

## Pe1b

<b>Cat. No.:</b>	HY-P5809
<b>Molecular Formula:</b>	C <sub>175</sub> H <sub>237</sub> N <sub>47</sub> O <sub>49</sub> S <sub>6</sub>
<b>Molecular Weight:</b>	3975.43
<b>Sequence:</b>	Glu-Cys-Arg-Tyr-Trp-Leu-Gly-Gly-Cys-Ser-Lys-Thr-Gly-Asp-Cys-Cys-Glu-His-Leu-Ser-Cys-Ser-Pro-Lys-Trp-His-Trp-Cys-Val-Trp-Asp-Gly-Thr-Phe (Disulfide bridge: Cys2-Cys16, Cys9-Cys21, Cys15-Cys28)
<b>Sequence Shortening:</b>	ECRYWLGGSCKTGDCCEHLSCSPKWHWCVWDGTF (Disulfide bridge: Cys2-Cys16, Cys9-Cys21, Cys15-Cys28)
<b>Target:</b>	Sodium Channel
<b>Pathway:</b>	Membrane Transporter/Ion Channel
<b>Storage:</b>	Please store the product under the recommended conditions in the Certificate of Analysis.

### BIOLOGICAL ACTIVITY

<b>Description</b>	Pe1b ( $\mu$ -TrTx-Pe1b) is a selective Na <sub>v</sub> 1.7 inhibitor with an IC <sub>50</sub> of 167 nM <sup>[1]</sup> .
<b>IC<sub>50</sub> &amp; Target</b>	IC <sub>50</sub> : 167 nM (Na <sub>v</sub> 1.7), 696 nM (Na <sub>v</sub> 1.6), 3.49 $\mu$ M (Na <sub>v</sub> 1.2), $\approx$ 10 $\mu$ M (Na <sub>v</sub> 1.5) <sup>[1]</sup>

### REFERENCES

[1]. Rupasinghe DB, et al. Mutational analysis of ProTx-I and the novel venom peptide Pe1b provide insight into residues responsible for selective inhibition of the analgesic drug target Na<sub>v</sub>1.7. *Biochem Pharmacol.* 2020 Nov;181:114080.

**Caution: Product has not been fully validated for medical applications. For research use only.**

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