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Print Date: Jan 31st 2025

Certificate of Analysis

www.tocris.com

Product Name: LDV

CAS Number: 1155866-55-9

Catalog No.: 7020 Batch No.: 5

1. PHYSICAL AND CHEMICAL PROPERTIES

Batch Molecular Formula: Batch Molecular Weight: Physical Appearance: Counter Ion: Solubility: Storage: Peptide Sequence: $C_{48}H_{70}N_{10}O_{12}$ 979.13 White lyophilised solid TFA Soluble to 1 mg/ml in 0.01M PBS (pH 7.4) Store at -20°C $O_{10}H_{-}Leu-Asp-Val-Pro-Ala-Ala-Lys}$

2. ANALYTICAL DATA

HPLC:

Mass Spectrum:

Shows 99.8% purity Consistent with structure

3. AMINO ACID ANALYSIS DATA

Amino	Acid Theoreti	cal Actual	Amino	Acid Theoreti	cal Actual
Ala	2 00	1.06	LVC	1 00	1 01

Ala	2.00	1.96	Lys	1.00	1.01
Arg			Met		
Asx	1.00	0.99	Phe		
Cys			Pro	1.00	1.02
Glx			Ser		
Gly			Thr		
His			Trp		
lle			Tyr		
Leu	1.00	0.99	Val	1.00	1.01

Caution - Not Fully Tested • Research Use Only • Not For Human or Veterinary Use

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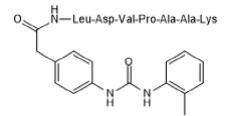
Description:

LDV is a $\alpha_4\beta_1$ integrin (VLA-4) ligand (K_d ~ 12 nM). Non-fluorescent derivative of LDV FITC (Cat. No. 4577).

Physical and Chemical Properties:

Batch Molecular Formula: C₄₈H₇₀N₁₀O₁₂ Batch Molecular Weight: 979.13 Physical Appearance: White Iyophilised solid

Peptide Sequence:



Storage: Store at -20°C

Solubility & Usage Info:

Soluble to 1 mg/ml in 0.01M PBS (pH 7.4)

This product is supplied in lyophilized form. It may appear as a solid, gel or film and be very hard to visualize. Solutions should be made by adding solvent directly to the vial. The vial should then be vortexed vigorously to ensure the product has completely dissolved.

Counter Ion: TFA

Stability and Solubility Advice:

Some solutions can be difficult to obtain and can be encouraged by rapid stirring, sonication or gentle warming (in a 45-60°C water bath).

Peptides in solution are much less stable than in lyophilized form. This is especially true for peptides whose sequences contain amino acids such Cys, Met,Trp, Asn, Gln, and N-terminal Glu.

Therefore we recommend storing peptides in solution for as short a time as possible. Avoid repeated freeze thaw cycles by dividing the peptide solution into aliquots and storing the aliquots at -20°C. Any portion of an aliquot unused after thawing should be discarded.

Peptides stored in solution can occasionally be susceptible to bacterial degradation. We recommend using sterile solutions or passing the peptide solution through a 0.2 μ m filter to remove potential bacterial contamination whenever possible.

References:

Chigaev *et al* (2009) Real-time analysis of conformation-sensitive antibody binding provides new insights into integrin conformational regulation. J.Biol.Chem. **284** 14337. PMID: 19251697.

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