

Supplementary Material

A microscopic insight from conformational thermodynamics to functional ligand binding in proteins

Samapan Sikdar ¹, J. Chakrabarti ^{1,2} * and Mahua Ghosh ¹ *

¹
Department of Chemical, Biological And Macromolecular Sciences, S. N. Bose National
Centre for Basic Sciences, Sector III, Block JD, Salt Lake, Kolkata 700098, INDIA

²
Also at Unit of Nanoscience and Technology-II and The Thematic Unit of Excellence on
Computational Materials Science, S. N. Bose National Center for Basic Sciences, Sector III,
Block JD, Salt Lake, Kolkata 700098, INDIA

*Corresponding authors:

E.mail: jaydeb@bose.res.in and mahuaghosh@bose.res.in

Supplementary Material:

Supplementary Tables S1-S3

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Table S1. Conformational thermodynamic changes (kJ/mol) of all the dihedrals of the metal binding loop aspartates upon Ca^{2+} binding to native apo-aLA. The ΔG_i^{conf} and $T\Delta S_i^{conf}$ in the last two columns indicate the total change for each residue by adding individual dihedral contributions.

	ΔG_{ϕ}^{conf}	$T\Delta S_{\phi}^{conf}$	ΔG_{ψ}^{conf}	$T\Delta S_{\psi}^{conf}$	$\Delta G_{\chi^1}^{conf}$	$T\Delta S_{\chi^1}^{conf}$	$\Delta G_{\chi^2}^{conf}$	$T\Delta S_{\chi^2}^{conf}$	ΔG_i^{conf}	$T\Delta S_i^{conf}$
D82	-0.75	-1.25	2.0	-0.2	-0.5	-1.1	-0.05	-1.0	0.7	-3.5
D84	0.5	0.1	-0.1	1.65	1.5	-0.2	0.25	2.75	2.15	4.3
D87	-0.4	-0.3	-0.3	-0.3	-0.25	-0.3	0.25	2.2	-0.7	1.3
D88	-0.1	-0.05	0	0	0.2	0.35	0.2	1.9	0.3	2.2

Table S2. Conformational thermodynamic changes (kJ/mol) in residues from A1 and A2 helices upon Ca^{2+} / Mg^{2+} binding to native apo-aLA.

Residue #	Residue	$\Delta G_i^{conf} (\text{Ca}^{2+})$	$T\Delta S_i^{conf} (\text{Ca}^{2+})$	$\Delta G_i^{conf} (\text{Mg}^{2+})$	$T\Delta S_i^{conf} (\text{Mg}^{2+})$
5	LYS	1.0	6.4	0.3	-0.8
6	CYS	0.3	0.0	0.3	0.2
7	GLU	0.6	4.9	0.9	3.7
8	VAL	-0.3	-0.9	0.2	0.2
9	PHE	1.3	7.0	0.6	2.8
10	ARG	-1.5	-6.2	2.8	2.9
11	GLU	0.6	5.6	0.2	0.2
12	LEU	0.05	-1.1	-1.1	-4.9
13	LYS	0.12	-4.0	-0.8	-3.5
14	ASP	-0.6	-1.4	-2.8	-2.3
15	LEU	-0.3	-3.1	0.1	-1.6
16	LYS	-0.7	-5.3	0.3	2.3
17	GLY	0.0	-0.3	0.7	1.0
18	TYR	-0.6	-4.2	1.0	1.2
19	GLY	-0.3	-0.2	-0.5	-0.4
20	GLY	0.2	0.2	0.3	0.5
21	VAL	-0.9	-1.7	-0.4	-3.4
22	SER	-0.3	-5.0	0.0	0.0
23	LEU	-0.5	-0.5	-0.5	-3.5
24	PRO	0.1	1.1	0.9	1.3
25	GLU	-0.9	-2.0	-0.8	-4.1
26	TRP	0.9	0.1	-0.4	-0.7
27	VAL	-0.4	-3.0	-0.3	-3.3
28	CYS	0.1	0.0	0.0	-0.1
29	THR	0.0	-1.5	-0.1	-0.7
30	THR	-0.2	0.6	-1.3	-2.8
31	PHE	-0.6	-4.9	-0.4	-2.1
32	HSE	0.2	2.1	0.2	0.6
33	THR	0.1	-0.4	-0.9	-2.1
34	SER	-0.8	2.2	-0.6	-2.0

Table S3. Conformational thermodynamic changes (kJ/mol) at the interfacial residues upon $\text{Ca}^{2+}/\text{Mg}^{2+}$ binding to native apo-aLA

Residue #	Residue	$\Delta G_i^{conf} (\text{Ca}^{2+})$	$T\Delta S_i^{conf} (\text{Ca}^{2+})$	$\Delta G_i^{conf} (\text{Mg}^{2+})$	$T\Delta S_i^{conf} (\text{Mg}^{2+})$
50	TYR	-0.6	1.6	-1.1	-4.0
51	GLY	0.0	-0.7	0.1	-0.7
52	LEU	-0.4	-0.5	-1.1	0.4
53	PHE	0.0	1.4	-0.6	-3.2
54	GLN	-0.4	-5.7	-0.5	-3.5
55	ILE	0.2	-1.4	0.5	-0.4
56	ASN	-0.5	-5.2	0.3	2.6
57	ASN	0.0	1.5	0.0	-1.9
58	LYS	-0.7	-0.7	-1.3	-2.4
59	ILE	0.9	0.8	0.6	-0.5
60	TRP	1.4	-0.4	3.6	2.3
86	THR	-0.8	-1.7	-0.9	-1.9
87	ASP	-0.7	1.3	-1.3	-1.4
88	ASP	0.3	2.2	-0.7	-1.6
89	ILE	-0.6	-1.5	0.0	2.3
90	MET	0.0	-0.1	0.1	0.5
91	CYS	0.1	-0.1	0.0	0.0
92	VAL	-1.0	-3.6	-1.2	-5.2
93	LYS	-1.2	-2.8	-0.2	-0.1
94	LYS	-0.9	-4.0	-0.4	0.1
95	ILE	3.4	2.9	-1.0	-2.7
96	LEU	-0.5	-5.3	-0.5	-3.7
97	ASP	0.1	-0.2	-0.2	-0.7
98	LYS	-0.5	-2.8	-0.6	-1.1
99	VAL	-0.6	-2.4	-0.8	-1.4
100	GLY	-0.7	-1.1	-0.8	-0.7
101	ILE	-1.0	-5.7	-0.2	0.2
102	ASN	2.3	5.7	1.2	0.5
103	TYR	0.3	5.0	0.0	-0.1
104	TRP	-1.0	1.2	-1.9	-1.3

Figure Captions:

Fig. S1. The Ramachandran plot of simulated Mg^{2+} -aLA complex.

Fig. S2. The convergence of equilibrium dihedral distributions corresponding to five samples generated randomly from the 80-100 ns trajectory, for a few representative dihedrals in apo-, Ca^{2+} -, Mg^{2+} -aLA.

Fig. S3. The conformational thermodynamic changes (in kJ/mol) at the residues of the MBL upon metal binding are illustrated. (a) $\Delta G_{\text{MBL},i}^{\text{conf}}$, (b) $T\Delta S_{\text{MBL},i}^{\text{conf}}$ of Ca^{2+} -aLA (c) $\Delta G_{\text{MBL},i}^{\text{conf}}$ and (d) $T\Delta S_{\text{MBL},i}^{\text{conf}}$ upon Mg^{2+} binding.

Fig. S4. Representative equilibrium dihedral distributions in native-apo, Ca^{2+} - and Mg^{2+} -aLA: (a) ψ of K79. (b) ψ of D82. (c) ψ of D84. (d) ϕ of D87. (e) ψ of F80. (f) ϕ of L81.

Fig. S5. Representative equilibrium dihedral distributions in native-apo, Ca^{2+} - and Mg^{2+} -aLA: (a) χ_1 and (b) χ_2 of K79. (c) χ_1 of D82. (d) χ_2 of D84. (e) χ_2 of D87. (f) χ_1 of D88.

Fig. S6. Multiple Sequence Alignment (MSA) of aLA sequences from different species. The predicted oleic acid binding residues, H32, W60, I95 and W104 are conserved; however, I59 shows weak conservation across the sequences.

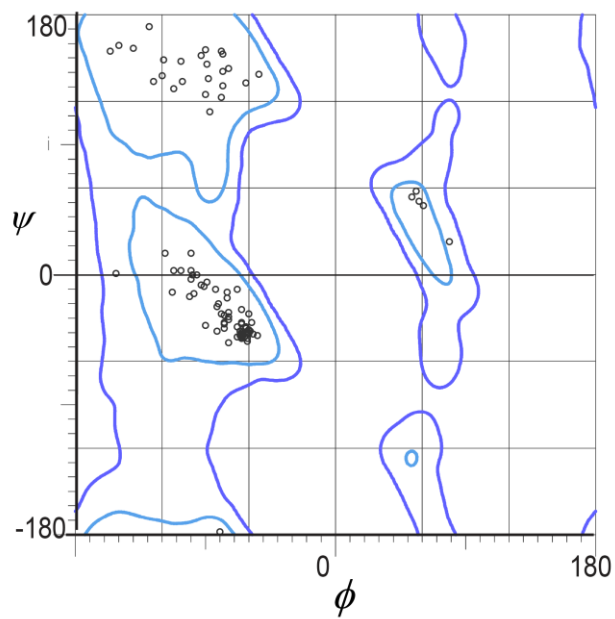


Fig S1

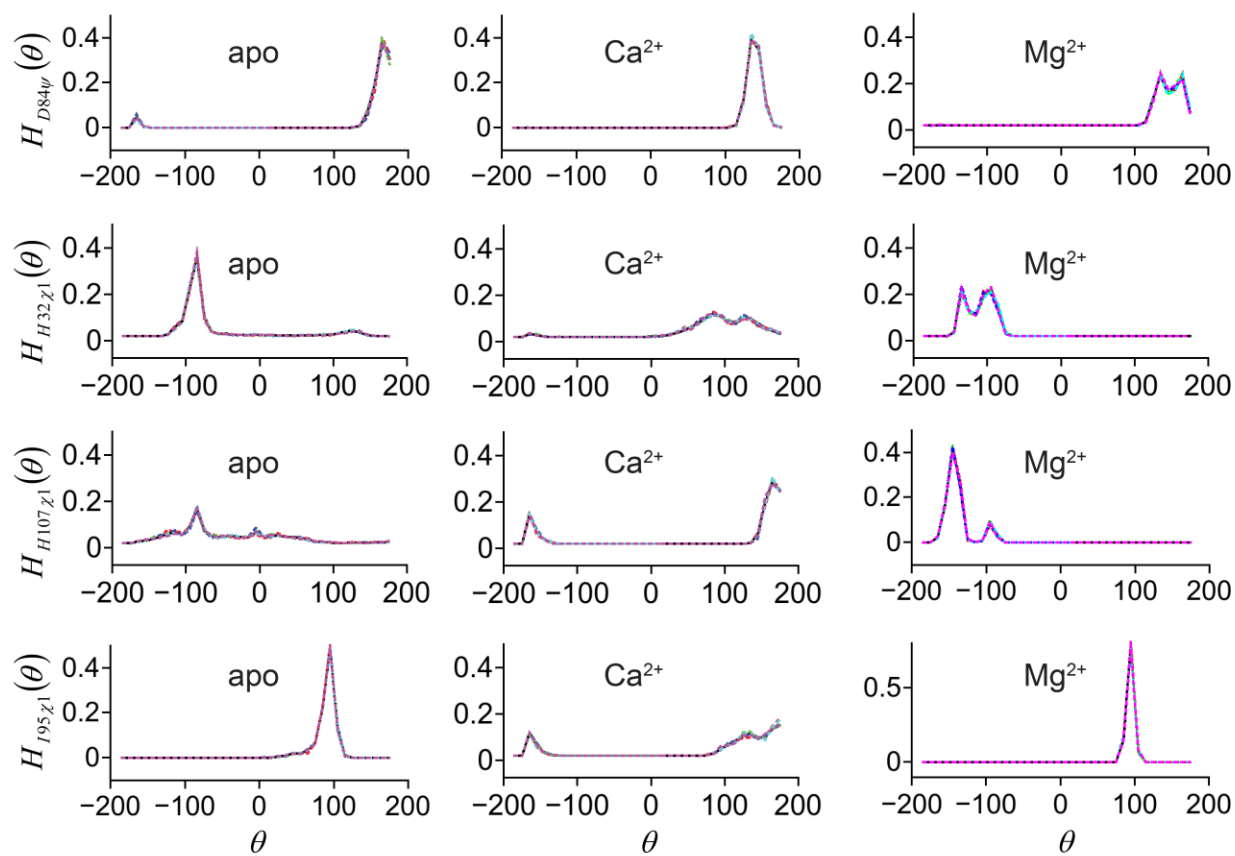


Fig S2

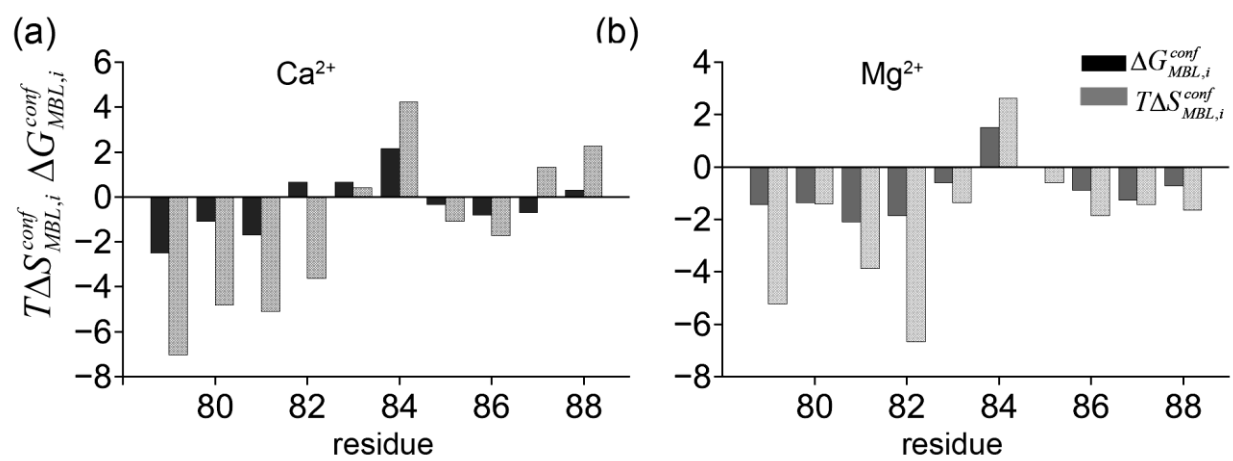


Fig S3

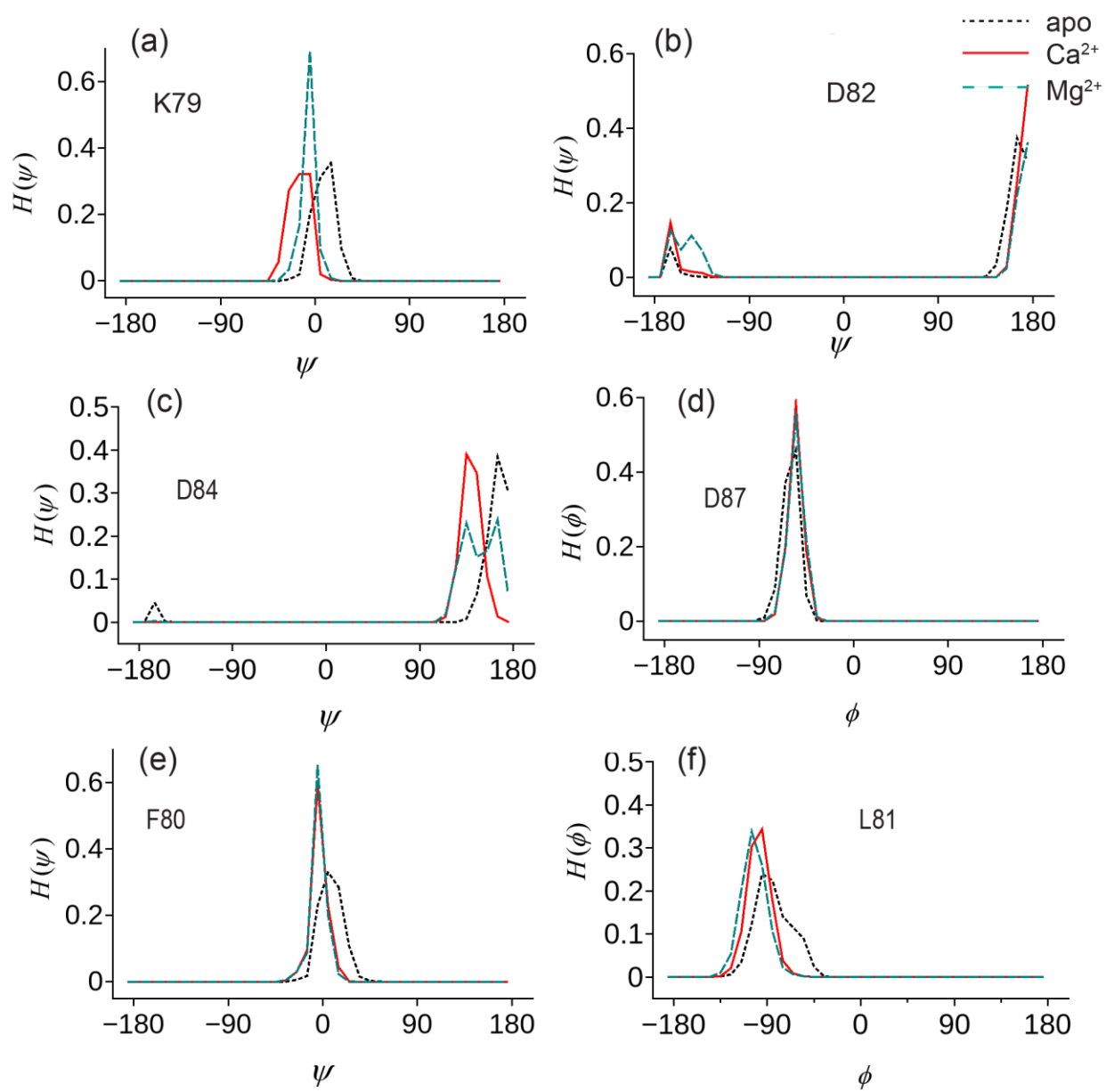


Fig S4

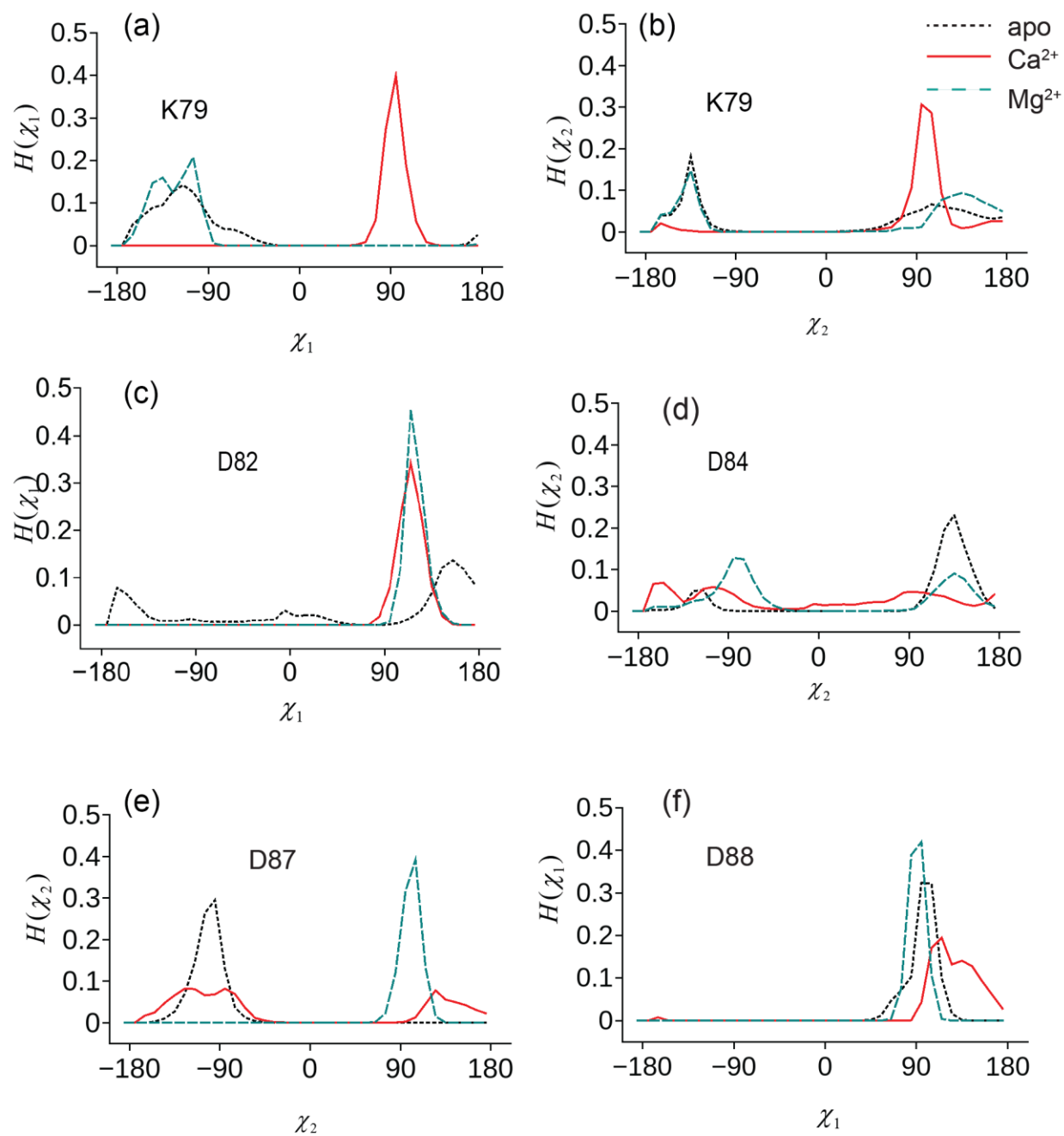


Fig S5

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Bos mutus      --MMSFVSLLLVGIL-FHATQAEQLTKCEVFRELKDLKYGGSVLPWVCTTHTSGYDTQAI VQNNDSTEYGLFQINNIIWCKDDQNPHSSNICNISCDKFLDDDLTDDIMCVKEI LDKVGINTWLAHKALCSEKLDQWLCEKL
Bos grunniens  --MMSFVSLLLVGIL-FHATQAEQLTKCEVFRELKDLKYGGSVLPWVCTTHTSGYDTQAI VQNNDSTEYGLFQINNIIWCKDDQNPHSSNICNISCDKFLDDDLTDDIMCVKEI LDKVGINTWLAHKALCSEKLDQWLCEKL
Bos indicus    --MMSFVSLLLVGIL-FHATQAEQLTKCEVFRELKDLKYGGSVLPWVCTTHTSGYDTQAI VQNNDSTEYGLFQINNIIWCKDDQNPHSSNICNISCDKFLDDDLTDDIMCVKEI LDKVGINTWLAHKALCSEKLDQWLCEKL
Bubalus bubalis--MMSFVSLLLVGIL-FHATQAEQLTKCEVFRELKDLKYGGSVLPWVCTTHTSGYDTQAI VQNNDSTEYGLFQINNIIWCKDDQNPHSSNICNISCDKFLDDDLTDDIMCVKEI LDKVGINTWLAHKALCSEKLDQWLCEKL
Ovis aries     --MMSFVSLLLVGIL-FHATQAEQLTKCEVFRELKDLKYGGSVLPWVCTTHTSGYDTQAI VQNNDSTEYGLFQINNIIWCKDDQNPHSSNICNISCDKFLDDDLTDDIMCVKEI LDKVGINTWLAHKALCSEKLDQWLCEKL
Capra hircus   --MMSFVSLLLVGIL-FHATQAEQLTKCEVFRELKDLKYGGSVLPWVCTTHTSGYDTQAI VQNNDSTEYGLFQINNIIWCKDDQNPHSSNICNISCDKFLDDDLTDDIMCVKEI LDKVGINTWLAHKALCSEKLDQWLCEKL
Cannis familiaris--MMSFVSLLLVSI L-FPAIQAKQFTKCELSQVLKMDMGFGGIALPEWICTIHTSGYDTQTI VNNNGSTGYGLFQISNIIWCKDDQNQLQSRNICDISCDKFLDDDLTDDIMCAKEI LDKEGIDTWLAHKPLCSEKLEQWRCEKL
Mustelo furo   --MMSFISLLLVGIM-FPAIQAKQFTKCELSQVLKMDMGFGGIALPEWICTIHTSGYDTQTI VNNNGSTGYGLFQISNIIWCKDNKILQSRNICDISCDKFLDDDLTDDMTCAKEI LNKEGIDTWLAHKPLCSEKLEQWHCEKL
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Sus scrofa     --MMSFVSLLLVGIL-FPAIQAKQFTKCELSQVLKMDMGYGDITLPEWICTIHTSGYDTKTI VHDNGSTEYGLFQINNIIWCKDNQ- IQSKNICGISCDKFLDDDLTDDMMCAKEI LDNEGIDTWLAHKALCSEKLDQWLCEKM
Equus caballus--KMMSFASLLLVGIL-FSATQAKQFTKCELSQVLKMDMGYKGVTLPEWICTIHTSGYDTQTI VKNNGKTEYGLFQINNIIWCKDNQILQSRNICGISCNKFLDDDLTDDVMCAKEI LDSEGIDTWLAHKPLCSEKLEQWLCEEL
Macaca fascicularis--MRSFVPLFLVGIL-FPAIPAKQFTKCELSQVLKMDIDYGGIALPEFICTMHTSGYDTQAI VESNGSTEYGLFQISNIIWCKSSQVPQSRNICDISCDKFLDDDLTDDIMCAKEI LDIKGIDTWLAHKALCSEKLEQWLCEKL
Macaca mulatta--MRSFVPLFLVGIL-FPAIPAKQFTKCELSQVLKMDIDYGGIALPEFICTMHTSGYDTQAI VESNGSTEYGLFQISNIIWCKSSQVPQSRNICDISCDKFLDDDLTDDIMCAKEI LDIKGIDTWLAHKALCSEKLEQWLCEKL
Gorilla gorilla--MRFFVPLFLVGIL-FPAILAKQFTKCELSQVLKMDIDYGGIALPELICTMHTSGYDTQAI VENNESTEYGLFQISNIIWCKSSQVPQSRNICDISCDKFLDDDLTDDIMCAKEI LDIKGIDTWLAHKALCSEKLEQWLCEKL
Pusa hispida   --MMSFVSLLLVGIM-FPAIQAKQFRKCELSQVLKMDMGFRGIALPKWICTIHTSGYDTQTI VSNNGSTEYGLFQINNIIWCKDNQILQSRNICDISCDKFLDDDLTDDIMCAKEI LDKEGIDTWLAHKPLCSEKLEQWHCEKL
Homo sapiens   --MRFFVPLFLVGIL-FPAILAKQFTKCELSQVLKMDIDYGGIALPELICTMHTSGYDTQAI VENNESTEYGLFQISNIIWCKSSQVPQSRNICDISCDKFLDDDLTDDIMCAKEI LDIKGIDTWLAHKALCSEKLEQWLCEKL
Myotis brandtii--MMSFSLLLVGIL-FPALEAKQFTKCELSQVLKMDMGYGGVTLPEWICNIHTSGYDTQTI VSNNGSTEYGLFQINNIIWCKDNQ- IQSRNICDISCDKFLDDDLTDDMMCAKEI LDSEGIDTWLAHKPLCSEKLEQWLCEKL
Ailuropoda melanoleuca--MMFFVSLLLVGIM-CPAIQAKQFTKCELSQVLKMDMGFGGIALSEWICTIHTSGYDTQTI VNNNGSTEYGLFQINNIIWCKDNQILQSRNICDISCDKFLDDDLTDDIMCAKEI LDKEGIDTWLAHKPLCSEKLEQWHCEKL
Camelus ferus  --MMSLVSLLLVGIL-FPTIQAKQFTKCKLSDELKMDMNGHGGITLAEWICIIHTSGYDTETVVSNNNGNREYGLFQINNIIWCKDNENLQSRNICDISCDKFLDDDLTDDKMAKEI LDKEGIDTWLAHKPLCSEKLEQWQCEKW
Cervus elaphus-----VCTAHTSGYDTQAI VQNDSTEYGLFQINNIIWCKDDQNPHSSNICNISCDKFLDDDLTDDIMCVKEI LDKVGINTWLAHKALCSEKLDQWLCEKL
Balaenoptera musculus---MSFVSLLLVGNL-FHAIQAEQLTKCEVFQRLKDLGYGGVTLPEWVCTVHTSGCDTQTI VNNNGSTEYGLFQINNIIWCKDNHIPS RDI CGISCDKFLDDDLTDDIMCVKEI LDNV-----
Cavia porcellus-----MMSFFPLLLVGIL-FPAVQAKQLTKCALSHELNDLAGYRDTLPEWLCIIHTSGYDTQAI VKNSDHKEYGLFQINDKDFCESSTTVQSRNICDISCDKFLDDDLTDDIMCVKEI LDIKGIDTWLAHKPLCSDKLEQWYCEAQ
Sotalia fluviatilis-----LLLVGIL-FHAVQAEQLTKCELFQRLKDLGYGGVTLPEWVCTVHTSGCDTQTI VNNSDSTEYGLFQINNIIWCKDNQIPS RDI CGISCDKFLDDDLTDDIMCVKEI LDNV-----
Heterocephalus glaber-----KQFTKCELSQNLIDYGYRIALPELICTMHTSGYDTQAI VENNESTEYGLFQISNIIWCKSSQSPQSRNICDITCDKFLDDDLTDDIMCAKEI LDIKGIDTWLAHKALCSEKLEQWLCEKE
Papio cynocephalus--MMSFPFLLLVSI L-FSALQAKQFTKCSLSQELNDLAGYRNITLPEWICIIHTSGYDTQTI IRNNGNTEYGLFQINDKDFCDSSQNLQSRNICDISCDKFLDDDLTDDIMCAKEI LDIKGIDTWLAHKPLCSDKLEQWYCKVL
Delphinus delphis-----LLVGIL-FHAVQAEQXTKCELFQRLKDLGYGGVTLPEWVCTVHTSGCDTQTI VNNNDSTEYGLFQINNIIWCKDNQIPS RDI CGISCDKFLDDDLTDDIMCVKEI LDNV-----
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Camelus dromedarius-----KQFTKCKLSDELKMDMNGHGGITLAEWICIIHTSGYDTETVVSNNNGNREYGLFQINNIIWCKDNENLQSRNICDISCDKFLDDDLTDDKMAKEI LDK-----
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Oryctolagus cuniculus--MMPLVPLLLVSIV-FPGIATQLTRCELTEKLKLDGYRDISMSEWICTLHTSGLDTKITVNNNGSTEYGI FQISDI LKLVSKQNPQSKNICDTPCENFLDDNLTDDVCKAMEI LDKEGIDTWLAHKPLCSENLEQWCKKL
Rattus norvegicus--MMRFVPLFLACI-SLPAFQATEFTKCEVSHAIEDMDGYQGISLLEWTCTVHTSGYDSQAI VKNNGSTEYGLFQISNIIWCKSSEFPESENICDISCDKFLDDELADDIVCAKEI LVAIKGIDTWLAHKPMCSEKLEQWRCEKP
Loxodonta africanaAKMMSFVPLLLVGIL-FPAIQAKQFTKCELSQVLKMDIDYAGITLPEFTCTIHTSGYDTQTI VNNNGSTEYGLFQISNIIWCKDHQIPQSRNICDISCDKFLDDDLTDDMMCAKEI LKSGKIDTW-----
Mus musculus--MMHFVPLFLVLCILSLPAFQATELTCKKVSHAIKDIDYQGISLLEWACVLHTSGYDTQAVVNDNGSTEYGLFQISDI LKLVCKSSEFPESENICDISCDKFLDDELDDDIACAKEI LVAIKGIDTWLAHKPMCSEKLEQWRCEKP
Phocoenoides phocoena-----QAEQLTKCELFQRLKDLGYGGVTLPEWVCTVHTSGCDTQTI VNNNDSTEYGLFQINNIIWCKDNQIPS RDI CGISCDKFLDDDLTDDIM-----
Cricetulus griseus--MMFFIPLILVCIL-FPAIQATQLTKCEVQYAMRMDMGHEGISSLEWTCIHTSGCDTQATVKNNNGSTEYGLFQISNIIWCKSSEIPESENICDISCDKFLDDDLTDDKMAKEI LVAIKGIDTW-----

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Fig S6