

Insecticidal toxin LaIT1, recombinant venom peptide

(Liocheles australasiae)

Catalogue number: VP0007, 50 μg

(0.5 mL at 0.1 mg/mL)

Description

Insecticidal LalT1 venom peptide is a recombinant peptide purified from *Escherichia coli* that was originally isolated from the venom of *Liocheles australasiae* (Dwarf wood scorpion). The endogenous Insecticidal LalT1 peptide affects the activity of both ryanodinesensitive calcium-release channels RyR1 and RyR2. This venom peptide binds to different sites on the RyRs channels with high-affinity, mediating the full openings of these channels. Matsushita *et al.* described that insecticidal LalT1 venom peptide has insect toxicity activity against crickets but no toxicity was observed against mice, suggesting that the effect of this toxin is insect-selective. The recombinant peptide is provided in 50 mM NaHepes buffer, pH 7.5, 300 mM NaCl, at a 0.1 mg/mL concentration.

Purity

Insecticidal LaIT1 venom peptide is produced recombinantly and subjected to a variety of highly stringent purification protocols to reach a degree of purity > 90%, as evaluated by SDS-PAGE and ESI-Q-ToF-MS.

Recombinant Peptide sequence

DFPLSKEYETCVRPRKCQPPLKCNKAQICVDPKKGW

Specifications

Peptide Length	36 aa
Molecular weight	4207 Da
Number of Cys	4
Disulfide bonds	Cys ¹¹ -Cys ²³ , Cys ¹⁷ -Cys ²⁹
Source	Recombinant peptide from Liocheles australasiae
Format provided	Liquid
Uniprot Access	POC5F2
PDB Code	2LDS

Storage Temperature

Insecticidal LalT1 venom peptide should be stored at 4°C and is stable for 12 months.

Reference

Biological and biochemical properties of this peptide are described in Matsushita, N. *et al.*, Toxicon 50 (6), 861-867 (2007).

Quality Control Assays

Purity

Recombinant Insecticidal LalT1 venom peptide is >90% pure as judged by SDS polyacrylamide gel electrophoresis followed by BlueSafe staining (MB15201).

Molecular weight determination

To confirm molecular weight, oxidation pattern, molecular integrity and degree of purification, the recombinant peptide was analysed through ElectroSpray Ionization Quadrupole Time-of-Flight Mass Spectrometry (ESI-Q-ToF-MS) using a Synapt G2 HDMS (Waters) instrument. The resulted mass spectra was deconvoluted using MassLynx software and the obtained mass was compared with the theoretical peptide mass considering that all cysteine residues are oxidized.

V2101

Certificate of Analysis		
Test	Result	
Peptide purity	Pass	



Patrícia Ponte Senior Manager, Quality Systems

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