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ABN: 14 754 514 569
Ozcaniumanalytics.com.au
+61 407 260 481

Certificate of Analysis

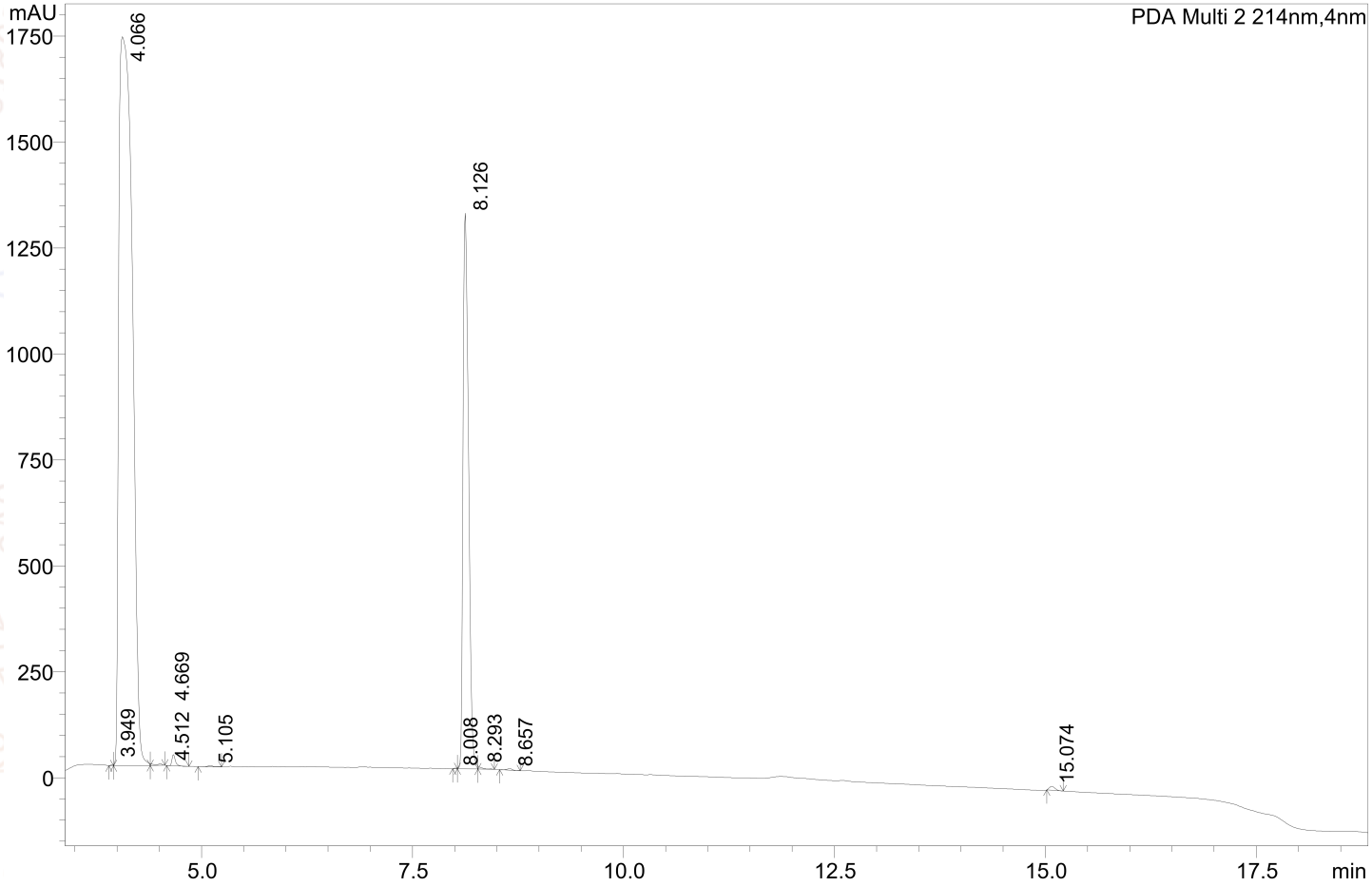
CJC-1295 + Ipamorelin 5mg / 5mg

Compound : CJC-1295 + Ipamorelin
Lot Number : 08/09/2026
Analysis date : 12/09/2026
Method : UPLC-MS/MS
Verification : OZ-HPLCMS-CYX8
Purity : 99.24%

Net Content : Ipamorelin: 5.10mg
CJC-1295 No DAC: 5.29mg
Client : Coregenyx Pty Ltd
Coregenyx.com



Ultra Performance Liquid Chromatography (UPLC-UV) Chromatogram



PEAK LIST

Peaks Detected: 10

#	Time (min)	Area	%Area	Assignments
1	3.949	5972	0.024	
2	4.066	19192037	75.851	Ipamorelin
3	4.512	22019	0.087	
4	4.669	68993	0.273	
5	5.105	14808	0.059	
6	8.008	3664	0.014	
7	8.126	5916942	23.385	CJC-1295 No DAC
8	8.293	20756	0.082	
9	8.657	14072	0.056	
10	15.074	42866	0.169	

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

Report Compiled by Zein S. Sabini
13/09/2026

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

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

Liquid Chromatography Mass Spectrometry (LC-MS) Identity Analysis

Identity confirmation with SIM **Analysis date** : 12/09/2026

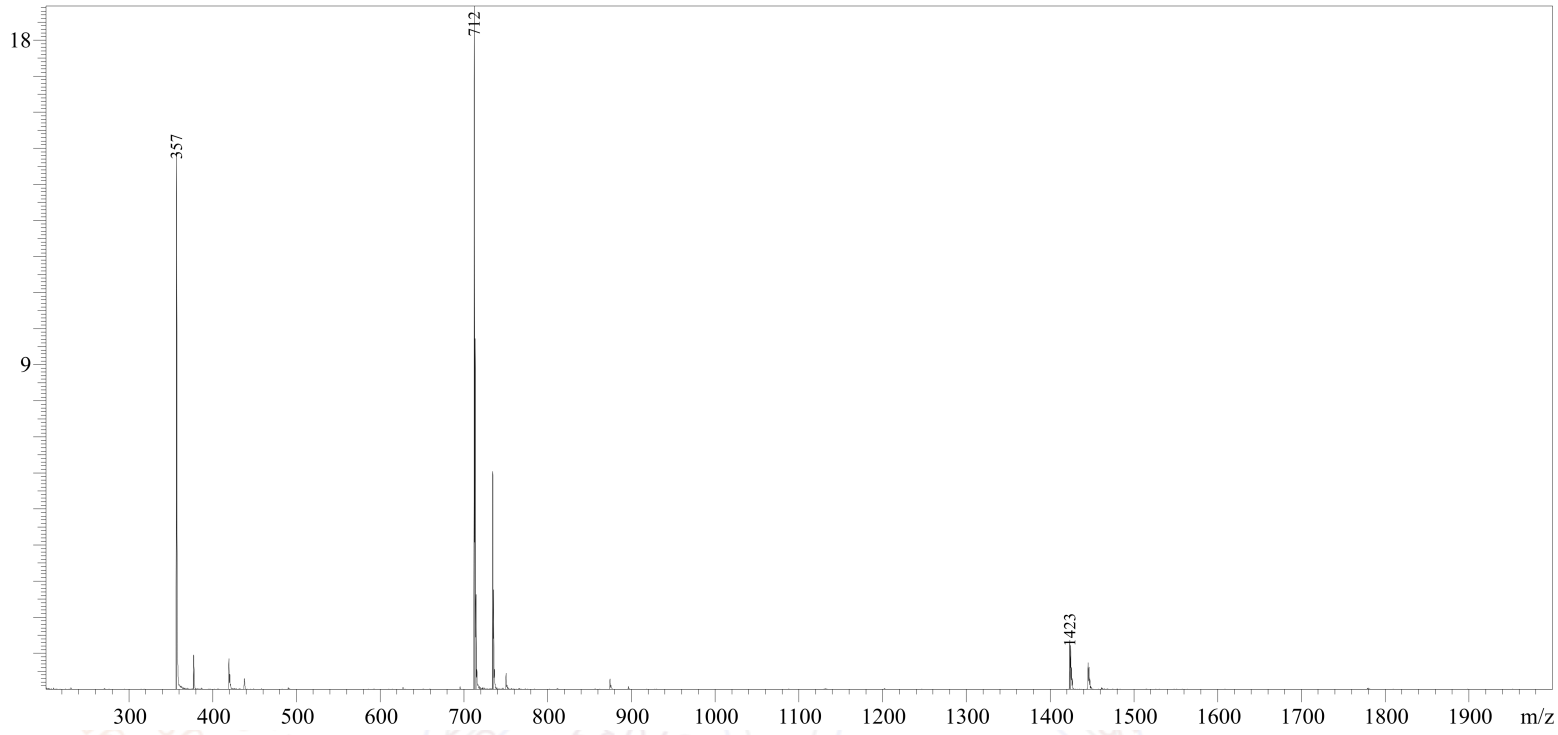
Monitored Ion Channels (SIM):

Monitored (m/z):	Detected (m/z):
356.9	357.10
712.9	712.45
734.9	734.65

Detection Assignment: Ipamorelin

Collected Mass Spectrum

Intensity 10^7



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.

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

Report Compiled by Zein S. Sabini
13/09/2026

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CJC-1295 No DAC

CAS Number : 446036-97-1

PubChem CID : 91976842

Liquid Chromatography Mass Spectrometry (LC-MS) Identity Analysis

Identity confirmation with SIM

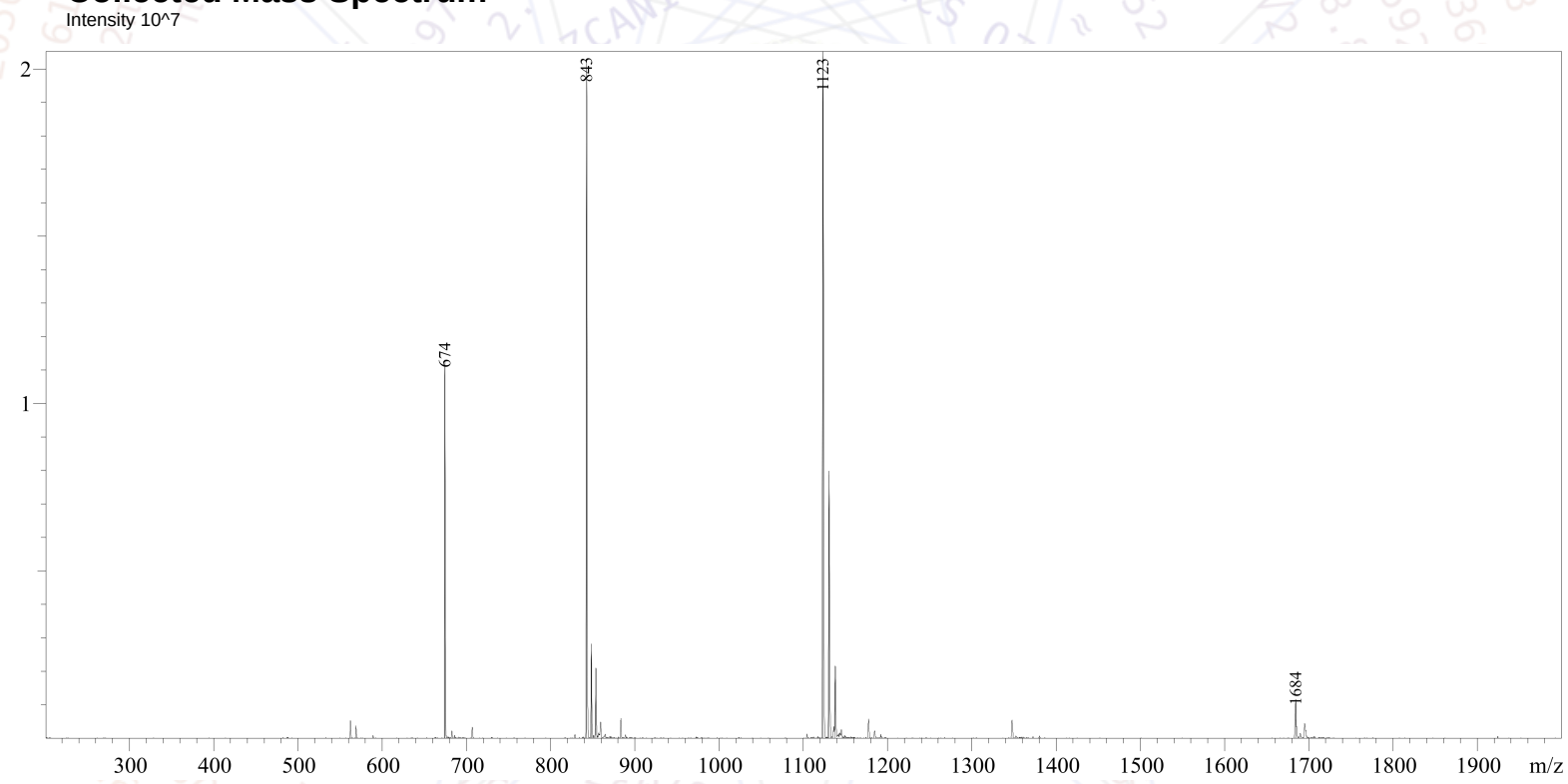
Analysis date : 12/09/2026

Monitored Ion Channels (SIM):

Monitored (m/z):	Detected (m/z):
842	842.55
854	854.20
1123	1123.10

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

Collected Mass Spectrum



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.

Analysis by Ibrahim A. Ozcan,
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Signature

Report Compiled by Zein S. Sabini
13/09/2026

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