

PRODUCT

CJC-1295 (No DAC)

Modified GRF (1-29), Mod GRF 1-29

CATALOG	VA-CJC1295NODAC-5MG
CAS	863288-34-0
FORMULA	C152H252N44O42
MOL. WEIGHT	3367.9 avg · 3365.89 monoisotopic
SEQUENCE	Tyr-D-Ala-Asp-Ala-Ile-Phe-Thr-Gln-Ser-Tyr-Arg-Lys-Val-Leu-Ala-Gln-Leu-Ser-Ala-Arg-Lys-Leu-Leu-Gln-Asp-Ile-Leu-Ser-Arg-NH2
FORM	Lyophilized powder
FILL WEIGHT	5 mg
CONTAINER	3 mL vial

BATCH

BATCH IDENTIFIER

VA-2512-9601

MANUFACTURED	2026-04-15
FILLED	2026-05-12
ANALYSED	2026-05-14
RETEST BY	2028-04-15
STORAGE	Refrigerated, 2-8 °C, protected from light

RESULTS

TEST / METHOD	SPECIFICATION	RESULT	STATUS
Appearance Visual inspection (Vantage Aminos)	White lyophilized powder	Conforms	PASS
Identity HPLC-UV-MS, positive polarity (+MS)	Theoretical mass ± 1.0 Da	3365.96 Da (theor. 3365.89)	PASS
Purity HPLC-UV, area % · DAD 209.9-288.3 nm	≥ 98.0 %	98.60 %	PASS
Quantity Quantitation as reported	90-110 % of label (5 mg)	5.026 mg (100.5 % of label)	PASS
Largest single impurity HPLC-UV, area %	Report	0.5 % at 6.88 min	REPORT
Water content Karl Fischer titration	≤ 8 % w/w	3.64 % w/w	PASS

HOW TO VERIFY THIS CERTIFICATE

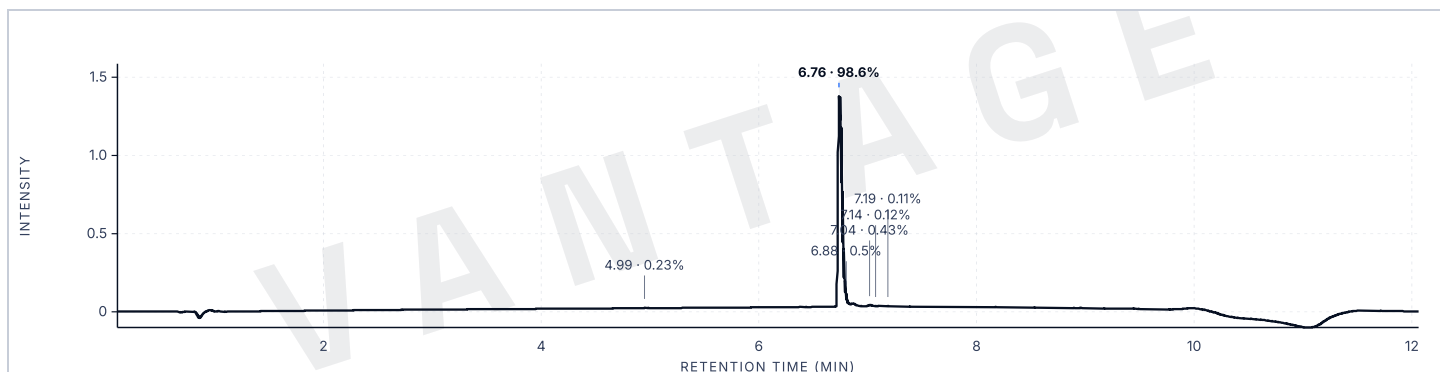
- 1

All units are shipped from the most recent batch. Confirm the compound and fill weight shown above match the product you ordered.
- 2

This certificate is published on the product page at <https://vantageaminos.co/product/cjc-1295-no-dac/>, alongside the certificate for every other batch of this product.
- 3

Any question about this analysis: contact support@vantageaminos.co quoting the batch identifier above.

HPLC CHROMATOGRAM



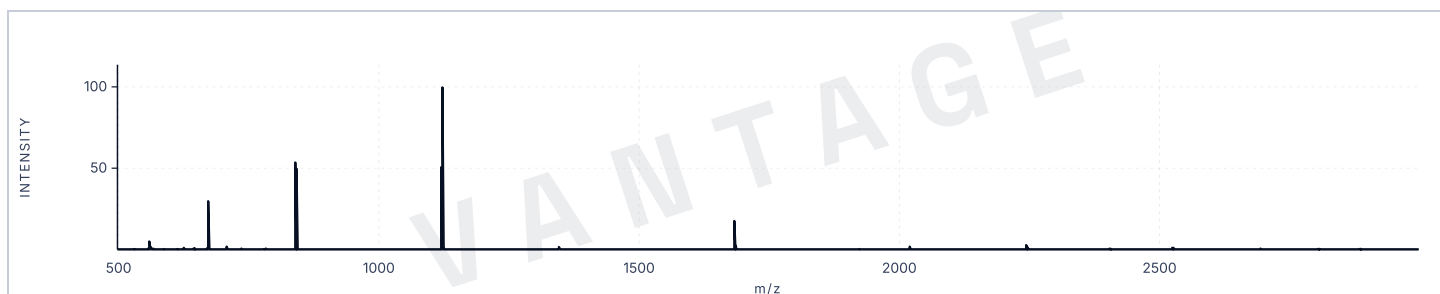
PEAK INTEGRATION

RT (MIN)	AREA	AREA %	ASSIGNMENT
4.99	8.50E+01	0.23	unassigned
6.76	3.66E+04	98.6	CJC-1295 no DAC
6.88	1.87E+02	0.5	unassigned
7.04	1.59E+02	0.43	unassigned
7.14	4.49E+01	0.12	unassigned
7.19	4.18E+01	0.11	unassigned

MASS SPECTROMETRY

Technique	HPLC-UV-MS (as printed on certificate)
Ionisation	Positive polarity (+MS)
Scan range	m/z 497.7 – 2997.6 (as plotted)
Theoretical mass	3365.89 Da
Observed mass	3365.96 Da
Deviation	+0.07 Da

MASS SPECTRUM



Full-scan mass spectrum. Base peak m/z 1123.32

HPLC METHOD

Technique	Reverse-phase HPLC with UV detection, MS confirmation
Detection	209.9-288.3 nm (DAD range, not a single wavelength)