Supplementary information

An Unprecedented alteration in mode of action of IsCT resulting its translocation into bacterial cytoplasm and inhibition of macromolecular syntheses

Jitendra K. Tripathi¹, Manoj Kathuria², Amit Kumar¹, Kalyan Mitra² and Jimut K. Ghosh¹#

¹Molecular and Structural Biology Division ²Electron Microscopy Unit CSIR-Central Drug Research Institute, Sector 10, Jankipuram Extension, Sitapur Road Lucknow–226 031, India

[#]To whom correspondence should be addressed: Jimut K. Ghosh, Molecular and Structural Biology Division, CSIR-Central Drug Research Institute, Sector 10, Jankipuram Extension, Sitapur Road, Lucknow–226 031, India,Tel: 091-522-2771940 (Ext.-4451); Fax: 091-522-2771941, E-mail: jighosh@yahoo.com; jk_ghosh@cdri.res.in.

Supplemental Tables

Supplementary Table S1. Thysicoencinical parameters of ise 1 and its analogs						
	µ−rel H	μH	<h></h>	Aliphatic	GRAVY	
Peptides	(Mean relative	(Mean H	(Mean	Index		
	H moment)	moment)	Hydrophobicity)			
IsCT	0.81	5.13	2.15	150.00	0.777	
E7K-IsCT	0.83	5.25	2.03	150.00	0.746	
I9K-IsCT	0.59	3.71	0.72	120.00	0.131	
E7K,I9K-IsCT	0.61	3.83	0.59	120.00	0.100	

Supplementary Table S1: Physicochemical parameters of IsCT and its analogs

Peptide	% Helix	% Beta	% Turn	% Random				
A. SSE of peptides in PBS (pH 7.4)								
IsCT	0	39.0	0.0	61.0				
E7K- IsCT	0	32.7	7.3	60.0				
I9K- IsCT	0	25.5	3.0	71.5				
E7K,I9K- IsCT	0	27.0	5.7	67.3				
B. SSE of peptides in presence of PC/Chol lipid vesicles								
IsCT	28.3	20.8	18.0	32.9				
E7K- IsCT	15.3	30.1	13.9	40.7				
I9K-IsCT	0.0	24.8	5.3	69.9				
E7K,I9K- IsCT	0.0	25.5	10.7	64.0				
C. SSE of peptides in presence of PC/PG lipid vesicles								
IsCT	29.2	21.2	17.8	31.0				
E7K- IsCT	25.9	20.1	19.7	34.3				
I9K- IsCT	0.0	32.2	4.5	63.2				
E7K,I9K- IsCT	0.0	24.7	8.7	66.6				
*Data analyzed by Structure Estimati	y-JASCO CD on) software.	, Model Nar	me-J-815, SS	E (Secondary				

Supplementary Table S2: Secondary Structure Estimation (SSE)* of IsCT and its analogs

Supplemental Figure:



Supplementary Figure S1: Helical wheel projections of IsCT and its analogues. Substituted amino acids are underlined





Supplementary Figure S2: Intrinsic Tryptophan fluorescence of peptides alone (A) and of PC/Chol and PC/PG alone (B), in PBS. Concentration of the each peptides used was \sim 2.0 μ M and lipid concentration was \sim 450 μ M.



Supplementary Figure S3: Change in HT voltage of the buffer as well different peptides in

PBS, pH 7.4. Concentration of the each peptides used was \sim 43 μ M.



Supplementary Figure S4: Quenching of NBD fluorescence of *E. coli ATCC 25922* bacteria bound to NBD-labeled IsCT or E7K,I9K-IsCT as marked in the Fig. by trypan blue. *E. coli* $(1x10^{6} \text{ CFU/ml})$ were incubated with or without NBD-labeled peptides (either NBD-labeled IsCT or E7K,I9K-IsCT at 2.0µM peptide concentration), washed, and analyzed by flow cytometry; Symbols: with NBD-labeled peptide, green histograms and without peptide, black histograms. *E.coli* bacteria treated with each of the NBD-labeled peptides were separately incubated with Trypan Blue (1mg/ml) for 10 min at 37°C and further analyzed by flow cytometry (pink histograms).