



Ozcanium Analytics
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Certificate of Analysis

CJC-1295 + Ipamorelin 5mg / 5mg

Compound Mix : CJC-1295 + Ipamorelin

Lot Number : APL-CJCIP10-2608-01

Analysis date : 27/08/2026

Method : UPLC-MS/MS

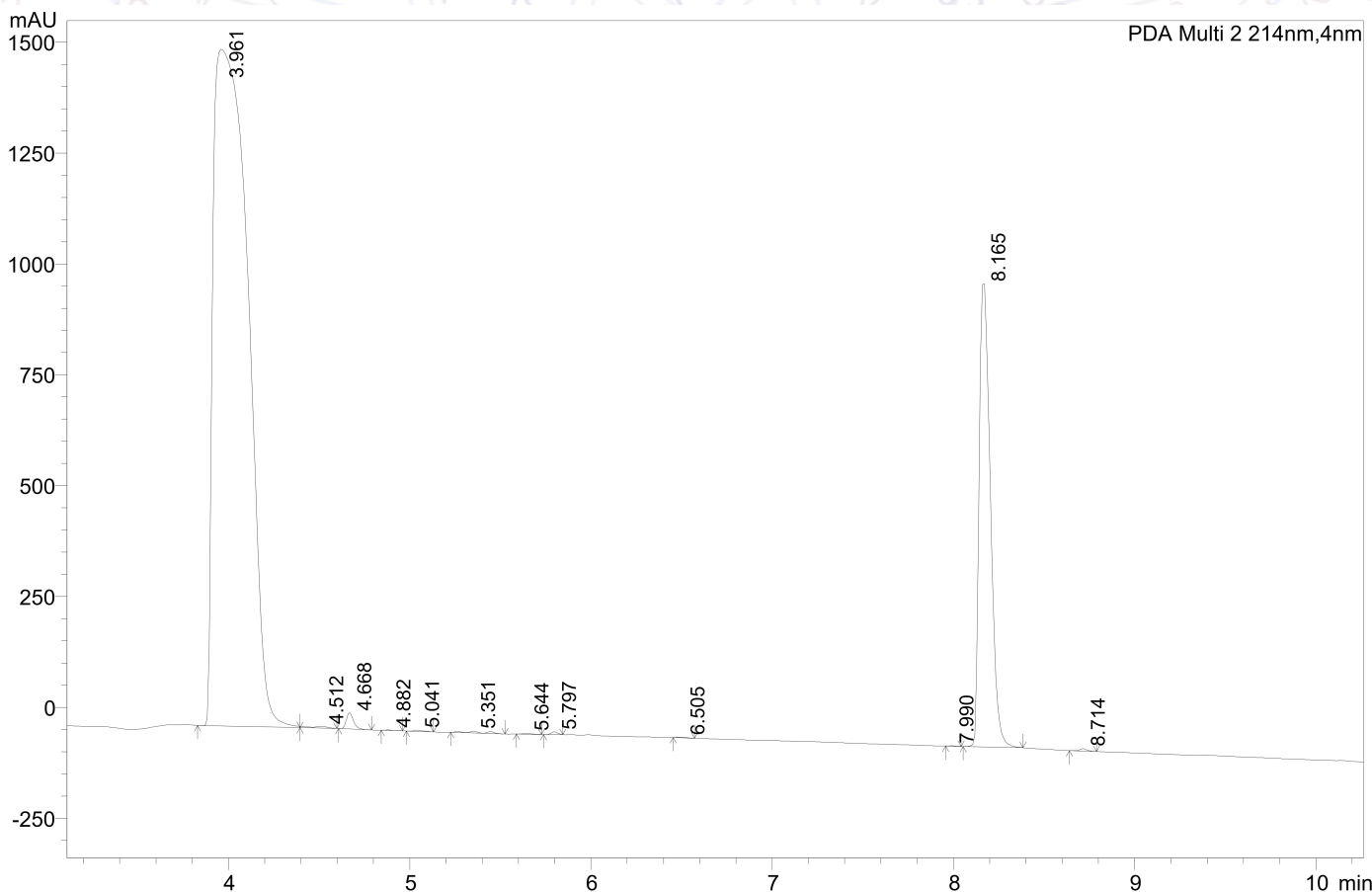
Verification Code : OZ-HPLCMS-JKJ2

Purity : 99.16%



Net Content : Ipamorelin: 5.10mg
CJC-1295 No DAC: 5.29mg
Client : Australian Peptide Labs

Ultra Performance Liquid Chromatography (UPLC-UV) Chromatogram



PEAK LIST

Peaks Detected: 12

#	Time (min)	Area	%Area	Assignments
1	3.961	20192466	81.054	Ipamorelin
2	4.512	24278	0.097	
3	4.668	100353	0.403	
4	4.882	2352	0.009	
5	5.041	9642	0.039	
6	5.351	26447	0.106	
7	5.644	7607	0.031	
8	5.797	16004	0.064	
9	6.505	5569	0.022	
10	7.990	2303	0.009	
11	8.165	4511709	18.110	CJC-1295 No DAC
12	8.714	13692	0.055	

Analysis by Ibrahim A. Ozcan,
Lead Chemist Ozcanium Analytics
Signature

Report Compiled by Zein S. Sabini
29/08/2026

Note: Sample may contain salts and sugars, as standard additives. These are not considered impurities and are not included in the peak list



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Ipamorelin

PubChem CID : 9831659
CAS Number : 170851-70-4

Liquid Chromatography Mass Spectrometry (LC-MS) Identity Analysis

Identity confirmation with SIM

Analysis date : 27/08/2026

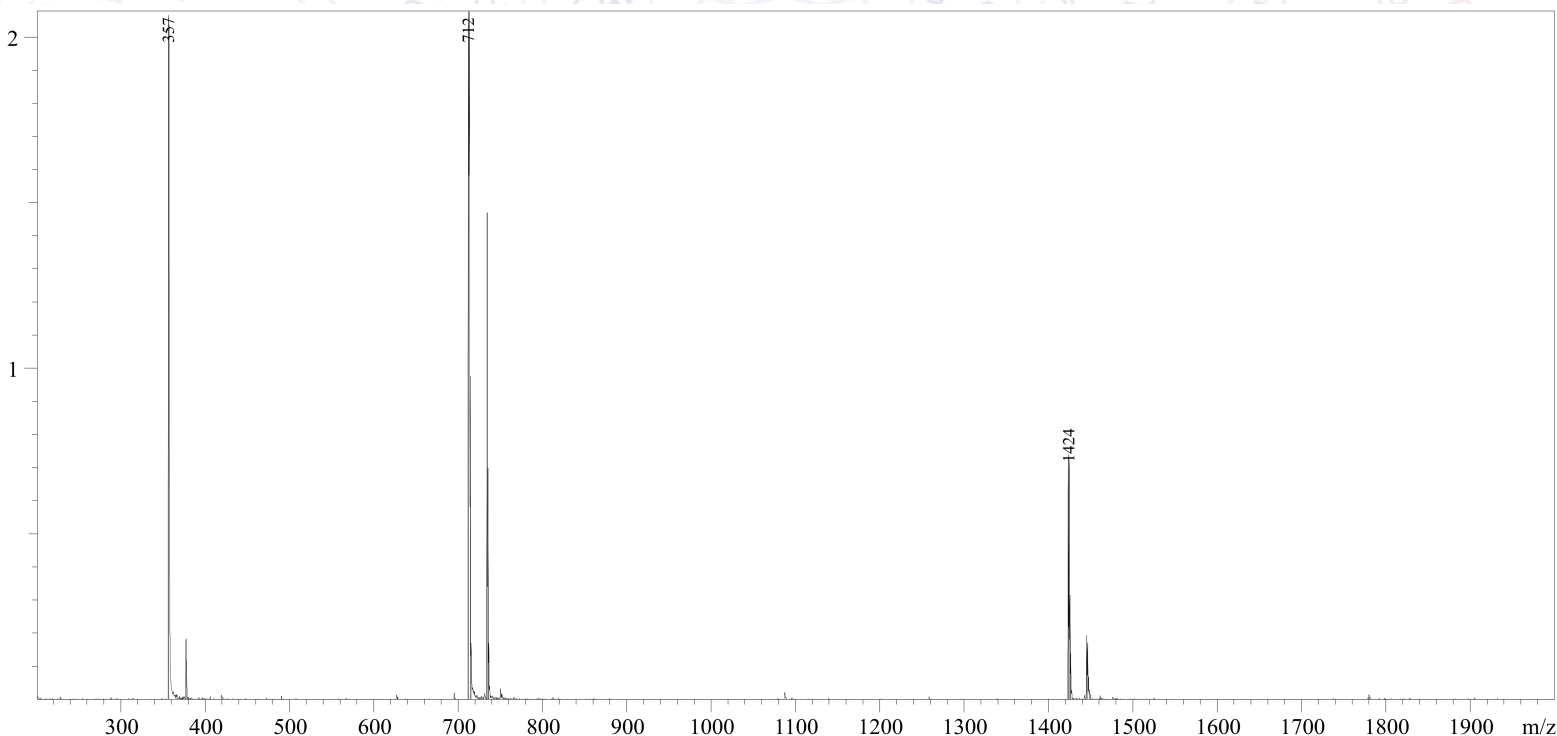
Monitored Ion Channels (SIM):

Monitored (m/z):	Detected (m/z):
356.9	357.10
712.9	712.40
734.9	735.05

Detection Assignment: **Ipamorelin**

Collected Mass Spectrum

Intensity 10⁶



Note : Monoisotopic m/z values are not easily seen in full spectrum view for larger molecules and peptides.

The recorded fragmentation pattern may differ between systems, and vary based on method.

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CJC-1295 No DAC

PubChem CID : 91976842
CAS Number : 446036-97-1

Liquid Chromatography Mass Spectrometry (LC-MS) Identity Analysis

Identity confirmation with SIM

Analysis date : 27/08/2026

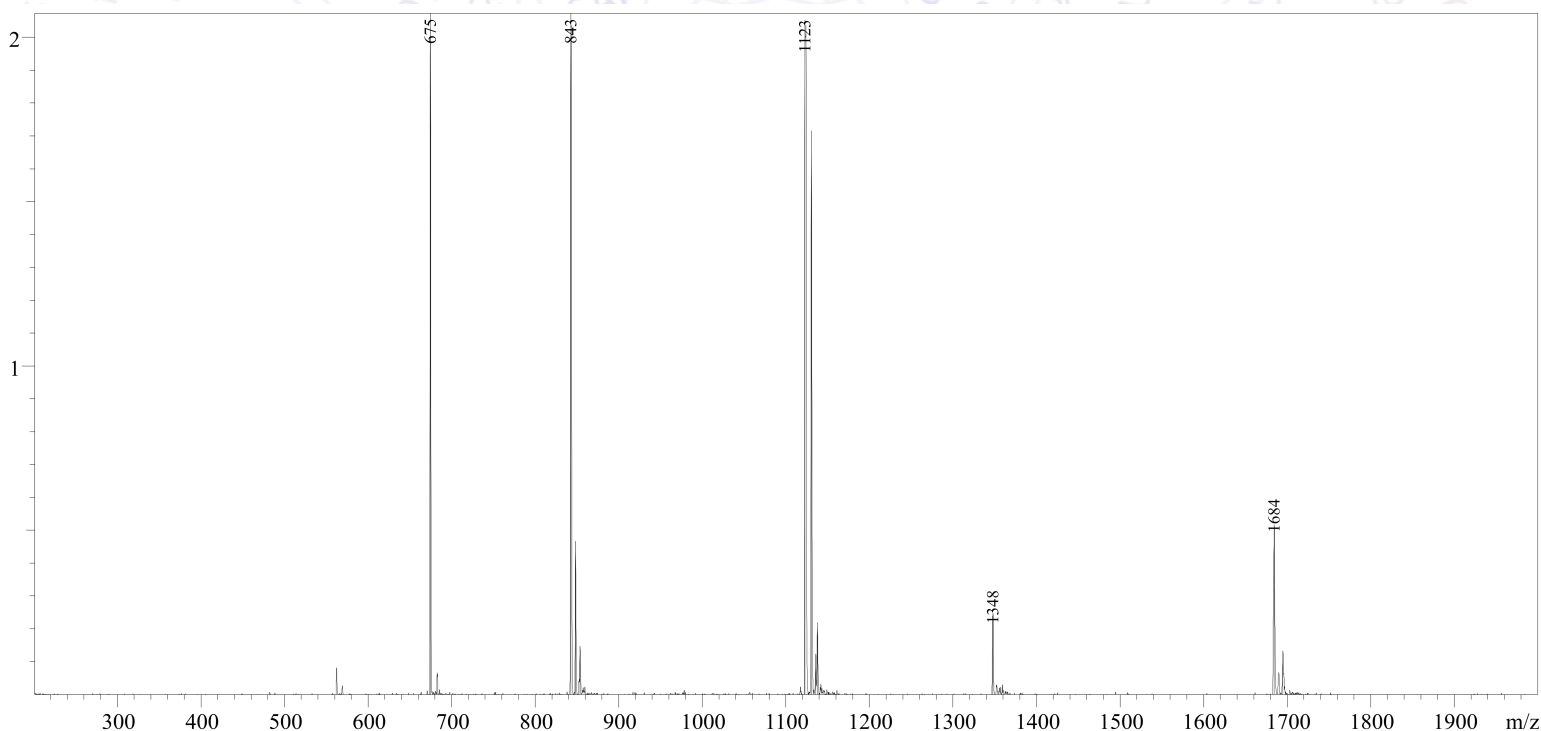
Monitored Ion Channels (SIM):

Monitored (m/z):	Detected (m/z):
842	842.60
854	854.10
1123	1123.35

Detection Assignment: **CJC-1295 No DAC**

Collected Mass Spectrum

Intensity 10⁶



Note : Monoisotopic m/z values are not easily seen in full spectrum view for larger molecules and peptides.

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