHAVKARFTSTDYQGNL-NHQ-DTDYAIIVERH	3335
PhacH1	KEMPWAYERLMKIDITETVKASSIDV-SHAVKARFTSTDYTGSK-IKE-AEDYAI FVERR	3337
PocaH1	QERWHERAYKLDVTEAARLELDP-YSTFDHNVSLFDYTGQP-LPY-TLPYPLVLYRP	3322
MetuMH	NEMPWAYERVYKYEITEAAKKYGLDH-DSVFNHFKVVKYDQGE-LSA-QFKAPILVERP	3344
LisaH1	EEMPWAYERPKLDITDAISKAGLAH-GDVTVHLSTSVARYDGTPLNV-TFPSSLIERP	3333
<b>CrfoH1</b>	<b>SEMPWAYERIYKLDMTEAASKLGLSS-ESTFHFKLTVTKYDGTALNV-TFPDPVIVKRA</b>	<b>3324</b>
RaveH1	SEMPWAYERIYKLDMTEAASKLGLSG-SSHDFDKLTVTKYDGSALDV-KFPDPVIVKRR	3339
OcvuHA	-----	2896
TopaH1	-----	3314
EuscH1	-----	3342
SeofH1	-----	3342
SemaH	-----	3343
LijaH2	TEMPWAFERPFKYEVTELFNKLGKY-TDDIKVRTEVKATNGSA-L-NIEFPLPVIYVRP	3253
NunuH1	KEMPWAYERLFKYDITEVLENNL NAGHNDITFRCEVTALDGTALDNPNTLSDPVLVLRP	3302
MecrH1	KEMPWAYERLYRFDITETVHN--LNLTDHVKFRFDLKKYDHTTE-LDASVLPAPIIVRRP	3312
HacrH1	KEMPWAYERLLKLDITDAVHR--LHLKDEEIRFRMEVTAYNGVP-VS-TKLADPLIVHRP	3319
HarufH1	KEMPWAYERLLKLDITDAVHR--LHLKDEEIRFRMEVTAYNGVP-VS-TKLADPLIVHRP	3319
HadishaH1	KEMPWAYERLLKLDITDAVHH--LHLKDEEIRFRMEVTAYNGVP-VS-TKLADPLIVHRP	3319
HativH1	KEMPWAYERLLKLDITDAVHH--LHLKDEEIRFRMEVTAYNGVP-VS-TKLADPLIAHRP	3234
HaasH1	KEMPWAYERMLKLDITDAVNK--LHVKDEDIRFRMEVTAYNGDI-VT-TKLSQPFIVHRP	3319
HarubH1	KEMPWAYERMLKLDITDAVNK--LHLKDEDIRFRMDVTA YNGDI-VT-TQLSQPFIVHRP	3320
CoasHaN	AYTDYLTNIPVGG-NLPLPSKLVVPKKT HVRFLPVSEDY--NV-PIENLVSYTNYNKC	3386
HepoHaN	AYTDYLTNIPVGV-GYPLPSKLVVPKKT HVRFLPVSEDY--NI-PIENLVSYTNYNKC	3385
HeluHaD	ADTDYDIVVIVPSK-DNTLVPKIVVKKGT RIEFVT--SDL---TD-PLEDLGSYTMMHK	3382
ApcaH1	AGVDYDIINIPVGR-NSSINPKTVVKKGT RFRFVT--DGV---ET-PMENLGSFTNFHFC	3386
LystH1	AEQDYDVVEIPVGR-KYPLPPKIVLKKGSRVKFYPGSDTF--NS-PLENLGSYTSFSKC	3390
PhacH1	ANTDYDVVNIPIGR-EQPLPPKISLKKGT RVRFI PVSDSF---TK-PMEDLGSYTSYNRC	3392
PocaH1	ASVDFDVLVYPLYV-DKALPPKVTVRRTKIRFHAADPSL---QGRRI RTFGSYTLFIKC	3378
MetuMH	AGVDYDVAIFQLKK-DNDLAPKVVLP RGT RIRFHS AEDGV---SS-VLREMGSYTNNAVY	3399
LisaH1	ANADYDVLVVQFGA-EQAKDYKVRV KRGT RVRFHSTLEGY---VG-QVKELGTYTNSQLC	3388
<b>CrfoH1</b>	<b>ANSQHDVLVPLSV-ANQLPPKIVVRQGTQVVFHASESGV---S-SLREVGSYTNSVHC</b>	<b>3378</b>
RaveH1	TNAEFDELILPIRK-ENKLPKIVVRGT RVL FHPTEEGI---LG-PIRELGSFTNSKLC	3394
OcvuHA	-----	2896
TopaH1	-----	3314

Eusch1	-----	3342
SeofH1	-----	3342
SemaH	-----	3343
LijaH2	PISENDVYVISIGQ-GRTPAHKIVAKRGRTRFIVNPFDAESE--GKRYIMDLSSYTPFYRC	3310
NunuH1	ARVNYDVII FQMAPTNHNPPAKVIVKKGTKLQYHPHAAIVQMGLSDAVAELGSYTSWSKC	3362
MecrH1	NNAVFDIIEIPIGK-DVNLPPKVVVKRGTKIMFMSVDEA----VTPMLNLGSYTAMFKC	3367
HacrH1	AHASHDILVIPVGM-GHELPPKVVVKSGTKIEFTPIDSS----VDRAMVELGSFTAMAKC	3374
HarufH1	AHASHDILVIPVGM-GHQLEPPKVVVKSGTKIEFTPIDSS----VDRAMVELGSFTAMAKC	3374
HadishaH1	AHASHDILVIPVGM-GHELPPKVVVKSGTKIEFTPIDSS----VDRAMVELGSFTAMAKC	3374
HadivH1	AHASHDILVIPVGM-GHELPPKVVVKSGTKIEFTPIDSS----VDRAMVELGSFTAMAKC	3289
HaasH1	AHVSHDILVIPVGA-GHDLPPKVVVKSGTKIEFTPIDSS----VNRPMVELGSFTAMAKC	3374
HarubH1	AHVSHDILVIPVGA-GHALPPKVVVKGGTKIEFTPIDSS----VDRSMVELGSFTALAKC	3375

CoasHaN	SLPPFAFDSYAVNVVHALQPGDYFFTSFS--KEHCTKGN-KIYIHVQDH--	3432
HepoHaN	SLPPFAFDSYAVNVVHALQPGDYFFTSFN--KEYCTQDK-KIYIHVQDH--	3431
HeluHaD	KIPPFNSNSYALNTVYKLSPGDYFFVPKN--VDLCNAGR-RIQITVEDD--	3428
ApcH1	NIPPFYSYHSFPFGEVHALSPGDYFFVNRD--VDLCNSGR-KLQITVADE--	3432
LystH1	SIPPFYSYALGVVHTLQPGDYFFVPKD--KALCEAGK-RIQITVEDE--	3436
PhacH1	AVPPFSYQSYALNTVHSLEPGEHFFVAAD--KALCAEGK-RVIAVEDE--	3438
PocaH1	EALPGNADALSLDVTYSLNPGEYLLALDGDLPKCLEAG-RTILVIDEE--	3426
MetuMH	SIPPGDANTYKLEVEYTLPEGDYFFVSN--EEKCKQGD-RIQISIDEE--	3445
LisaH1	VIPPGLANSYKLEMEHKLEPGDYFFTDFD--RSGCLAGH-RIMISVDQDWN	3436
<b>CrfoH1</b>	<b>AIPPGQANLYDLDVAYSLEAGDYFFTS--KTKCQGS-RIQITVDE--</b>	<b>3424</b>
RaveH1	AIPPGHAHAYELDRLVLEPGDYFFVSN--VESCKAGS-RIQISVDEE--	3440
OcvuHA	-----	2896
TopaH1	-----	3314
Eusch1	-----	3342
SeofH1	-----	3342
SemaH	-----	3343
LijaH2	DVPEFGFQKYNFNQBYTLAPGTYFFTTAD--SETCKQGA-KFILEIAE--	3355
NunuH1	IVPPFGYRRYQFNTVYSLSPGNYYFSAPT--VEMCKQNR-KLILAVEEE--	3408
MecrH1	KVPPFSFHAFELGKMYSVESGDYFMTAST--TELCNDNLRIVHVHVDDE--	3414
HacrH1	IVPPFTYNAFELNKVYSVDHGDYYITAGT--HELCEK-NVRLNVHVEEE--	3420
HarufH1	IVPPFTYNAFELNKVYSVDHGDYYITAGT--HELCEQ-NVRLNVHVEEE--	3420
HadishaH1	IVPPFTYNAFELNKVYSVDHGDYYITAGT--HELCEQ-NVRLNVHVEEE--	3420
HadivH1	IVPPFTYNAFELNKVYSVDHGDYYITAGT--HELCEQ-NVRLNVHVEDE--	3335
HaasH1	IVPPFSYNGFELDKVYSVEHGDYYITAGT--TGLCEQ-NLKLNIHVEHE--	3420
HarubH1	IVPPFTYHGFEELNKVYSVHHGDYYITAGT--KELCKQ-NLRLNIHVEHE--	3421