

SUPPORTING INFORMATION

Insights into the Molecular Flexibility of Theta-defensins by NMR

Relaxation Analysis.

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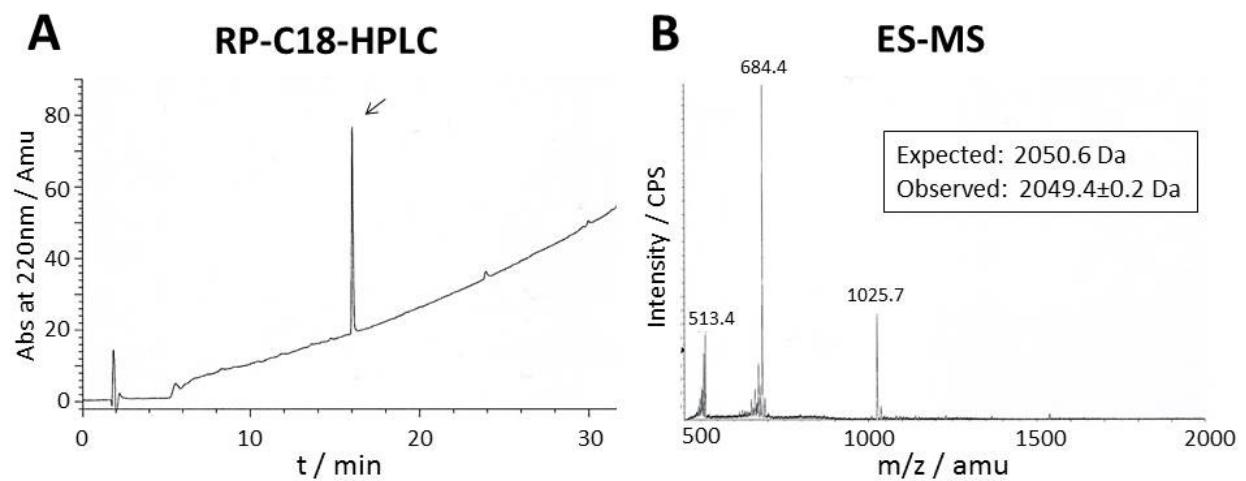
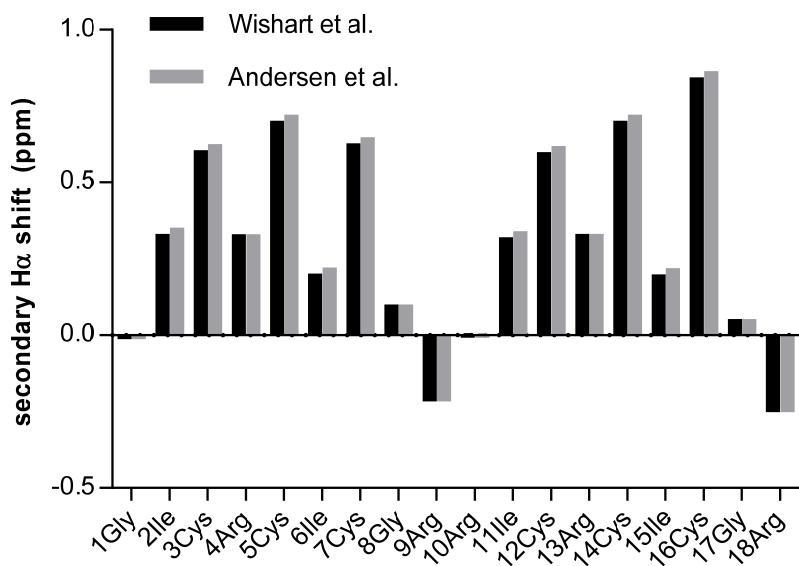


Figure S1. Characterization of ^{15}N -labeled HTD-2. Reverse-phase HPLC analysis (A) and ES-MS analysis (B) of purified ^{15}N -labeled HTD-2.

Supplementary Figure S2: Secondary H α chemical shifts of HTD-2



Supplementary Table S1. Forward (p5) and reverse (p3) 5'-phosphorylated oligonucleotides used to clone the different TEV-HTD2-intein linear precursors into the pTXB1 expression plasmid.

Peptide		Oligonucleotide Sequence
HTD-2	p5	5'-TATGGAAAACCTGTACTTCCAGTGCATCTGCGGTCGTAC TGCCGTTGCATCTGCGGTCGTGGTATCTGCCGT-3'
	p3	5'-GCAACGGCAGATAACCACGACCGCAGATGCAACGGCAGATACG ACGACCGCAGATGCACTGGAAGTACAGGTTTCCA-3'

Supplementary Table S2: T1 relaxation times of ^{15}N -labelled HTD-2.

Residue	500 MHz		600 MHz		900 MHz	
	mean	s.d.	mean	s.d.	mean	s.d.
1 Gly	0.57	0.04	0.65	0.02	0.71	0.05
2 Ile	0.57	0.02	0.60	0.01	0.61	0.01
3 Cys						
4 Arg	0.59	0.02	0.58	0.02	0.57	0.01
5 Cys	0.56	0.01	0.60	0.01	0.63	0.01
6 Ile	0.54	0.01	0.56	0.03	0.56	0.00
7 Cys	0.53	0.01	0.57	0.01	0.57	0.00
8 Gly	0.55	0.01	0.57	0.01	0.63	0.00
9 Arg	0.61	0.01	0.65	0.01	0.64	0.01
10 Arg	0.67	0.04	0.69	0.02		
11 Ile	0.62	0.02	0.64	0.01	0.64	0.02
12 Cys						
13 Arg	0.58	0.01	0.57	0.01	0.60	0.00
14 Cys	0.56	0.03	0.59	0.01	0.59	0.01
15 Ile	0.54	0.01	0.55	0.01	0.58	0.01
16 Cys	0.52	0.01	0.58	0.02	0.59	0.02
17 Gly	0.61	0.03	0.58	0.01	0.61	0.01
18 Arg	0.60	0.02	0.68	0.04	0.71	0.06

Supplementary Table S3: NOE values of ^{15}N -labelled HTD-2.

Residue	500 MHz		600 MHz		900 MHz	
	mean	s.d.	mean	s.d.	mean	s.d.
1 Gly	-0.26	0.09	0.03	0.00	0.57	0.03
2 Ile	-0.36	0.03	0.05	0.00	0.60	0.01
3 Cys	-0.23	0.10	0.07	0.01	0.58	0.08
4 Arg	-0.47	0.08	0.04	0.00	0.63	0.02
5 Cys	-0.39	0.06	0.06	0.00	0.67	0.03
6 Ile	-0.38	0.05	0.13	0.01	0.65	0.02
7 Cys	-0.35	0.03	0.12	0.00	0.61	0.01
8 Gly	-0.30	0.07	0.09	0.01	0.62	0.01
9 Arg	-0.31	0.03	0.08	0.00	0.56	0.00
10 Arg	-0.29	0.12	-0.04	0.01	0.47	0.01
11 Ile	-0.28	0.10	-0.07	0.01	0.56	0.01
12 Cys	-0.21	0.05	0.09	0.03	0.51	0.04
13 Arg	-0.38	0.05	0.04	0.01	0.59	0.03
14 Cys	-0.39	0.06	0.06	0.01	0.65	0.03
15 Ile	-0.38	0.05	0.13	0.01	0.64	0.02
16 Cys	-0.25	0.06	0.08	0.00	0.62	0.02
17 Gly	-0.32	0.00	0.08	0.00	0.65	0.01
18 Arg	-0.25	0.04	0.04	0.00	0.44	0.02

Supplementary Table S4. Experimental NMR relaxation data for HTD-2 and fitted data using ModelFree.

Residue	Data ^[a]	T ₁ (s)			NOE			θ (°)	τ _i (ps)	S ²	RMSD ^[b]
		500	600	900	500	600	900				
1 Gly	Exp	0.57	0.65	0.71	-0.26	0.03	0.57	111			
	IsoRigid ^[c]	0.46	0.47	0.46	-0.31	0.04	0.57			0.33	
	Ax Rigid ^[d]	0.46	0.47	0.45	-0.32	0.04	0.56	111		0.33	
	Iso + Int ^[e]	0.64	0.65	0.64	-0.30	0.03	0.51		19	0.68	0.11
	Ax + Int ^[f]	0.64	0.65	0.64	-0.29	0.03	0.50	111	21	0.68	0.11
2 Ile	Exp	0.57	0.60	0.61	-0.36	0.05	0.60	80			
	IsoRigid	0.46	0.47	0.46	-0.31	0.04	0.57			0.24	
	Ax Rigid	0.46	0.47	0.45	-0.31	0.04	0.57	80		0.24	
	Iso + Int	0.60	0.61	0.60	-0.28	0.05	0.53		19	0.73	0.08
	Ax + Int	0.60	0.61	0.60	-0.28	0.05	0.53	80	18	0.73	0.08
3 Cys ^[g]	Exp				-0.23	0.07	0.58	91			
	IsoRigid				-0.31	0.04	0.57			0.06	
	Ax Rigid				-0.31	0.04	0.57	91		0.06	
	Iso + Int				-0.26	0.07	0.57		167	0.73	0.03
	Ax + Int				-0.26	0.07	0.56	91	125	0.66	0.03
4 Arg	Exp	0.59	0.58	0.57	-0.47	0.04	0.63	82			
	IsoRigid	0.46	0.47	0.46	-0.31	0.04	0.57			0.24	
	Ax Rigid	0.46	0.47	0.45	-0.31	0.04	0.57	82		0.24	
	Iso + Int	0.57	0.58	0.57	-0.29	0.04	0.52		26	0.77	0.15
	Ax + Int	0.57	0.58	0.57	-0.29	0.04	0.52	82	25	0.77	0.15
5 Cys	Exp	0.56	0.60	0.63	0.39	0.06	0.67	79			
	IsoRigid	0.46	0.47	0.46	-0.31	0.04	0.57			0.26	

	Ax Rigid	0.46	0.47	0.45	-0.31	0.04	0.57	79		0.26
	Iso + Int	0.61	0.62	0.61	-0.27	0.06	0.55		13	0.72 0.14
	Ax + Int	0.61	0.62	0.61	-0.27	0.06	0.55	79	13	0.73 0.13
6 Ile	<i>Exp</i>	<i>0.54</i>	<i>0.56</i>	<i>0.56</i>	<i>-0.38</i>	<i>0.13</i>	<i>0.65</i>	<i>67</i>		
	IsoRigid	0.46	0.47	0.46	-0.31	0.04	0.57			0.20
	Ax Rigid	0.46	0.47	0.45	-0.32	0.03	0.56	67		0.20
	Iso + Int	0.56	0.57	0.56	-0.23	0.10	0.59		2	0.80 0.12
	Ax + Int	0.55	0.57	0.56	-0.21	0.12	0.60	67	2	0.80 0.13
7 Cys	<i>Exp</i>	<i>0.53</i>	<i>0.57</i>	<i>0.57</i>	<i>-0.35</i>	<i>0.12</i>	<i>0.61</i>	<i>86</i>		
	IsoRigid	0.46	0.47	0.46	-0.31	0.04	0.57			0.19
	Ax Rigid	0.46	0.47	0.45	-0.31	0.04	0.57	86		0.19
	Iso + Int	0.56	0.57	0.56	-0.23	0.10	0.59		1	0.79 0.10
	Ax + Int	0.56	0.57	0.56	-0.23	0.10	0.59	86	1	0.79 0.09
8 Gly	<i>Exp</i>	<i>0.55</i>	<i>0.57</i>	<i>0.63</i>	<i>-0.30</i>	<i>0.09</i>	<i>0.62</i>	<i>87</i>		
	IsoRigid	0.46	0.47	0.46	-0.31	0.04	0.57			0.23
	Ax Rigid	0.46	0.47	0.45	-0.31	0.04	0.57	87		0.23
	Iso + Int	0.60	0.62	0.60	-0.23	0.10	0.59		1	0.74 0.09
	Ax + Int	0.60	0.62	0.60	-0.24	0.10	0.59	87	1	0.74 0.09
9 Arg	<i>Exp</i>	<i>0.61</i>	<i>0.65</i>	<i>0.64</i>	<i>-0.31</i>	<i>0.08</i>	<i>0.56</i>	<i>85</i>		
	IsoRigid	0.46	0.47	0.46	-0.31	0.04	0.57			0.31
	Ax Rigid	0.46	0.47	0.45	-0.31	0.04	0.57	85		0.31
	Iso + Int	0.64	0.65	0.64	-0.25	0.08	0.56		7	0.69 0.05
	Ax + Int	0.64	0.65	0.64	-0.26	0.07	0.56	85	6	0.69 0.05
10 Arg ^[h]	<i>Exp</i>	<i>0.67</i>	<i>0.69</i>		<i>-0.29</i>	<i>-0.04</i>	<i>0.47</i>	<i>76</i>		
	IsoRigid	0.46	0.47	0.46	-0.31	0.04	0.57			0.32
	Ax Rigid	0.46	0.47	0.45	-0.31	0.04	0.57	76		0.32
	Iso + Int	0.67	0.68		-0.35	-0.03	0.45		28	0.64 0.04
	Ax + Int	0.67	0.68	0.67	-0.35	-0.03	0.45	76	28	0.64 0.04
11 Ile	<i>Exp</i>	<i>0.62</i>	<i>0.64</i>	<i>0.64</i>	<i>-0.28</i>	<i>-0.07</i>	<i>0.56</i>	<i>117</i>		

	IsoRigid	0.46	0.47	0.46	-0.31	0.04	0.57					0.32
	Ax Rigid	0.46	0.47	0.46	-0.32	0.03	0.56	117				0.31
	Iso + Int	0.63	0.64	0.63	-0.35	-0.03	0.45		34	0.68	0.10	
	Ax + Int	0.63	0.64	0.63	-0.34	-0.03	0.44	117	37	0.67	0.10	
12 Cys ^[g]	<i>Exp</i>				-0.21	0.09	0.51	89				
	IsoRigid				-0.31	0.04	0.57					0.09
	Ax Rigid				-0.31	0.04	0.57	89				0.09
	Iso + Int				-0.25	0.08	0.57		127	0.68	0.05	
	Ax + Int				-0.25	0.08	0.57	89	127	0.72	0.06	
13 Arg	<i>Exp</i>	0.58	0.57	0.60	-0.38	0.04	0.59	91				
	IsoRigid	0.46	0.47	0.46	-0.31	0.04	0.57					0.22
	Ax Rigid	0.46	0.46	0.45	-0.31	0.04	0.57	91				0.22
	Iso + Int	0.59	0.60	0.59	-0.29	0.04	0.52		22	0.74	0.09	
	Ax + Int	0.59	0.60	0.59	-0.29	0.04	0.52	91	20	0.74	0.09	
14 Cys	<i>Exp</i>	0.56	0.59	0.59	-0.39	0.06	0.65	101				
	IsoRigid	0.46	0.47	0.46	-0.31	0.04	0.57					0.23
	Ax Rigid	0.46	0.47	0.45	-0.31	0.04	0.57	101				0.23
	Iso + Int	0.57	0.59	0.58	-0.26	0.07	0.56		13	0.77	0.12	
	Ax + Int	0.57	0.59	0.58	-0.26	0.07	0.56	101	12	0.77	0.12	
15 Ile	<i>Exp</i>	0.54	0.55	0.58	-0.38	0.13	0.64	114				
	IsoRigid	0.46	0.47	0.46	-0.31	0.04	0.57					0.20
	Ax Rigid	0.46	0.47	0.45	-0.32	0.03	0.56	114				0.20
	Iso + Int	0.55	0.56	0.55	-0.23	0.10	0.59		1	0.81	0.12	
	Ax + Int	0.55	0.56	0.55	-0.21	0.12	0.60	114	2	0.81	0.13	
16 Cys	<i>Exp</i>	0.52	0.58	0.59	-0.25	0.07	0.62	98				
	IsoRigid	0.46	0.47	0.46	-0.31	0.04	0.57					0.20
	Ax Rigid	0.46	0.47	0.45	-0.31	0.04	0.57	98				0.20
	Iso + Int	0.55	0.56	0.55	-0.25	0.08	0.56		13	0.80	0.06	
	Ax + Int	0.55	0.56	0.55	-0.25	0.08	0.57	98	11	0.80	0.06	

17 Gly	<i>Exp</i>	0.61	0.58	0.61	-0.32	0.08	0.65	94			
	IsoRigid	0.46	0.47	0.46	-0.31	0.04	0.57				0.26
	Ax Rigid	0.46	0.47	0.45	-0.31	0.04	0.57	94			0.26
	Iso + Int	0.61	0.62	0.61	-0.27	0.06	0.55		13	0.72	0.09
	Ax + Int	0.61	0.62	0.61	-0.27	0.06	0.55	94	12	0.73	0.09
18 Arg	<i>Exp</i>	0.60	0.68	0.71	-0.25	0.04	0.44	127			
	IsoRigid	0.46	0.47	0.46	-0.31	0.04	0.57				0.38
	Ax Rigid	0.46	0.47	0.46	-0.34	0.02	0.56	127			0.38
	Iso + Int	0.62	0.63	0.62	-0.29	0.04	0.52		18	0.70	0.12
	Ax + Int	0.62	0.63	0.63	-0.27	0.04	0.50	127	26	0.68	0.11

RMSD ^[i]	IsoRigid	0.50	0.58	0.65	0.20	0.16	0.20				
	Ax Rigid	0.50	0.58	0.65	0.21	0.16	0.19				
	Iso + Int	0.12	0.09	0.13	0.29	0.05	0.21				
	Ax + Int	0.12	0.09	0.13	0.29	0.04	0.21				

^[a] Abbreviations used: Exp, experimental data; IsoRigid, isotropic rigid; Ax Rigid, axial rigid; Iso + Int, isotropic with internal motion, Ax + Int, axial with internal motion.

$$^{[b]} RMSD = \sqrt{\sum(T_1 exp - T_1 calc)^2 + \sum[0.5(NOE exp - NOE calc)^2]}$$

RMSD is the root mean square deviation between the experimental and fitted (calc) T₁ and NOE values. The NOE deviations were weighted by 50% because of their intrinsically higher errors compared to the T₁ values.

^[c]For isotropic rigid motion, the overall correlation time, $\tau_0 = 1.07$ ns.

^[d]For axial rigid motion, the overall correlation time, $\tau_0 = 1.06$ ns and the D_{ratio} = 0.90.

^[e]For isotropic with internal motion, the overall correlation time, $\tau_0 = 1.13$ ns.

^[f]For axial with internal motion, the overall correlation time, $\tau_0 = 1.16$ ns and the D_{ratio} = 1.27.

^[g] The ¹⁵N HSQC peaks for Cys3 and Cys12 were overlapped with those of Ile6 and Ile15 and a satisfactory exponential fit could not be obtained to determine T₁.

^[h] Broadening of the Arg10 NH peak resulted in a poor exponential fit so Arg10 T₁ was excluded from the analysis.

$$^{[i]} RMSD = \sqrt{\sum(exp - calc)^2}$$

RMSD is the root mean square deviation between the experimental and fitted (calc) T_1 and NOE values calculated over all the residues at each field strength.

Supplementary Table S5: Fitted parameters and errors using an in-house spreadsheet.

Model	Residue(s)	τ_0 (ps)	s.d.	D_{ratio}	s.d.	τ_i (ps)	s.d.	S^2	s.d.
<i>IsoRigid</i>									
All		1.05	0.004						
<i>Ax Rigid</i>									
All		1.08	NA	2.44	NA				
All		1.02	0.03	0.87	0.07				
<i>Iso + Int</i>									
All		1.09	0.007						
1 Gly				2.45	3.82			0.71	0.03
2 Ile				6.15	3.37			0.76	0.01
3 Cys				24.92	16.94			0.93	0.15
4 Arg				17.56	8.43			0.77	0.02
5 Cys				1.34	3.10			0.77	0.01
6 Ile				0.00	0.00			0.82	0.02
7 Cys				0.00	0.00			0.82	0.01
8 Gly				1.00	2.94			0.78	0.01
9 Arg				2.44	1.72			0.72	0.01
10 Arg				18.10	10.10			0.66	0.02
11 Ile				11.72	7.01			0.71	0.01
12 Cys				24.41	13.39			0.96	0.15
13 Arg				11.55	7.45			0.8	0.01
14 Cys				3.62	4.01			0.77	0.02
15 Ile				0.26	0.76			0.82	0.01
16 Cys				0.00	0.00			0.81	0.01
17 Gly				0.00	0.00			0.76	0.01
18 Arg				10.11	3.46			0.67	0.02

Supplementary Table S6: Fitted parameters and errors using MODELFREE.

Model	Residue(s)	τ_0 (ps)	s.d.	D_{ratio}	s.d.	τ_i (ps)	s.d.	S^2	s.d.
<i>IsoRigid</i>									
All		1.07	0.000						
<i>Ax Rigid</i>									
All		1.06	0.002	0.90	0.01				
<i>Iso + Int</i>									
All		1.13	0.002						
1 Gly				18.91	1.65	0.68	0.02		
2 Ile				18.52	1.01	0.73	0.01		
3 Cys				166.56	185.00	0.73	0.29		
4 Arg				26.17	1.84	0.77	0.01		
5 Cys				13.09	1.16	0.72	0.01		
6 Ile				1.70	2.01	0.80	0.01		
7 Cys				1.04	1.19	0.79	0.00		
8 Gly				1.11	1.25	0.74	0.01		
9 Arg				7.21	0.52	0.69	0.01		
10 Arg				27.95	2.18	0.64	0.01		
11 Ile				34.06	1.63	0.68	0.01		
12 Cys				127.38	178.02	0.68	0.32		
13 Arg				21.97	1.78	0.74	0.01		
14 Cys				12.68	2.24	0.77	0.01		
15 Ile				1.44	1.89	0.81	0.01		
16 Cys				12.99	2.03	0.80	0.01		
17 Gly				13.35	1.00	0.72	0.01		
18 Arg				17.71	1.78	0.70	0.02		
<i>Ax + Int</i>									

All	1.16	0.009				
All	1.24	0.07				
1 Gly			21.10	1.91	0.68	0.02
2 Ile			17.91	0.94	0.73	0.01
3 Cys			124.61	176.66	0.66	0.31
4 Arg			24.88	1.82	0.77	0.01
5 Cys			12.60	1.09	0.73	0.01
6 Ile			1.61	2.09	0.80	0.01
7 Cys			1.00	1.20	0.79	0.00
8 Gly			0.83	1.10	0.74	0.01
9 Arg			6.39	0.57	0.69	0.01
10 Arg			27.92	2.30	0.64	0.01
11 Ile			36.97	1.95	0.67	0.01
12 Cys			127.37	177.14	0.72	0.30
13 Arg			19.99	1.78	0.74	0.30
14 Cys			11.79	2.33	0.77	0.01
15 Ile			1.60	2.05	0.81	0.01
16 Cys			11.22	1.96	0.80	0.01
17 Gly			11.66	1.04	0.73	0.01
18 Arg			26.09	3.00	0.68	0.02
