

Supplementary Information for

“Salt Bridges in the Prion Protein: Implications for Stability and Misfolding”

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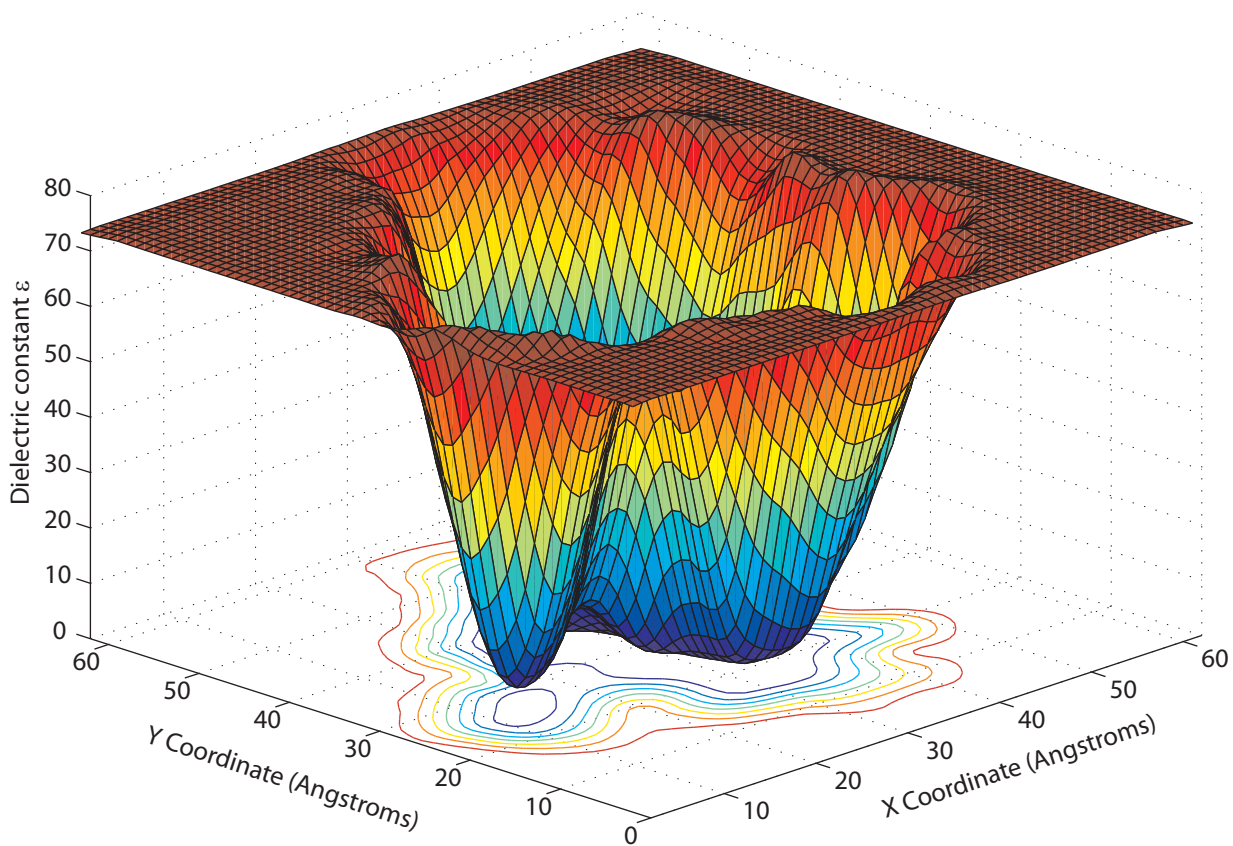


Figure 1: Local heterogeneous dielectric map of human PrP as calculated by mesoscopic dielectric theory (Guest et al., 2009). This is a slice taken through the geometric centre of the molecule in the X-Y plane.

Table 1: Salt bridges in 13 structures of the prion protein with a repulsive or attractive energy greater than 3 kJ/mol. The number of NMR conformers in which the participating charged groups are within a distance of 12Å is n . $E_{sb}(1)$ is the salt bridge energy calculated according to Equation 2 with the heterogeneous dielectric theory, and $E_{sb}(2)$ is the same energy calculated with a constant protein dielectric of 4. The standard deviation of salt bridge energy over the structures in the NMR ensemble containing the salt bridge is given by δE_{sb} and as a fraction of the total salt bridge energy by $\frac{\delta E_{sb}}{E_{sb}}$.

PDB	Species	Residues Involved	n (/20)	$E_{sb}(1)$ (kJ/mol)	$\delta E_{sb}(1)$ (kJ/mol)	$\frac{\delta E_{sb}(1)}{E_{sb}(1)}$	$E_{sb}(2)$ (kJ/mol)	$\delta E_{sb}(2)$ (kJ/mol)	$\frac{\delta E_{sb}(2)}{E_{sb}(2)}$
1DX0	Bovine	147 151	20	-6.4	3.5	0.54	-15.2	12.6	0.83
1DX0	Bovine	164 178	20	-3.1	1.9	0.60	-6.2	6.3	1.02
1DX0	Bovine	156 196	20	-6.0	2.8	0.47	-12.1	8.7	0.72
1DX0	Bovine	156 202	20	-7.8	5.5	0.71	-13.0	10.7	0.82
1DX0	Bovine	200 204	20	-3.1	1.4	0.46	-4.1	5.9	1.44
1DX0	Bovine	208 211	20	-5.1	1.9	0.38	-10.4	10.2	0.98
1FKC	Human E200K	164 167	19	-3.0	1.1	0.38	-3.2	2.0	0.64
1FKC	Human E200K	164 178	20	-4.6	1.0	0.23	-7.0	3.8	0.55
1FKC	Human E200K	156 196	20	-3.1	0.7	0.22	-3.2	1.5	0.46
1FKC	Human E200K	156 202	20	-3.2	0.7	0.21	-4.6	1.5	0.34
1FKC	Human E200K	146 204	20	-5.2	1.6	0.31	-8.6	5.1	0.59
1FKC	Human E200K	204 208	20	3.4	0.5	0.15	3.7	0.9	0.25
1FKC	Human E200K	208 211	20	-4.2	0.6	0.14	-5.2	1.8	0.33
1QLZ	Human WT	144 147	19	4.4	0.8	0.19	5.4	1.7	0.32
1QLZ	Human WT	144 151	16	-3.5	1.0	0.29	-4.1	2.5	0.60
1QLZ	Human WT	147 151	20	-9.9	3.8	0.39	-30.4	16.2	0.53
1QLZ	Human WT	148 152	19	-5.0	1.2	0.25	-14.6	7.5	0.51
1QLZ	Human WT	164 168	12	-3.3	1.7	0.53	-4.9	6.5	1.32
1QLZ	Human WT	164 178	15	-4.8	5.9	1.23	-8.2	11.8	1.44
1QLZ	Human WT	156 196	20	-5.1	1.6	0.32	-11.8	11.4	0.96
1QLZ	Human WT	146 204	18	-3.0	0.9	0.29	-3.4	1.4	0.43
1QLZ	Human WT	200 204	20	-3.7	1.5	0.39	-6.5	5.5	0.85
1QLZ	Human WT	204 207	15	-3.1	1.4	0.45	-3.7	5.9	1.58
1QLZ	Human WT	208 211	20	-5.8	1.8	0.30	-14.9	9.9	0.66
1QLZ	Human WT	167 228	11	-13.1	0.9	0.07	-75.5	5.7	0.07
1QLZ	Human WT	221 228	12	-13.7	1.0	0.07	-76.6	5.7	0.07
1U3M	Chicken	150 153	20	-5.0	1.3	0.25	-10.1	7.3	0.72
1U3M	Chicken	155 159	20	-3.8	0.7	0.19	-5.2	2.3	0.43
1U3M	Chicken	171 185	20	-5.4	1.6	0.29	-12.4	6.7	0.54
1U3M	Chicken	159 215	20	5.2	0.8	0.15	6.2	2.7	0.44
1U3M	Chicken	163 219	20	-8.4	4.3	0.51	-20.6	15.9	0.77
1U3M	Chicken	217 221	20	-5.7	1.1	0.20	-15.3	7.6	0.50
1U3M	Chicken	225 229	20	-4.1	1.8	0.44	-5.9	7.7	1.30
1U3M	Chicken	139 237	20	-4.1	2.0	0.48	-6.0	6.0	0.99
1U5L	Turtle	143 146	20	-3.3	0.7	0.21	-5.8	2.3	0.40
1U5L	Turtle	165 169	18	-3.0	2.2	0.74	-5.0	7.1	1.42
1U5L	Turtle	164 179	20	-5.1	2.8	0.55	-7.9	7.9	1.00
1U5L	Turtle	175 179	20	-4.2	2.8	0.68	-6.9	7.3	1.06

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PDB	Species	Residues Involved	n (/20)	$E_{sb}(1)$ (kJ/mol)	$\delta E_{sb}(1)$ (kJ/mol)	$\frac{\delta E_{sb}(1)}{E_{sb}(1)}$	$E_{sb}(2)$ (kJ/mol)	$\delta E_{sb}(2)$ (kJ/mol)	$\frac{\delta E_{sb}(2)}{E_{sb}(2)}$
1U5L	Turtle	156 202	16	-3.1	2.5	0.79	-4.2	6.8	1.63
1U5L	Turtle	146 204	20	-5.7	1.8	0.32	-15.1	10.8	0.71
1U5L	Turtle	191 207	20	-4.6	2.6	0.56	-7.5	8.7	1.15
1XU0	Frog	148 152	20	-3.3	1.3	0.39	-4.7	6.2	1.31
1XU0	Frog	169 176	19	-3.1	1.6	0.52	-5.6	6.6	1.17
1XU0	Frog	175 179	20	-3.1	1.3	0.43	-4.8	7.0	1.45
1XU0	Frog	169 180	19	3.0	1.4	0.46	4.1	2.8	0.69
1XU0	Frog	188 192	20	-9.4	1.8	0.19	-19.3	6.3	0.33
1XU0	Frog	197 206	20	-5.3	3.4	0.63	-10.2	11.9	1.16
1XU0	Frog	197 210	19	4.7	2.3	0.50	7.7	6.5	0.85
1XU0	Frog	206 210	20	-21.4	6.7	0.32	-42.2	15.9	0.38
1XU0	Frog	163 221	20	-9.4	6.1	0.64	-16.9	13.6	0.80
1XU0	Frog	221 224	20	-5.0	3.6	0.72	-11.5	12.1	1.05
1XYJ	Cat	147 151	20	-11.6	3.1	0.27	-40.9	13.4	0.33
1XYJ	Cat	164 178	20	-4.7	2.5	0.54	-11.7	11.9	1.01
1XYJ	Cat	156 202	20	-11.8	5.3	0.44	-21.5	15.0	0.70
1XYJ	Cat	146 204	20	-3.9	2.0	0.51	-5.7	8.7	1.51
1XYJ	Cat	200 204	20	-3.5	0.9	0.26	-4.0	3.4	0.85
1XYJ	Cat	208 211	20	-5.8	3.0	0.52	-11.8	11.3	0.96
1XYK	Dog	144 147	20	3.1	1.0	0.32	3.9	2.0	0.52
1XYK	Dog	144 148	20	-4.1	2.1	0.51	-9.3	9.1	0.97
1XYK	Dog	147 151	20	-8.4	4.5	0.54	-22.5	16.5	0.73
1XYK	Dog	148 152	20	-3.1	1.2	0.39	-5.3	5.8	1.09
1XYK	Dog	164 178	20	-3.6	2.2	0.60	-7.4	9.3	1.25
1XYK	Dog	136 202	2	-4.0	0.8	0.19	-6.1	0.8	0.12
1XYK	Dog	200 204	20	-3.1	1.2	0.40	-3.8	4.7	1.22
1XYK	Dog	204 207	20	-3.8	1.4	0.38	-5.6	6.2	1.11
1XYK	Dog	207 211	20	4.0	0.7	0.16	4.8	1.2	0.26
1XYK	Dog	208 211	20	-5.3	2.5	0.48	-11.5	11.3	0.98
1XYQ	Pig	147 151	20	-4.3	1.3	0.29	-6.0	4.1	0.69
1XYQ	Pig	148 151	20	3.5	0.3	0.09	4.1	1.0	0.24
1XYQ	Pig	164 178	20	-5.9	1.5	0.25	-11.9	7.0	0.59
1XYQ	Pig	200 204	20	-4.0	1.3	0.33	-7.0	6.4	0.91
1XYQ	Pig	204 207	20	-3.3	1.2	0.37	-3.0	2.1	0.71
1XYQ	Pig	207 211	20	3.4	0.8	0.23	4.0	1.2	0.30
1XYQ	Pig	208 211	20	-5.9	2.5	0.42	-9.8	12.1	1.24
1XYQ	Pig	220 223	19	-3.8	1.2	0.33	-3.6	2.4	0.67
1XYX	Mouse	144 147	12	3.2	0.8	0.25	3.2	1.3	0.41
1XYX	Mouse	147 151	16	-5.1	2.0	0.38	-9.2	7.9	0.85
1XYX	Mouse	148 152	12	-3.8	1.7	0.44	-7.8	9.3	1.18
1XYX	Mouse	164 178	18	-5.4	2.4	0.45	-10.8	11.0	1.02
1XYX	Mouse	156 196	13	-3.7	1.0	0.28	-4.2	2.5	0.61
1XYX	Mouse	194 196	16	-3.5	0.7	0.21	-3.0	0.7	0.23
1XYX	Mouse	156 202	20	-10.4	3.9	0.37	-18.9	8.8	0.46
1XYX	Mouse	146 204	20	-9.7	2.5	0.26	-27.2	11.9	0.44
1XYX	Mouse	146 208	10	-3.2	0.3	0.10	-3.8	0.9	0.23
1XYX	Mouse	204 208	18	3.2	0.4	0.13	3.6	0.9	0.24
1XYX	Mouse	207 211	10	3.5	0.5	0.15	4.0	0.8	0.20

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PDB	Species	Residues Involved	n (/20)	$E_{sb}(1)$ (kJ/mol)	$\delta E_{sb}(1)$ (kJ/mol)	$\frac{\delta E_{sb}(1)}{E_{sb}(1)}$	$E_{sb}(2)$ (kJ/mol)	$\delta E_{sb}(2)$ (kJ/mol)	$\frac{\delta E_{sb}(2)}{E_{sb}(2)}$
1XYX	Mouse	208 211	10	-4.4	1.1	0.24	-6.2	5.8	0.94
1XYX	Mouse	167 221	2	4.5	0.1	0.03	5.7	0.8	0.14
1XYX	Mouse	227 230	10	-3.7	1.9	0.51	-6.9	7.9	1.15
2K1D	Human D178N	144 147	20	3.3	0.4	0.13	3.6	0.9	0.23
2K1D	Human D178N	147 151	20	-3.7	1.1	0.30	-5.1	3.2	0.63
2K1D	Human D178N	164 185	19	3.3	1.5	0.45	3.7	2.8	0.78
2K1D	Human D178N	156 194	20	3.4	1.4	0.40	4.9	4.1	0.85
2K1D	Human D178N	156 202	16	-3.5	1.9	0.53	-5.5	5.1	0.93
2K1D	Human D178N	208 211	19	-3.6	1.2	0.32	-3.0	1.4	0.46
2KFL	Wallaby	147 151	20	-5.4	2.7	0.51	-14.6	11.3	0.77
2KFL	Wallaby	148 152	20	-3.6	1.4	0.40	-7.9	7.6	0.96
2KFL	Wallaby	164 178	20	-4.8	1.9	0.39	-8.3	7.8	0.93
2KFL	Wallaby	178 185	19	-3.3	0.9	0.27	-3.7	1.6	0.42
2KFL	Wallaby	156 196	20	-10.9	3.5	0.32	-29.6	15.1	0.51
2KFL	Wallaby	156 202	20	-13.3	11.7	0.88	-18.0	16.2	0.90
2KFL	Wallaby	196 202	20	4.9	0.7	0.13	7.4	1.6	0.22
2KFL	Wallaby	200 204	20	-4.6	1.1	0.24	-7.4	5.5	0.74
2KFL	Wallaby	204 207	20	-4.2	0.7	0.17	-4.4	1.2	0.28
2KFL	Wallaby	146 208	20	-3.7	0.3	0.09	-4.6	0.9	0.20
2KFL	Wallaby	208 211	20	-3.8	0.9	0.25	-4.8	5.0	1.04

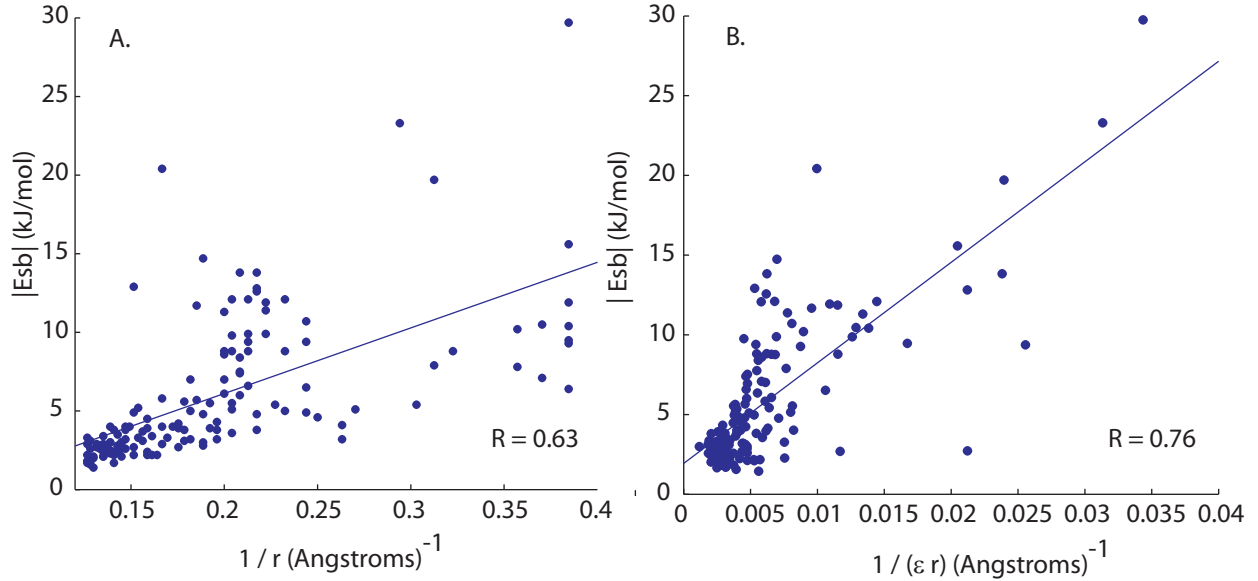


Figure 2: A. Absolute salt bridge energies as a function of the reciprocal of the separation distance between the participating charged groups. B. Absolute salt bridge energies as a function of the reciprocal of the separation distance multiplied by the dielectric constant ϵ at the midpoint of the line joining the charged groups. In a homogeneous dielectric environment, the energy of a salt bridge connecting charges q_A and q_B would be given by $E_{sb} = \frac{1}{4\pi\epsilon_0} \frac{q_A q_B}{\epsilon r}$. Including $1/\epsilon$ in the fit in addition to $1/r$ improves the correlation from 0.63 to 0.76, but even taken together the quantities ϵ and r do not reliably predict salt bridge energies. Thus a direct calculation of energies from the full heterogeneous dielectric map is necessary to obtain accurate results.

Table 2: Total electrostatic energies in kJ/mol for every residue in a selection of 13 prion protein structures at pH 7. The energies are calculated according to Equation 3.

Res	2KFL Wallaby	1QLX Human WT	1FKC Human E200K	2K1D Human D178N	1DX0 Bovine	1U3M Chicken	1U5L Turtle	1XU0 Frog	1XYJ Cat	1XYK Dog	1XYQ Pig	1XYX Mouse	1AG2 Mouse			
120	S	-11					S	-7								
121	V	-9					V	-2								
122	V	-2					V	-4								
123	G	-1					G	0								
124	G	-1					G	0								
125	L	-19				L	-7						L	-6		
126	G	-1	G	-1	G	-1	G	-1	G	-2	L	-5	L	-23	L	-6
127	G	-1	G	-1	G	-2	G	0	S	-7	G	-1	G	-1	G	-1
128	Y	-42	Y	-32	Y	-28	Y	-18	Y	-52	Y	-2	Y	-23	Y	-27
129	M	-11	M	-9	M	-8	V	-6	M	-7	V	-2	A	-2	M	-8
130	L	-15	L	-10	L	-10	L	-4	L	-12	G	-1	L	-9	L	-14
131	G	-2	G	-5	G	-5	G	-1	G	-2	G	0	G	-2	G	-4
132	S	-21	S	-18	S	-16	S	-7	S	-11	L	-3	S	-11	N	-13
133	A	-1	A	-3	A	-1	A	-1	A	-1	A	-1	A	-1	A	-3
134	M	-8	M	-6	M	-24	M	-9	M	-30	G	0	M	-10	V	-22
135	S	-8	S	-8	S	-14	S	-9	S	-15	Y	-10	S	-16	G	-11
136	R	-382	R	-151	R	-140	R	-55	R	-63	A	-3	G	-1	R	-119
137	P	-8	P	-10	P	-15	P	-4	P	-7	M	-7	M	-16	M	-8
138	V	-6	I	-11	I	-10	I	-5	L	-7	G	-3	R	-35	S	-4
139	M	-15	I	-20	I	-18	I	-37	I	-43	R	-59	M	-7	Y	-26
140	H	-23	H	-25	H	-27	H	-20	V	-13	N	-14	Q	-13	H	-15
141	F	-29	F	-18	F	-47	F	-24	F	-32	M	-13	F	-23	F	-19
142	G	1	G	1	G	-1	G	0	G	0	S	-16	D	-26	N	0
143	N	-17	S	-9	S	-19	S	-10	S	-26	G	-1	R	-32	N	-15
144	E	-46	D	-36	D	-39	D	-40	D	-36	M	-10	P	-4	P	-45
145	Y	-13	Y	-11	Y	-12	Y	-10	Y	-11	N	-11	E	-29	M	-18
146	E	-40	E	-39	E	-80	E	-43	E	-78	Y	-40	E	-90	E	-45
147	D	-75	D	-196	D	-70	D	-61	D	-91	H	-14	R	-76	S	-86
148	R	-54	R	-34	R	-48	R	-59	R	-34	F	-12	Q	-26	R	-70
149	Y	-52	Y	-20	Y	-42	Y	-23	Y	-27	D	-32	W	-57	Y	-106
150	Y	-86	Y	-171	Y	-132	Y	-68	Y	-143	R	-42	W	-81	Y	-114
151	R	-97	R	-43	R	-55	R	-43	R	-69	P	-2	N	-47	N	-46
152	E	-39	E	-31	E	-39	E	-57	E	-31	D	-33	E	-45	D	-78
153	N	-53	N	-25	N	-49	N	-31	N	-30	E	-33	N	-36	Y	-158
154	Q	-25	M	-24	M	-16	M	-4	M	-8	Y	-17	S	-10	Y	-13
155	Y	-17	H	-19	H	-27	H	-19	H	-18	R	-41	N	-15	N	-21
156	R	-128	R	-29	R	-41	R	-60	R	-97	W	-36	R	-77	Q	-242
157	Y	-145	Y	-75	Y	-148	Y	-18	Y	-140	W	-52	Y	-9	M	-228
158	P	-16	P	-12	P	-9	P	-6	P	-37	S	-13	P	-9	P	-12
159	N	-21	N	-32	N	-52	N	-14	N	-38	E	-54	N	-59	N	-22
160	Q	-23	Q	-34	Q	-18	Q	-24	Q	-16	N	-56	Q	-27	R	-24
161	V	-63	V	-53	V	-64	V	-26	V	-54	S	-20	V	-34	V	-49
162	M	-19	Y	-25	Y	-25	Y	-34	Y	-32	A	-1	Y	-29	Y	-50
163	Y	-59	Y	-51	Y	-40	Y	-31	Y	-39	R	-30	Y	-42	R	-30
164	R	-59	R	-99	R	-63	R	-62	R	-74	Y	-15	K	-30	P	-182
165	P	-16	P	-5	P	-5	P	-8	P	-5	P	-5	E	-47	M	-4
166	I	-25	M	-19	M	-21	M	-2	V	-13	N	-48	Y	-152	Y	-34
167	D	-34	D	-41	D	-58	D	-48	D	-31	R	-67	N	-12	R	-39
168	Q	-20	E	-31	E	-39	E	-70	Q	-11	V	-47	D	-34	G	-20
169	Y	-137	Y	-16	Y	-8	Y	-7	Y	-8	Y	-8	Y	-31	R	-10
170	G	-1	S	-25	S	-27	S	-9	S	-18	Y	-52	S	-10	E	-10
171	S	-12	N	-20	N	-17	N	-13	N	-19	R	-36	V	-22	Y	-13
172	Q	-22	Q	-11	Q	-21	Q	-15	Q	-30	D	-29	P	-3	V	-70
173	N	-15	N	-11	N	-16	N	-13	N	-18	Y	-131	E	-36	S	-17
174	S	-14	N	-12	N	-14	N	-17	N	-15	S	-7	G	-1	E	-34

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Res	2KFL Wallaby	1QLX Human WT	1FKC Human E200K	2K1D Human D178N	1DX0 Bovine	1U3M Chicken	1U5L Turtle	1XU0 Frog	1XYJ Cat	1XYK Dog	1XYQ Pig	1XYX Mouse	1AG2 Mouse
175	F -79	F -53	F -54	F -28	F -45	S -12	R -134	D -36	F -30	F -18	F -73	F -24	F -79
176	V -18	V -9	V -19	V -19	V -14	P -3	F -68	R -47	V -6	V -14	V -46	V -10	V -10
177	H -17	H -17	H -16	H -16	H -18	V -4	V -9	F -56	H -14	R -29	H -15	H -11	H -17
178	D -69	D -96	D -63	D -32	D -56	P -3	R -39	V -10	D -42	D -52	D -39	D -41	D -78
179	C -14	C -15	C -12	C -11	C -13	Q -24	D -127	R -46	C -6	C -8	C -11	C -9	C -16
180	V -22	V -11	V -12	V -18	V -9	D -31	C -13	D -60	V -16	V -15	V -38	V -9	V -6
181	N -19	N -14	N -18	N -17	N -14	V -3	V -21	C -11	N -13	N -20	N -24	N -9	N -19
182	I -30	I -29	I -14	I -31	I -18	F -60	N -19	Y -13	I -26	I -9	I -26	I -10	I -31
183	T -212	T -197	T -172	T -158	T -193	V -20	I -36	N -18	T -166	T -157	T -165	T -174	T -219
184	V -14	I -34	I -37	I -22	V -14	A -1	T -178	M -15	V -41	V -8	V -48	I -12	I -10
185	K -21	K -19	K -23	K -29	K -23	D -37	V -31	S -133	R -50	K -16	K -21	K -18	K -18
186	Q -51	Q -23	Q -26	Q -20	E -88	C -13	T -19	V -9	Q -20	Q -58	Q -24	Q -28	Q -33
187	H -29	H -37	H -43	H -34	H -41	F -22	E -59	T -26	H -26	H -76	H -19	H -50	H -39
188	T -29	T -22	T -34	T -56	T -34	N -15	Y -20	E -84	T -66	T -11	T -56	T -18	T -24
189	T -21	V -4	V -6	V -4	V -4	I -14	K -13	Y -107	V -7	V -3	V -8	V -3	V -4
190	T -16	T -18	T -11	T -12	T -26	T -133	I -29	I -13	T -18	T -17	T -26	T -15	T -28
191	T -19	T -23	T -19	T -73	T -99	V -20	D -149	I -8	T -22	T -31	T -51	T -54	T -26
192	T -49	T -51	T -50	T -33	T -16	T -18	P -4	K -21	T -27	T -8	T -20	T -19	T -26
193	T -11	T -17	T -21	T -8	T -16	E -43	N -15	P -4	T -13	T -8	T -13	T -10	T -15
194	K -16	K -19	K -15	K -16	K -20	Y -43	E -50	A -2	K -14	K -25	K -19	K -18	K -16
195	G 0	G 0	G 0	G 0	G 0	S -8	N -36	E -29	G 0	G 0	G 0	G 0	G -1
196	E -53	E -30	E -41	E -37	E -48	I -14	Q -13	G 0	E -83	E -67	E -46	E -40	E -78
197	N -22	N -12	N -18	N -18	N -13	G -1	N -13	K -19	N -11	N -13	N -11	N -19	N -18
198	F -34	F -10	F -34	F -11	F -42	P -2	V -19	N -17	F -24	F -21	F -16	F -35	F -23
199	T -37	T -29	T -55	T -32	T -16	A -2	T -14	N -22	T -17	T -34	T -16	T -30	T -63
200	E -35	E -37	K -17	E -33	E -30	A -1	Q -17	S -8	E -31	E -31	E -33	E -36	E -33
201	T -34	T -20	T -25	T -20	T -14	K -9	V -6	E -27	T -14	T -25	T -13	T -23	T -25
202	D -304	D -137	D -290	D -122	D -235	K -14	E -104	L -14	D -123	D -340	D -86	D -161	D -372
203	I -9	V -5	V -5	V -8	I -7	N -14	V -18	N -16	M -8	M -5	V -7	V -6	V -7
204	K -12	K -9	K -12	K -8	K -7	T -12	R -46	Q -15	K -15	K -11	K -13	K -12	K -11
205	I -58	M -20	M -40	M -13	M -42	S -10	V -74	L -14	I -39	I -70	M -15	M -40	M -18
206	M -49	M -37	M -46	M -17	M -39	E -27	M -28	D -119	M -24	M -34	I -42	M -47	M -32
207	E -61	E -44	E -36	E -60	E -42	A -2	K -23	T -14	E -51	E -36	E -69	E -39	E -45
208	R -35	R -35	R -32	R -73	R -41	V -4	Q -16	T -13	R -58	R -30	R -45	R -32	R -27
209	V -86	V -66	V -58	V -30	V -56	A -1	V -60	V -51	V -50	V -41	V -84	V -47	V -63
210	V -80	V -73	V -73	V -55	V -75	A -3	I -81	K -95	V -67	V -61	V -82	V -66	V -86
211	E -52	E -41	E -42	E -59	E -44	A -2	Q -16	S -11	E -53	E -47	E -94	E -37	E -56
212	Q -24	Q -17	Q -21	Q -104	Q -20	N -18	E -46	Q -19	Q -33	Q -23	Q -25	Q -14	Q -23
213	M -31	M -27	M -38	M -30	M -41	Q -13	M -31	I -82	M -36	M -26	M -35	M -36	M -38
214	C -14	C -12	C -16	C -13	C -14	T -12	C -12	I -54	C -8	C -12	C -11	C -7	C -19
215	I -7	I -7	I -17	I -28	I -10	E -44	M -5	R -40	V -10	V -7	I -24	V -5	V -33
216	T -17	T -24	T -57	T -51	T -20	V -9	Q -29	E -50	T -23	T -24	T -30	T -28	T -50
217	Q -66	Q -53	Q -82	Q -21	Q -54	E -25	Q -36	M -52	Q -43	Q -53	Q -60	Q -45	Q -46
218	Y -34	Y -23	Y -51	Y -91	Y -22	M -3	Y -22	C -14	Y -40	Y -37	Y -48	Y -22	Y -122
219	Q -17	E -37	E -44	E -26	Q -15	E -55	Q -28	I -10	Q -19	Q -17	Q -27	Q -15	Q -32
220	N -25	R -53	R -77	R -33	R -56	N -13	Q -18	T -23	K -16	K -15	K -16	K -20	K -16
221	E -220	E -92	E -77	E -88	E -91	K -9	Y -33	E -196	E -42	E -43	E -58	E -53	E -43
222	Y -18	S -19	S -40	S -14	S -11	V -29	Q -56	Y -64	S -52	S -25	Y -16	S -29	S -43
223	Q -22	Q -15	Q -17	Q -13	Q -17	V -19	L -8	R -41	E -39	E -43	E -79	Q -13	Q -16
224	A -2	A -2	A -2	A -2	A -3	T -24	A -2	R -63	A -2	A -1	A -2	A -2	A -2
225	A -2	Y -16	Y -26	Y -17	Y -21	K -15	V -15	G 0	Y -11	Y -11	Y -11	Y -11	Y -15
226	Q -21	Y -9	Y -22	Y -9	Y -13	V -16	V -16		Y -35	Y -11	A -1	Y -11	
227	R -46	Q -14	Q -24	Q -14		I -56			Q -18	Q -22	Q -13	D -29	
228	Y -9		R -41	R -30		R -47			R -28	R -33	R -37	G 0	
229	Y -11		G 0	G 0		E -47			R -29	G 1	G -1	R -39	
230		S -8	S -9			M -41			A -1	A -1	A -1	R -26	
231						C -9						S -8	
232						V -12							
233						Q -22							
234						Q -67							
235						Y -41							

Continued...

Res	2KFL Wallaby	1QLX Human WT	1FKC Human E200K	2K1D Human D178N	1DX0 Bovine	1U3M Chicken	1U5L Turtle	1XU0 Frog	1XYJ Cat	1XYK Dog	1XYQ Pig	1XYX Mouse	1AG2 Mouse
236						R	-35						
237						E	-55						
238						Y	-22						
239						R	-36						
240						L	-3						
241						A	-1						

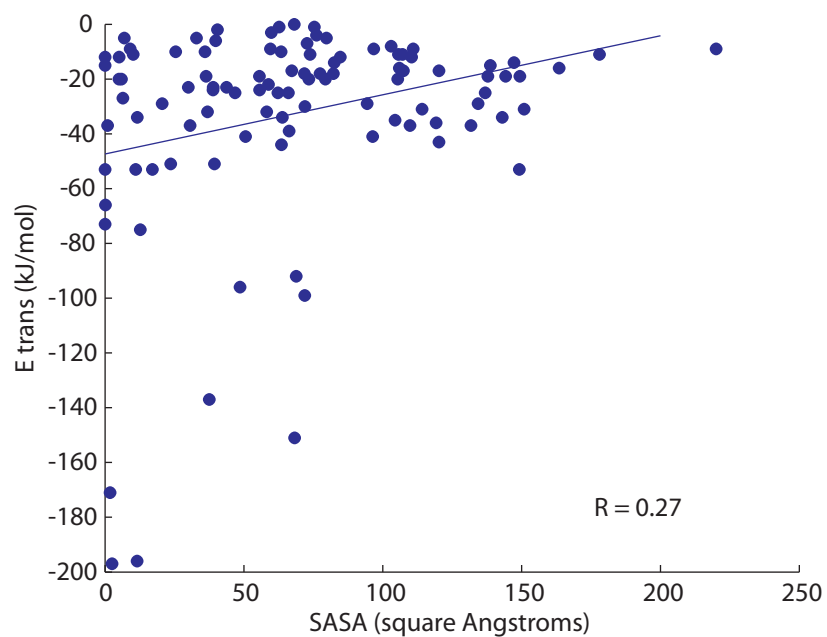


Figure 3: Total electrostatic energies of side chains in human PrP calculated from Equation 3 plotted as a function of solvent-accessible surface area (SASA), a statistic related to burial in the core of the protein. There is a weak but significant ($p = 0.01$) correlation between the energy and SASA, but SASA alone is a poor proxy for the electrostatic energy.

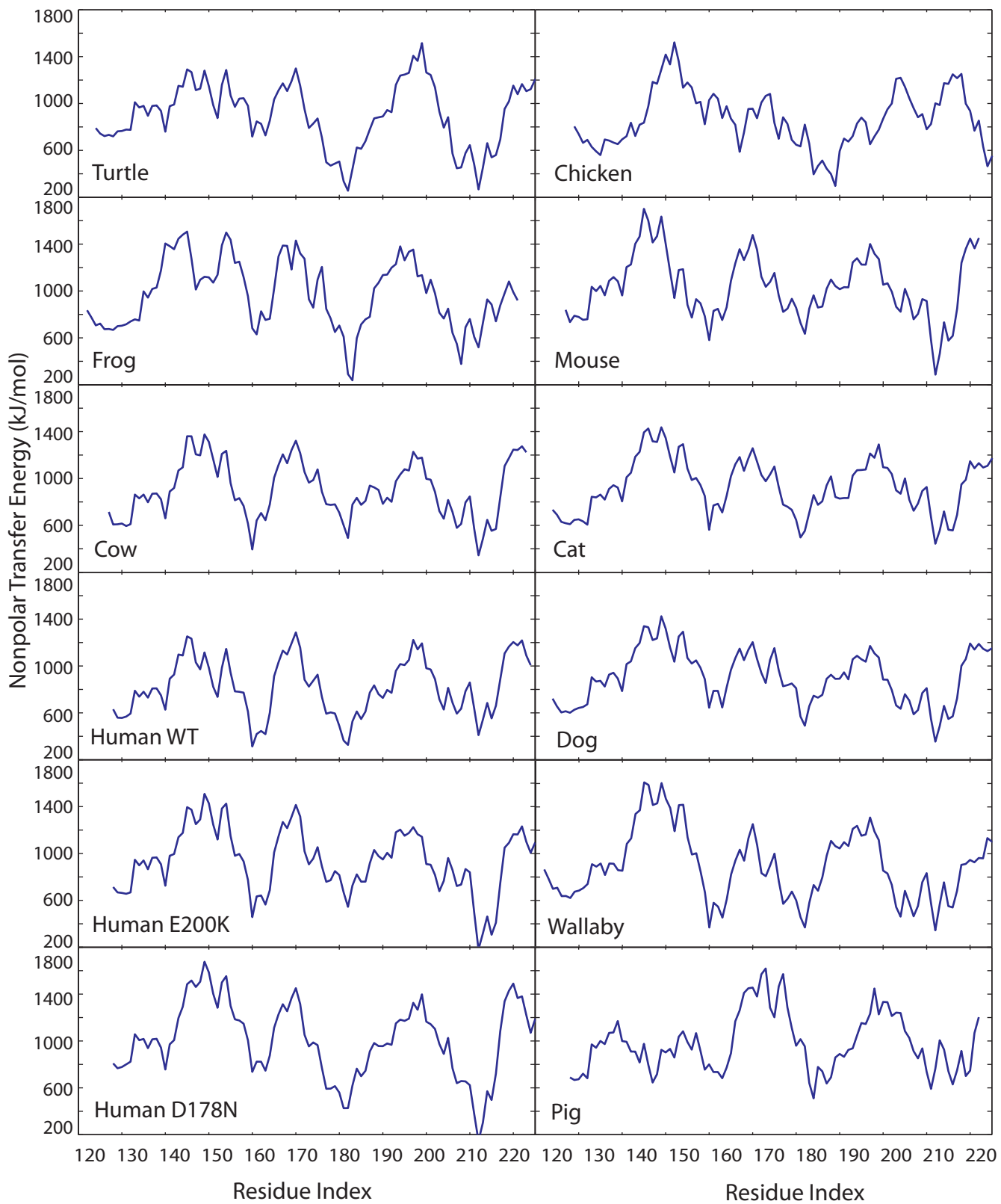


Figure 4: Hydrophobic transfer energies from Equation 4 for seven residue contiguous segments of 12 prion protein structures. The residue index gives the position of the middle residue in each segment.