

Erratum: Reviewing Biophysical and Cell Biological Methodologies in Cell-Penetrating Peptide (CPP) Research (Volume 24, Issue 3)

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Please note that the following table was printed incorrectly in *Critical Reviews in Therapeutic Drug Carrier Systems*, Volume 24, Issue 3, on page 206. There should be a connection between the 'K' in the sequence on the second line and the 'A' in the sequence on the third line. Below is a corrected version of Table 1.

Table 1. Name, Amino Acid Sequence, Origin and Representative Studies of Reviewed CPPs

CPP	Sequence	Origin	Biological studies	Biophysical studies
hCT(9-32)	LGTYTODFNKHFHTFPQTAIGYGAP-amide	human calcitonin	[16, 23, 107]	[88, 89, 108, 211]
hCT(9-32)-br	LGTYTODFNKHFHTFPQTAIGYGAP-amide	human calcitonin	[16, 147]	
KLA1	AFGVGPDEVKRRKKKP-amide KLALKLALKAWKAALKLA-amide	NLS of SV-40 large T antigen amphipathic model peptide	[166]	
MAP	KLALKLALKLAKAALKLA-amide	amphipathic model peptide	[3, 149]	
MPG	GALFLGLGAAGSTMGAWSQPKSRKV-Cya	fusion peptide of HIV-1 gp41 and the NLS of SV40 large T-antigen	[6, 7]	[96]
pAntip*	ROIKWFQNRMRKWKK	third helix of the DNA binding domain of Antennapedia, a <i>Drosophila</i> transcription factor	[1, 3, 26, 27, 138, 140, 149, 169]	[26, 27, 40-42, 47, 56, 58, 66-69, 71, 77, 79, 89, 103, 140, 210]
Pep-1	KETWWTWTEWSQPKSRKV-Cya	hydrophobic tryptophane-rich motif fused to the NLS of SV-40 large T antigen connected by a spacer	[5]	[44, 96]
pVEC	LLILRRIRIKOAHASK-amide	murine vascular endothelial cadherin	[163]	[89]
SAP	VLPPPVRLPPVRLPPP	modified maize zein sequence	[16]	
SynB3	RRLYSRRRF	protegrin 1, an antimicrobial peptide	[66]	[66]
Tat(49-57)*	RKKRRQRRR	human immunodeficiency virus 1	[2, 3, 10, 17, 18, 22, 24, 25, 137-139, 144, 146, 148, 161, 169, 175, 179, 208]	[70, 57, 56, 24, 70, 175]
Transportan	GWTLNSAGYLLGKINLKALAALAKISIL-amide	neuropeptide galanin coupled to the wasp venom peptide mastoparan via Lys	[3]	[42, 47, 67, 95]
TP10	AGYLLGKINLKALAALAKKIL-amide	modification of Transportan	[186]	
VP22	DAATATGRSAASRPTERRPAPARSAPRRPVD	herpes simplex virus	[164, 167]	

*minimal sequence required for uptake; frequently sequences elongated at their C- or N-terminus were used.