Supporting Information

Stapling strategy enables to improve antitumor activity and proteolytic stability of host-defense peptide hymenochirin-1B

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Table S1. Electrospray MS data for peptides (positive mode)

Peptide	Molecular	Found Mass	Yield
	Weight (g/mol)		
hymenochirin-	3061.87	$[M+2H]^{2+} = 1532.4541; [M+3H]^{3+} =$	31.23%
1B		1021.9723	
H-1	3100.97	$[M+2H]^{2+} = 1551.9954; [M+3H]^{3+} = 11.43\%$	
		1034.6734	
H-2	3069.96	$[M+2H]^{2+} = 1535.9828; [M+3H]^{3+} =$	6.43%
		1024.3264	
H-3	3183.03	$[M+2H]^{2+} = 1593.0213; [M+3H]^{3+} =$	6.51%
		1062.3504	
H-4	3227.04	$[M+3H]^{3+} = 1076.6888; [M+4H]^{4+} =$	6.72%
		807.7680;	
H-5	3170.97	$[M+3H]^{3+}= 1057.9979; [M+4H]^{4+}=$	19.94%
		793.7491	
H-6	3098.91	$[M+2H]^{2+}= 1550.9557; [M+3H]^{3+}=$	7.25%
		1033.9745	
H-7	3127.93	$[M+2H]^{2+}=1586.9935; [M+3H]^{3+}=$	13.07%
		1057.9994	
H-8	3113.92	$[M+3H]^{3+}= 1038.9788; [M+4H]^{4+}=$	14.06%
		779.4843	
H-9	3199.99	$[M+3H]^{3+}= 1067.3436; [M+4H]^{4+}=$	18.47%
		800.7592	
H-10	3177.05	$[M+2H]^{2+} = 1590.5354; [M+3H]^{3+} =$	5.43%
		1060.3598	



Figure S1. The dose response curves for stapled peptides against non-small cell lung adenocarcinoma A549. H-0 represented template peptide Hymenochirin-1B. All experiments were repeated three times. H-0 represents hymenochirin-1B.



Figure S2. The dose response curves for stapled peptides against hepatocarcinoma HepG2. H-0 represented template peptide Hymenochirin-1B. All experiments were repeated three times. H-0 represents hymenochirin-1B.



Figure S3. The dose response curves for stapled peptides against colon cancer cell HCT116. H-0 represented template peptide Hymenochirin-1B. All experiments were repeated three times. H-0 represents hymenochirin-1B.



Figure S4. Proteolytic stability of peptides incubated in trypsin solution (5 ng/µl in 50 mM PBS buffer, pH=7.4) at a final concentration of 0.1 mM. Date points are displayed as the mean value SEM of duplicate independent experiments. The percent of residual peptide was monitored by HPLC. All experiments were repeated three times. H-0 represents hymenochirin-1B.

peptide		IC ₅₀ (μM)	
	LO-2	BEAS-2B	293T
hymenochirin-1B	23.98±0.32	9.24±0.25	11.12±0.35
H-1	7.80±0.21	9.61±0.35	5.82±0.45
H-2	3.40±0.32	3.12±0.52	3.20±0.33
Н-3	7.61±0.43	7.95±0.36	4.58±0.34
H-4	8.36±0.42	25.01±0.23	5.41±0.32
Н-5	3.57±0.52	5.76±0.32	4.70±0.32
Н-6	49.77±0.22	46.58±0.42	60.11±0.35
H-7	6.52±0.32	7.50±0.32	7.25±0.26
H-8	24.82±0.44	12.82±0.25	15.58±0.36
H-9	8.56±0.32	7.72±0.36	8.66±0.23
H-10	8.78±0.32	5.52±0.42	10.42±0.25

Table S2. Toxicity of the peptides against normal cells

























