Supplementary Information

Positional effects of click cyclization on β-hairpin structure, stability, and function

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Proton Assignments (ppm) for Term-U						
	α	β	γ	δ	3	
AzK	4.32	1.78	1.43	1.60	3.30	
R	4.64	1.79	1.58	3.19		
W	5.35	3.28, 3.00				
V	4.73	2.00	0.91			
Κ	4.36	1.06, 1.16	-0.11	0.46	2.31	
V	4.14	1.85	0.87			
Ν	4.37	3.08, 2.75				
G*	0.74					
0	4.70	1.93	1.76	3.07		
W	5.05	3.20, 3.08				
Ι	4.79	1.98	1.25,1.46	0.96 (y,CH ₃)	0.83 (δ,CH ₃)	
Κ	4.03	1.26	0.39, 0.47	0.67	2.65	
Q	4.49	2.06, 1.87	2.25			
Pra	4.70	2.51				

*Glycine splitting value

Table S2

Proton Assignments (ppm) for Term-C							
	α	β	γ	δ	3		
K- N3	4.07	1.87	1.11	1.47, 1.57	4.36		
R	4.67	1.79	1.60	3.19			
W	5.30	3.29, 3.04					
V	4.69	2.00	0.90				
K	4.35	1.13, 1.06	-0.10	0.43	2.29		
\mathbf{V}	4.13	1.84	0.86				
Ν	4.36	3.08, 2.75					
G*	0.74						
0	4.70	1.92	1.75	3.06			
W	5.04	3.18, 3.07					
Ι	4.76	1.96	1.44, 1.24	0.93 (y,CH ₃)	0.84 (δ,CH ₃)		
K	4.18	1.18	0.41	0.70	2.47		
Q	4.46	2.00, 1.84	2.23				
Pra	4.80	2.92, 2.76					

	Proton Assignments (ppm) for Term-rev-U					
	α	β	γ	δ	3	
K- N3	4.38	1.62	1.34	1.47	3.24	
R	4.58	1.85, 1.77	1.58	3.19		
W	5.28	3.29, 3.00				
V	4.67	2.00	0.89			
K	4.32	1.06, 1.16	-0.10	0.46	2.32	
V	4.12	1.84	0.86			
Ν	4.36	3.06, 2.73				
G*	0.72					
0	4.67	1.91	1.74	3.06		
W	5.01	3.18, 3.05				
Ι	4.78	1.97	1.44, 1.23	0.95 (y,CH ₃)	0.82 (δ,CH ₃)	
K	4.21	1.22	0.46	0.86	2.56	
Q	4.50	2.05, 1.87	2.26			
Pra	4.46	2.71, 2.45				

*Glycine splitting value

Table S4

Proton Assignments (ppm) for Term-rev-C							
	α	β	γ	δ	3		
K- N3	4.09	1.65	0.43	1.42	4.35		
R	4.64	1.72	1.56	3.16			
W	5.27	3.19,2.95					
V	4.67	1.94	0.83				
K	4.31	1.00, 1.10	-0.19	0.41	2.24		
V	4.09	1.79	0.83				
Ν	4.32	3.04, 2.71					
G*	0.74						
0	4.66	1.94	1.74	3.03			
W	4.98	3.14, 3.03					
Ι	4.78	1.97	1.41, 1.24	0.91 (y,CH ₃)	0.82 (δ,CH ₃)		
K	4.07	1.39	0.60	0.99	2.71		
Q	4.46	1.89, 1.72	2.14				
Pra	4.70	3.28, 2.93					

	Proton Assignments (ppm) for NHB-U						
	α	β	γ	δ	3		
R	4.61	1.85	1.69	3.01			
K- N3	4.37	1.78	1.63	1.94			
V	4.57	1.99	0.88				
K	4.30	1.08, 1.18	0.09, 0.14	0.52	2.32		
V	4.10	1.83	0.85				
Ν	4.38	3.00, 2.72					
G*	0.63						
0	4.99	1.70, 1.81	1.39	3.05			
W	4.87	3.10					
Ι	4.58	1.94	1.39, 1.18	0.89 (y,CH ₃)	0.82 (d,CH3)		
Pra	4.76	2.71					
Q	4.38	2.11, 1.94	2.31				

*Glycine splitting value

Table S6

	Proton Assignments (ppm) for NHB-C						
	α	β	γ	δ	3		
R	4.30	1.78	1.51	3.21			
K- N3	4.29	1.64	0.71	0.71 1.06			
V	4.65	1.97	0.85				
K	4.37	1.12	-0.11, 0.16	0.40, 0.91	2.04		
V	4.09	1.79	0.84				
Ν	4.33	3.04, 2.71					
G*	0.72						
0	4.62	1.86	1.69	3.00			
W	4.84	3.12					
Ι	4.59	1.98	1.35, 1.28	0.89 (y,CH ₃)	0.78 (δ,CH ₃)		
Pra	4.09	2.42, 2.31					
Q	4.41	2.11, 2.01	2.33				

	Proton Assignments (ppm) for NHB-rev-U						
	α	β	γ	δ	3		
R	4.40	1.84, 1.79	1.65	3.20			
K- N3	4.54	1.77	1.43	1.60			
\mathbf{V}	4.49	2	0.87				
K	4.31		0.63	1.27, 1.17	2.42		
\mathbf{V}	4.10	1.87	0.85				
Ν	4.42	3.00, 2.73					
G*	0.55						
0	4.57	1.83	1.69	3.01			
W	5.08	2.73, 2.67					
Ι	4.45	1.91	1.40, 1.19	0.88 (y,CH ₃)	0.77 (δ,CH ₃)		
Pra	4.80	3.11					
Q	4.37	2.10, 1.97	2.33				

*Glycine splitting value

Table S8

Proton Assignments (ppm) for NHB-rev-C						
	α	β	γ	δ	3	
R	4.32	1.80	1.63	3.21		
K- N3	4.21	1.78, 1.98	0.94, 1.13	1.56	4.43, 4.37	
\mathbf{V}	4.65	1.98	0.86			
K	4.29	1.07, 1.16	0.00	0.44	2.29	
\mathbf{V}	4.10	1.81	0.84			
Ν	4.36	3.04, 2.71				
G*	0.71					
0	4.65	1.87	1.71	3.03		
W	5.34	3.17, 3.06				
Ι	4.57	1.92	1.36, 1.21	0.88 (y,CH ₃)	0.81 (δ,CH ₃)	
Pra	4.87	3.14				
Q	4.31	2.05, 1.96	2.29			

Proton Assignments (ppm) for HB-U							
	α	β	γ	δ	3		
R	4.32	1.69	1.54	3.14			
W	4.99	3.27, 3.10					
K-N3	4.63	1.74, 1.67	1.26	1.52	3.25		
K	4.00	1.39	0.68	0.86	2.73		
V	4.11	2.01	0.89				
Ν	4.45	2.98, 2.73					
G*	0.48						
0	4.58	1.88	1.70	3.03			
W	4.90	3.27, 3.12					
Pra	4.78	2.67, 2.41					
K	4.31	1.35, 1.19	0.48	0.79, 0.64	2.43		
Q	4.27	2.06, 1.92	2.30				

*Glycine splitting value

Table S10

	Proton Assignments (ppm) for HB-C						
	α	β	γ	δ	3		
R	4.49	1.75	1.63	3.22			
W	5.30	3.32, 3.13					
K-N3	4.82	1.93, 1.67	0.86	1.16	4.30, 4.26		
K	3.90	1.28	0.33, 0.57	1.17	2.68		
V	4.21	1.97	0.92				
Ν	4.41	3.05, 2.80					
G*	0.66						
0	4.80	1.75	1.63	3.12			
W	5.07	3.31, 3.02					
Pra	5.13	3.35					
K	4.54		0.44	1.06, 0.93			
Q	4.32	2.08, 1.94	2.33				

	Proton Assignments (ppm) for HB-rev-U						
	α	β	γ	δ	3		
R	4.41	1.68	1.58	3.17			
W	5.17	3.29, 3.06					
K-N3	4.71	1.82, 1.75	1.40	1.59	3.32		
K	4.34	1.28	0.53	1.15	2.39		
V	4.11	1.90	0.90				
Ν	4.41	3.02, 2.76					
G*	0.59						
0	4.64	1.91	1.75	3.06			
W	4.92	3.25, 3.12					
Pra	4.87	2.67					
K	3.92	1.37	0.63	0.82	2.72		
Q	4.28	2.04, 1.91	2.29				

*Glycine splitting value

Table S12

	α	β	γ	δ	3
R	4.48	1.77	1.61	3.18	
W	5.32	3.35, 3.07			
K-N3	4.79	1.96	1.26	1.75	4.32
K	4.34	1.00	0.03, 0.18	0.27	2.30
\mathbf{V}	4.15	1.88	0.82		
Ν	4.39	3.03, 2.77			
G*	0.69				
0	4.71	1.93	1.75	3.08	
W	4.92	3.26, 3.07			
Pra	5.20	3.35			
K	4.05	1.32, 1.20	0.43	0.7	2.71
Q	4.27	2.01, 1.86	2.20		

	Proton Assignments (ppm) for Term-pG-U						
	α	β	γ	δ	3		
Pra	4.68	2.48					
R	4.62	1.78	1.57	3.18			
W	5.31	3.24, 3.01					
\mathbf{V}	4.68	1.99	0.89				
K	4.30	1.19, 0.97	-0.07	0.5			
\mathbf{V}	4.50	1.90	0.88				
dP	4.33	2.32	2.08, 1.93	3.80, 3.73			
G*							
0	4.72	1.90	1.75	3.04			
W	5.09	3.17, 3.04					
Ι	4.76	1.95	1.42, 1.22	0.92 (y,CH ₃)	0.80 (δ,CH ₃)		
K	3.99	1.24	0.46. 0.31	0.60	2.61		
Q	4.46	2.02, 1.85	2.23				
K- N3	4.30	2.75	1.41	1.58	3.29		

	Proton Assignments (ppm) for Term-pG-C					
	α	β	γ	δ	3	
Pra	4.82	2.94, 2.77				
R	4.69	1.82	1.62	3.20		
W	5.27	3.29, 3.07				
\mathbf{V}	4.67	2.02	0.92			
K	4.31	1.20, 0.99	0.04	0.51	2.38	
\mathbf{V}	4.50	1.92	0.88			
dP	4.34	2.33	1.95, 2.11	3.81, 3.74		
G*						
0	4.74	1.93	1.77	3.07		
W	5.10	3.18, 3.03				
Ι	4.76	1.97	1.45, 1.24	0.95 (y,CH ₃)	0.84 (δ,CH ₃)	
K	4.15	1.17	0.45, 0.23	0.65	2.47	
Q	4.47	2.02, 1.84	2.24			
K- N3	4.06	1.89	1.12	1.58, 1.47	4.38, 4.36	

	Proton Assignments (ppm) for NHB-pG-U					
	α	β	γ	δ	3	
R	4.36	1.75	1.61	3.19		
K- N3	5.03		1.36	1.80, 1.71	3.03	
V	4.58	1.97	0.85			
K	4.22	1.16, 0.95	0.01	0.49	2.30	
V	4.46	1.94	0.87			
dP	4.31	2.29	1.90, 2.04	3.76, 3.69		
G*						
0	4.66	1.85	1.70	2.99		
W	4.94	3.07, 2.96				
Ι	4.62	1.94	1.38, 1.17	0.89 (y,CH ₃)	0.78 (δ,CH ₃)	
Pra	4.80	2.70				
Q	4.37	2.09, 1.94	2.30			

	α	β	γ	δ	3
R	4.30	1.80	1.59	3.22	
K- N3	4.30	1.66, 1.54	0.70	1.05	4.55, 4.54
\mathbf{V}	4.58	1.96	0.88		
K	4.30	1.21, 0.88	-0.06	0.28, 0.46	
V	4.46	1.86	0.86		
dP	4.30	2.3	2.07, 1.90	3.80, 3.69	
G*					
0	4.63	1.86	1.71	3.01	
W	4.91	3.06, 2.85			
Ι	4.58	1.96	1.27	0.88 (y,CH ₃)	0.80 (d,CH3)
Pra	4.92	3.23			
Q	4.41	2.14,2.01	2.34		

	Р	roton Assignments	(ppm) for HB-pG-	U	
	α	β	γ	δ	3
R	4.43	1.71	1.62	3.19	
W	5.12	3.28, 3.08			
K-N ₃	4.83	1.55		1.33	1.77
K	4.27	1.32	0.30, 0.57	1.04	2.40
V	4.48	2.04	0.90		
dP	4.36	2.31	2.07, 1.94	3.72	
G*					
0	4.71	1.91	1.77	3.07	
W	5.09	3.27, 3.07			
Pra	4.99	2.73, 2.66			
K	3.96	1.30	0.46, 0.58	0.76	2.67
Q	4.29	2.06, 1.91	2.31		

Proton Assignments (ppm) for NHB-pG-C						
	α	β	γ	δ	3	
R	4.47	1.77	1.59, 1.69	3.19		
W	5.35	3.28, 3.06				
K-N ₃	4.29	1.70	0.92	1.15		
K	3.88	1.18	0.32	0.53	2.64	
\mathbf{V}	4.53	2.01	0.92			
dP	4.37	2.32	2.07, 1.94	3.75, 3.67		
G*						
0	4.80	1.93, 1.84	1.67	3.07		
W	5.06	3.27, 3.01				
Pra	5.14	3.32				
K	4.53		0.05, 0.42	12	2.50	
Q	4.30	2.05, 1.89	2.31			

I	Proton Assignments for Random Coil strands				
	Ηα		Ηα		
R	4.25	G	3.98		
W	4.74	0	4.16		
Azk/V	4.36/4.00	W	4.70		
Κ	4.13	Pra/I	4.30/4.06		
\mathbf{V}	4.08	K	4.29		
Ν	4.67	Q	4.29		



Figure S1. Thermal stability assessed by circular dichroism of (A) HB at 213 nm, (B) NHB at 215 nm, and (C) Term at 212 nm from 5 °C to 95 °C in 10 mM phosphate buffer, pH 7.5.



Figure S2. Thermal stability assessed by circular dichroism of (A) **HB-pG**, (B) **NHB-pG**, and (C) **Term-pG** peptides from 20 °C to 90 °C in 10 mM phosphate buffer, pH 7.5.



Figure S3. (A) Circular dichrosim spectra of **Term-pG-U** (dark green) compared to **Term -pG-**C (light green) at 298 K in 10mM sodium phosphate pH 7.0 buffer. (B) H α chemical shift differences for **Term -pG-U** (dark green), **Term -pG-C** (light green), **WKWK-pG** (orange), and **WKWK-pG-Cys** (grey) from random coil peptides. Values calculated from data obtained at 293 K in 50 mM potassium phosphate-*d2*, pD 7.0 (uncorrected). *The Gly bars reflect the glycine splitting.



Figure S4. NOEs observed between cross-strand H α protons (blue), side-chain side-chain (red), and the triazole proton (green) in (A) **Term-pG-U** and (B) **Term-pG-C**.

Peptide	Sequence
Term-pG-AzK	Ac-(AzK)RWVK <u>VNGO</u> WIKQ(Pra)-NH ₂
Term-pG-AzO	Ac-(AzO)RWVK <u>VNGO</u> WIKQ(Pra)-NH ₂
Term-pG-AzB	Ac-(AzB)RWVK <u>VNGO</u> WIKQ(Pra)-NH ₂
Term-pG-AzP	Ac-(AzP)RWVK <u>VNGO</u> WIKQ(Pra)-NH ₂
Prefixes of "II" and	1 "C" denote linear unclicked and cyclic clicked pentides $\mathbf{n} = d\mathbf{P}\mathbf{r}_0 \mathbf{A}\mathbf{z}\mathbf{O} =$

Table S20	Sequences	of Term-pG-AzX	X peptides
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Prefixes of "U" and "C" denote linear unclicked and cyclic clicked peptides. p = dPro, $AzO = Orn(N_3)$ (azido-ornithine), $AzB = Dab(N_3)$ (azido-homoalanine), $AzP = Dap(N_3)$ (azido-alanine)



Figure S5. (A) Circular dichrosim spectra of **Term-pG-AzK-U** (red), **Term-pG-AzK-C** (blue), **Term-pG-AzO-C** (green), **Term-pG-AzB-C** (orange), and **Term-pG-AzP-C** (purple) at 298 K in 10mM sodium phosphate pH 7.0 buffer. (B) H α chemical shift differences for **Term-pG-AzK-U** (red), **Term-pG-AzK-C** (blue), **Term-pG-AzO-C** (green), **Term-pG-AzB-C** (orange), and **Term-pG-AzP-C** (purple) from random coil peptides. Values calculated from data obtained at 293 K in 50 mM potassium phosphate-*d2*, pD 7.0 (uncorrected). Glycine shifts were unclear due to overlap with the H δ protons of dPro. The residue N3 represents the azide that was incorporated.



Figure S6. (A) Circular dichrosim spectra of **NHB-pG-U** (dark red) compared to **NHB-pG-C** (light red) at 298 K in 10mM sodium phosphate pH 7.0 buffer. (B) Hα chemical shift differences for **NHB-pG-U** (dark red), **NHB-pG-C** (light red), **WKWK-pG** (orange), and **WKWK-pG-Cys** (grey) from random coil peptides. Values calculated from data obtained at 293 K in 50 mM potassium phosphate-*d2*, pD 7.0 (uncorrected). *The Gly bars reflect the glycine splitting.



Figure S7. NOEs observed between cross-strand H α protons (blue), side-chain side-chain (red), and the triazole proton (green) in (A) **NHB-pG-U** and (B) **NHB-pG-C**.



Figure S8. (A) Circular dichrosim spectra of **HB-pG-U** (dark blue) compared to **HB-pG-C** (light blue) at 298 K in 10mM sodium phosphate pH 7.0 buffer. (B) Hα chemical shift differences for **HB-pG-U** (dark blue), **HB-pG-C** (light blue), **WKWK-pG** (orange), and **WKWK-pG-Cys** (grey) from random coil peptides. Values calculated from data obtained at 293 K in 50 mM potassium phosphate-*d2*, pD 7.0 (uncorrected)



Figure S9. NOEs observed between cross-strand H α protons (blue), side-chain side-chain (red), and the triazole proton (green) in (A) **HB-pG-U** and (B) **HB-pG-C**.



Figure S10. Overlay of HPLC traces monitoring Pronase E degradation of **Scramble** at 280 nm over 24 hours.



Figure S11. Overlay of HPLC traces monitoring Pronase E degradation of **HB-pG-U** at 280 nm over 24 hours.



Figure S12. Overlay of HPLC traces monitoring Pronase E degradation of **HB-pG-C** at 280 nm over 24 hours.



Figure S13. Overlay of HPLC traces monitoring Pronase E degradation of **NHB-pG-U** at 280 nm over 24 hours.



Figure S14. Overlay of HPLC traces monitoring Pronase E degradation of **NHB-pG-C** at 280 nm over 24 hours.



Figure S15. Overlay of HPLC traces monitoring Pronase E degradation of **Term-pG-U** at 280 nm over 24 hours.



Figure S16. Overlay of HPLC traces monitoring Pronase E degradation of **Term-pG-C** at 280 nm over 24 hours.



Figure S17. Overlay of HPLC traces monitoring Trypsin degradation of **NHB-pG-U** at 280 nm over 24 hours.



Figure S18. Overlay of HPLC traces monitoring Trypsin degradation of **NHB-pG-C** at 280 nm over 24 hours.



Figure S19. Fluorescence quenching titrations of (A) **Term-C** (red) and **Term-rev-C** (blue) and (B) **HB-C** (blue) and **HB-rev-C** (red) with ATP at 5 μ M peptide in 10 mM acetate buffer, pH 5.0 at 25 °C.

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Figure S1 ¹H NMR spectrum for **WKWK-pG-Cys**: Ac-Cys-Arg-Trp-Val-Lys-Val-dPro-Gly-Orn-Trp-Ile-Lys-Gln-Cys-NH₂

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Figure S2 ¹H NMR spectrum for **HB-pG-U**: Ac-Arg-Trp-AzK-Lys-Val-dPro-Gly-Orn-Trp-Pra-Lys-Gln-NH₂

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Figure S3 ¹H NMR spectrum for **HB-pG-C**: Ac-Arg-Trp-AzK-Lys-Val-dPro-Gly-Orn-Trp-Pra-Lys-Gln-NH₂

Figure S4 ¹H NMR spectrum for **NHB-pG-U**: Ac-Arg-AzK-Val-Lys-Val-dPro-Gly-Orn-Trp-Ile-Pra-Gln-NH₂

\$10.6.8.8.6.6.5.4.8 Expresses" STATISTICS FOR

Figure S5 1 H NMR spectrum for NHB-pG-C: Ac-Arg-AzK-Val-Lys-Val-dPro-Gly-Orn-Trp-Ile-Pra-Gln-NH₂

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Figure S6 ¹H NMR spectrum for **Term-pG-U**: Ac-Pra-Arg-Trp-Val-Lys-Val-dPro-Gly-Orn-Trp-Ile-Lys-Gln-AzK-NH₂

\$10.6.6.8.6.6.6.2.8.4.6. U.4 Hores points " 0.38001 shtnp://///R/pask.ph7-3/ft.og.2010/00/06/99/2 [[]_8090.00 [12] 73/3880 66//6402271921 Hate

Figure S7 ¹H NMR spectrum for **Term-pG-C**: Ac-Pra-Arg-Trp-Val-Lys-Val-dPro-Gly-Orn-Trp-Ile-Lys-Gln-AzK-NH₂

Figure S8 1 H NMR spectrum for HB-U: Ac-Arg-Trp-AzK-Lys-Val-Asn-Gly-Orn-Trp-Pra-Lys-Gln-NH₂

Figure S9 1 H NMR spectrum for HB-C: Ac-Arg-Trp-AzK-Lys-Val-Asn-Gly-Orn-Trp-Pra-Lys-Gln-NH₂

Figure S10 ¹H NMR spectrum for **HB-rev-U**: Ac-Arg-Trp-Pra-Lys-Val-Asn-Gly-Orn-Trp-AzK-Lys-Gln-NH₂

.8.0.0.4

Figure S11 ¹H NMR spectrum for **HB-rev-C**: Ac-Arg-Trp-Pra-Lys-Val-Asn-Gly-Orn-Trp-AzK-Lys-Gln-NH₂

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Figure S12 ¹H NMR spectrum for **NHB-U**: Ac-Arg-AzK-Val-Lys-Val-Asn-Gly-Orn-Trp-Ile-Pra-Gln-NH₂

.8.6.6.6.4.8 tot: "presat"

Figure S13 ¹H NMR spectrum for **NHB-C**: Ac-Arg-AzK-Val-Lys-Val-Asn-Gly-Orn-Trp-Ile-Pra-Gln-NH₂

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Figure S14 $\,^1\mathrm{H}$ NMR spectrum for NHB-rev-U: Ac-Arg-Pra-Val-Lys-Val-Asn-Gly-Orn-Trp-Ile-Lys-AzK-Gln-NH_2

.6.0.6.6.0.5.4.8.4.0.6. ALL BORNES

Figure S15 1 H NMR spectrum for NHB-rev-C: Ac-Arg-Pra-Val-Lys-Val-Asn-Gly-Orn-Trp-Ile-Lys-AzK-Gln-NH₂

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Figure S16 ¹H NMR spectrum for **Term-U**: Ac-Pra-Arg-Trp-Val-Lys-Val-Asn-Gly-Orn-Trp-Ile-Lys-Gln-AzK-NH₂

0.8.8.4.0.6.4.8 Malex preset 1) Friterlick - 143, ph/2-5 8000 000 fiz - 213 9 880

Figure S17 1 H NMR spectrum for Term-C: Ac-Pra-Arg-Trp-Val-Lys-Val-Asn-Gly-Orn-Trp-Ile-Lys-Gln-AzK-NH₂

P.810 G. B. B. G. D. G. Z. B. 4. G. B. 2. B. 4. O. G. O. B. 0. O. 4. Shamiling revelue app. ant 5. **(BACATORICATIONAL STREET**) BIRGERT Will Volume 1. J. 1933 Solv **Barken US27192** J. **Home**, Const. **10**, 38001

Figure S18 ¹H NMR spectrum for **Term-rev-U**: Ac-AzK-Arg-Trp-Val-Lys-Val-Asn-Gly-Orn-Trp-Ile-Lys-Gln-Pra-NH₂

Figure S19 ¹H NMR spectrum for **Term-rev-C**: Ac-AzK-Arg-Trp-Val-Lys-Val-Asn-Gly-Orn-Trp-Ile-Lys-Gln-Pra-NH₂

Analytical HPLC trace for peptides: Lyophilized peptides were reconstituted in water for HPLC analysis. 5 μ L of sample was injected on a Waters 2695 HPLC using an Atlantis C-18 column with a gradient of 0 to 50% solvent B in 25 minutes, where solvent B is 95% acetonitrile, 5% water, and 0.1% trifluoroacetic acid and solvent A is 95:5:0.1 of H₂O:ACN:TFA.



Figure S20. HPLC trace of Term-rev-C



Figure S21. HPLC trace of Term-rev-U







Figure S23. HPLC trace of WKWK



Figure S24. HPLC trace of HB-rev-C



Figure S25. HPLC trace of HB-rev-U



Figure S26. HPLC trace of Term-U



Figure S27. HPLC trace of NHB-rev-U



Figure S28. HPLC trace of HB-C



Figure S30. HPLC trace of HB-U



Figure S31. HPLC trace of NHB-rev-C







Figure S33. HPLC trace of HB-pG-C



Figure S34. HPLC trace of Term-pG-U



Figure S35. HPLC trace of NHB-U



Figure S36. HPLC trace of Scramble



Figure S37. HPLC trace of NHB-C



Figure S38. ESI-MS of Term-rev-C, calculated mass: 1817.03



Figure S39. ESI-MS of Term-rev-U, calculated mass: 1817.03



Figure S40. ESI-MS of Term-C, calculated mass: 1817.03



Figure S41. ESI-MS of HB-rev-C, calculated mass: 1604.88



Figure S42. ESI-MS of WKWK, calculated mass: 1567.91



Figure S43. ESI-MS of HB-rev-U, calculated mass: 1604.88



Figure S44. ESI-MS of Term-U, calculated mass: 1817.03



Figure S45. ESI-MS of NHB-rev-U, calculated mass: 1502.86



Figure S46. ESI-MS of HB-C, calculated mass: 1604.88



Figure S47. ESI-MS of HB-U, calculated mass: 1604.88



Figure S48. ESI-MS of NHB-rev-C, calculated mass: 1502.86



Figure S49. ESI-MS of HB-pG-U, calculated mass: 1587.89



Figure S50. ESI-MS of Term-pG-C, calculated mass: 1800.04



Figure S51. ESI-MS of Term-pG-U, calculated mass: 1800.04



Figure S52. ESI-MS of NHB-pG-U, calculated mass: 1485.87



Figure S53. ESI-MS of NHB-U, calculated mass: 1502.86



Figure S54. ESI-MS of Scramble, calculated mass: 1800.04