

Certificate of Analysis



Grail Formula B.V.

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Sample Identification

Sample Name	Hexarelin 3 mg
Batch Number	GF-HEXAR3-B241
Date Published	2026-09-20 13:01

Results for LYO-0733

Peptides	Result	Unit	Uncertainty	Acceptable Range
Hexarelin Assay Peptide Screening 0.1% TFA	3.71	mg	[± 0.02]	
Hexarelin Purity Peptide Screening 0.1% TFA	99.2	%	[± 0.5]	
Hexarelin Identification by Spectrum (FTIR) Peptide Screening 0.1% TFA	994		[± 5]	
Hexarelin Identification by RT Peptide Screening 0.1% TFA	0.0		[±]	
Microbiology	Result	Unit	Uncertainty	Acceptable Range
Total Aerobic Microbial Count USP <61>/Eur. Ph. 2.6.12. Plate Count Method	0	CFU/g	[±]	
Total Yeast and Mold Count USP <61>/Eur. Ph. 2.6.12. Plate Count Method	0	CFU/g	[±]	
Bacterial Endotoxin Chromogenic USP<85>/ Eur. Ph. 2.6.14. Bacterial Endotoxin Chromogenic Test	0.040	EU/mg	[± 0.001]	
Elemental Impurities	Result	Unit	Uncertainty	Acceptable Range
Arsenic Elemental Impurities screening USP (232) / Ph. Eur. 5.20 / 2.4.20	< 0.001	ppm		
Cadmium Elemental Impurities screening USP (232) / Ph. Eur. 5.20 / 2.4.20	< 0.001	ppm		
Quicksilver Elemental Impurities screening USP (232) / Ph. Eur. 5.20 / 2.4.20	< 0.001	ppm		
Lead Elemental Impurities screening USP (232) / Ph. Eur. 5.20 / 2.4.20	< 0.001	ppm		
Nickel Elemental Impurities screening USP (232) / Ph. Eur. 5.20 / 2.4.20	< 0.001	ppm		
Vanadium Elemental Impurities screening USP (232) / Ph. Eur. 5.20 / 2.4.20	< 0.001	ppm		
Cobalt Elemental Impurities screening USP (232) / Ph. Eur. 5.20 / 2.4.20	< 0.001	ppm		
Mass Spectrometry	Result	Unit	Uncertainty	Acceptable Range
Molecular Ion Mass Identification (MS Deconvolution) Mass Spectrometry Identity	887	Da	[± 1]	

	Method Specification	
Determination of identity, content and purity of Hexarelin		
<i>Document number</i> HEX_008_2026	<i>Superseded document</i> -	<i>Number of pages</i> 4

1. Content Assessment

1.1. Instrumentation

Module	Name	Serial Number
System Controller	Shimadzu CBM-40 Lite	L221226351398
Degassing Unit	Shimadzu DGU-403	NA
Pump	Shimadzu LC-40B XR	L22146350580
Autosampler	Shimadzu SIL-40C XR	L22216351622
Colum Thermostat	Shimadzu CTO-40S	L22236351602
PDA Detector	Shimadzu SPD-M40	L22276352808
SQ MS Detector	Shimadzu LCMS-2050	O12476200760

1.2. Chromatographic conditions

Chromatographic conditions	
Eluent A	0.1% DFA in Water (HPLC, Gradient Grade)
Eluent B	0.1% DFA in Acetonitrile (HPLC, Gradient Grade)
Flow rate	0.8 mL/min
Program	Gradient elution
Injection volume	2 µL
Colum Temperature	60°C
Column	Waters XSelect CSH C18, 100x2.1mm 2.5µm
Detection wavelength	280nm

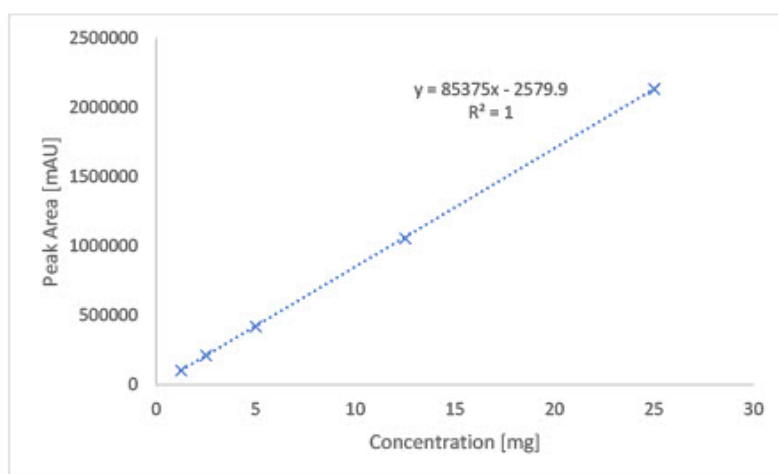
Gradient Program		
Time [min]	A [%]	B [%]
1	98	2
13.99	35	65
14	5	95
15	5	95
15.01	98	2
19	end	

1.3. Sample preparation

Whole amount of container was dissolved in 2mL of water (LCMS Grade). 100 µL of sample was transferred to HPLC vial and diluted by 900 µL water (LCMS Grade) and submitted for analysis.

1.4. Calibration curve

Calibration curve detail	
Quantitative method	External Standard
Calibration Type	Linear
Number of calibration points	5
Force through Zero	Disabled
Weighting Method	None



2. Purity assessment

2.1 Instrumentation

Module	Name	Serial Number
System Controller	Shimadzu CBM-40 Lite	L221226351398
Degassing Unit	Shimadzu DGU-403	NA
Pump	Shimadzu LC-40B XR	L22146350580
Autosampler	Shimadzu SIL-40C XR	L22216351622
Colum Thermostat	Shimadzu CTO-40S	L22236351602
PDA Detector	Shimadzu SPD-M40	L22276352808
SQ MS Detector	Shimadzu LCMS-2050	O12476200760

2.2 Chromatographic conditions

Chromatographic conditions	
Eluent A	0.1% DFA in Water (HPLC, Gradient Grade)
Eluent B	0.1% DFA in Acetonitrile (HPLC, Gradient Grade)
Flow rate	0.8 mL/min
Program	Gradient elution
Injection volume	2 µL
Colum Temperature	60°C
Column	Waters XSelect CSH C18, 100x2.1mm 2.5µm
Detection wavelength	225nm

Gradient Program		
Time [min]	A [%]	B [%]
1	98	2
13.99	35	65
14	5	95
15	5	95
15.01	98	2
19	end	

2.3 Purity assesment

Purity of compound assesed by area normalization method, comparing area of each peak to sum of area of all peaks detected at wavelenght of 225 nm.

3. Identity Assessment

3.1 Instrumentation

Module	Name	Serial Number
System Controller	Shimadzu CBM-40 Lite	L221226351398
Degassing Unit	Shimadzu DGU-403	NA
Pump	Shimadzu LC-40B XR	L22146350580
Autosampler	Shimadzu SIL-40C XR	L22216351622
Column Thermostat	Shimadzu CTO-40S	L22236351602
PDA Detector	Shimadzu SPD-M40	L22276352808
SQ MS Detector	Shimadzu LCMS-2050	O12476200760

3.2 Chromatographic conditions

Chromatographic conditions	
Eluent A	0.1% DFA in Water (HPLC, Gradient Grade)
Eluent B	0.1% DFA in Acetonitrile (HPLC, Gradient Grade)
Flow rate	0.8 mL/min
Program	Gradient elution
Injection volume	2 µL
Column Temperature	60°C
Column	Waters XSelect CSH C18, 100x2.1mm 2.5µm
Detection	MS 220-2000 Da

Gradient Program		
Time [min]	A [%]	B [%]
1	98	2
13.99	35	65
14	5	95
15	5	95
15.01	98	2
19	end	

3.3 Molecular Ion Mass evaluation

Molecular ion mass was determined by deconvolution of multiply charged ESI-MS spectra to calculate the average neutral (zero-charge) molecular mass by equation:

$$M(\text{neutral}) = (z_i((mz_i) - H)) - ME$$

Where:

mz_i - Measured mass of charged particle

z_i - charge

H - proton mass (1.0076 Da)

ME - mass error

Analysis Report

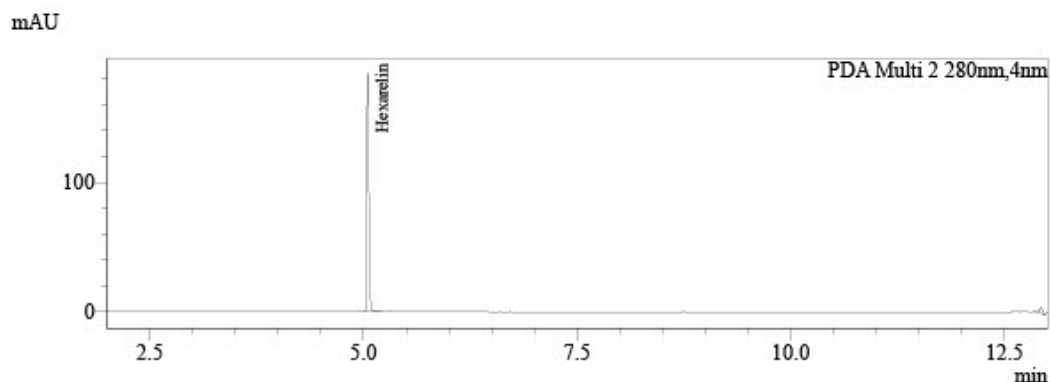
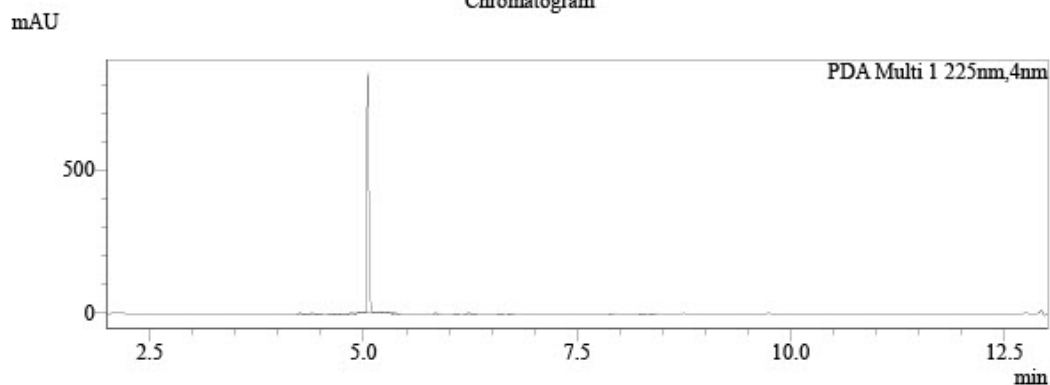


Analysis of quantity of active ingredient by UHPLC with MS detection

Sample Information

Injection Volume : 2
Data File : LYO-0733__026.lcd
Method File : Peptide Screening_DFA_Group B.lcm
Date Acquired : 9/1/2026 12:47:21 AM

Chromatogram



Peak Table

Peak#	Name	Ret. Time	Area	Area%	Conc.	Unit
1		4.259	410	0.028	0.000	
2		4.395	693	0.048	0.000	
3		4.486	110	0.008	0.000	
4		4.622	244	0.017	0.000	
5		4.691	87	0.006	0.000	
6		4.797	257	0.018	0.000	
7		4.866	1236	0.085	0.000	
8		4.921	186	0.013	0.000	
9		4.959	2127	0.146	0.000	
10		5.048	1440194	99.212	0.000	
11		5.253	655	0.045	0.000	
12		5.311	2063	0.142	0.000	
13		5.368	574	0.040	0.000	
14		5.841	758	0.052	0.000	
15		6.078	215	0.015	0.000	
16		6.233	927	0.064	0.000	

Peak#	Name	Ret. Time	Area	Area%	Conc.	Unit
17		6.594	154	0.011	0.000	
18		6.713	170	0.012	0.000	
19		7.890	105	0.007	0.000	
20		8.262	182	0.013	0.000	
21		8.382	293	0.020	0.000	
Total			1451639	100.000		

Peak Table

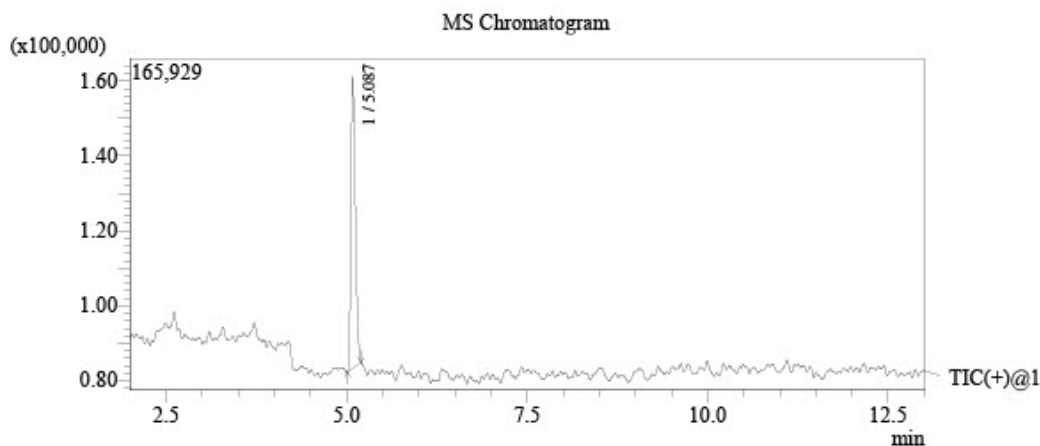
PDA Ch2 280nm

Peak#	Name	Ret. Time	Area	Area%	Conc.	Unit
1	Hexarelin	5.048	314524	95.011	3.714	mg
2		12.928	12024	3.632	0.000	
3		14.214	4490	1.356	0.000	
Total			331038	100.000		

Analysis Report



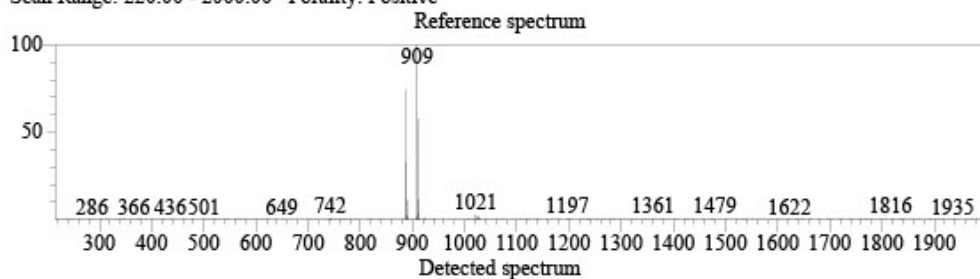
Identification of active ingredient by UHPLC Mass Spectrometr



Library Search

Base Peak: 909(24483)

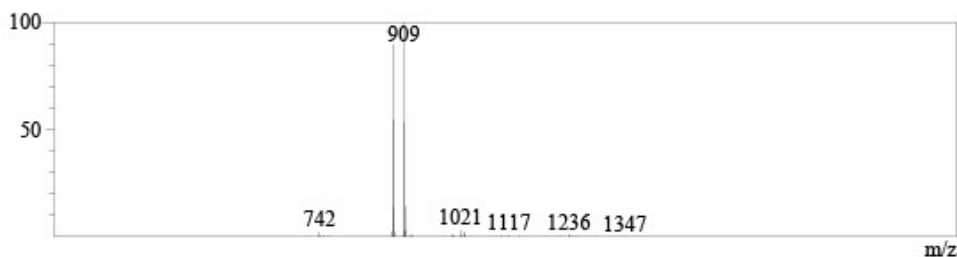
Scan Range: 220.00 - 2000.00 Polarity: Positive



Library: Peptide Screening_DFA.lib

Formula: $C_{47}H_{58}N_{12}O_6$ CAS: 140703-51-1 Mol.Weight: 887(0!=)

Compound Name: Hexarelin (examorelin)



Endotoxin Determination Report

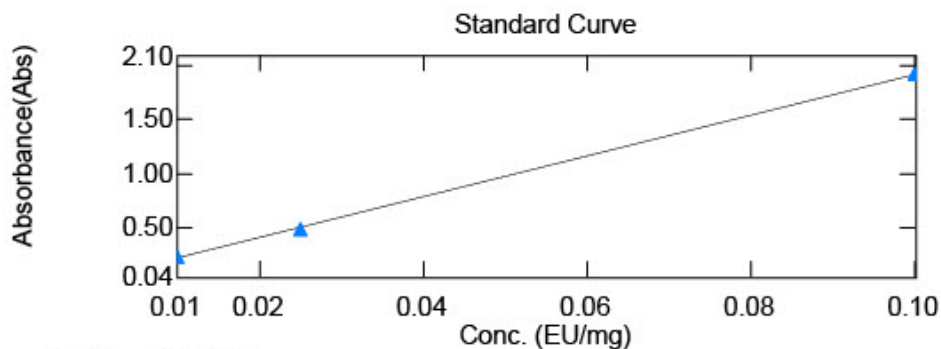


File Information

Filename: C:\UVVis-Data\Data\File_260831_145950.vqud
Date/Time: 08/31/2026 03:44:00 PM

Instrument Information

Instrument Type: UV-1900 Series
Model (S/N): UV-1900i (A12536253123)



$$y = 18.9790x + 0.0240870$$
$$r^2 = 0.99983$$

	Sample Name	Conc	Raw_WL545.0	Dilution Rate
1	LYO-0687	0.180	0.1945	20.00
2	LYO-0688	0.194	0.2080	20.00
3	LYO-0689	0.088	0.1075	20.00
4	LYO-0690	0.084	0.1039	20.00
5	LYO-0691	0.068	0.0884	20.00
6	LYO-0697	1.364	1.3188	20.00
7	LYO-0707	0.150	0.1667	20.00
8	LYO-0718	2.210	1.0729	40.00
9	LYO-0719	2.683	1.2971	40.00
10	LYO-0720	0.089	0.1087	20.00
11	LYO-0721	0.068	0.0890	20.00
12	LYO-0732	0.121	0.1390	20.00
13	LYO-0733	0.150	0.1662	20.00
14	LYO-0734	0.080	0.1003	20.00
15	LYO-0736	2.428	1.1761	40.00
16	LYO-0737	0.163	0.1787	20.00
17	LYO-0738	0.073	0.0937	20.00
18	LYO-0739	3.102	1.4960	40.00
19	LIQ-0066	0.072	0.7082	2.00

 LIQUILABS Analytical services	Method Specification	
Determination of bioburden of lyophilized samples		
Document number MIC_001_2025	Superseded document -	Number of pages 2

1. Instrumentation and chemicals

1.1. Instruments used

- Sterile Syringe 2mL Luer
- Sterile needles
- Ready made PCA Plate ROTI Aquatest
- Ready made Sab4 Plate ROTI Aquatest

1.2. Chemicals

Sterile physiological solution (0.9% NaCl)

2. Sample preparation and inoculation

2.1 Sample preparation

1. Fresh sterile needle and syringe was used for measuring exactly 2 mL of sterile physiological solution.
2. Needle was changed and by new needle rubber top of peptide container was penetrated and 2 mL of sterile physiological solution was dispensed.
3. Content of container was completely dissolved and left for 5 minutes to settle potentially created bubbles.
4. This procedure is repeated for two vials.

2.2 Total Aerobic microbial count inoculation and cultivation

1. By sterile needle 1 mL of solution was filled into the sterile syringe.
2. Needle was placed above the flame for few seconds to sterilize.
3. Consequently 1 mL of solution was poured into the ready to use sterile petri dish filled with PCA agar and petri dish was closed.
4. Proces was repeated for two petri dishes.
5. With sterile needle, 1 mL of sterile physiological solution was filled into the sterile needle and was inoculated onto one sterile petri dish filled with PCA agar as negative control sample.
6. Samples and negative control sample were placed in incubator at temperature 37°C for 120h.

2.3 Total Yeast and Mold count inoculation and cultivation

1. By sterile needle 1 mL of solution was filled into the sterile syringe.
2. Needle was placed above the flame for few seconds to sterilize.
3. Consequently 1 mL of solution was poured into the ready to use sterile petri dish filled with Sab4 agar and petri dish was closed.
4. Proces was repeated for two petri dishes.
5. With sterile needle, 1 mL of sterile physiological solution was filled into the sterile needle and was inoculated onto one sterile petri dish filled with Sab4 agar as negative control sample.
6. Samples and negative control sample were placed in incubator at temperature 25°C for 72h.

3. Evaluation of results

After incubation time, colonies are counted as cfu (colonies forming units) and result per 1g of sample is determined as:

$$CFU_{avg} = \frac{\sum CFU_n}{n}$$

CFU_{avg} = average CFU counted form n inoculations

CFU_n = CFU counted per inoculation

n = number of inoculations

$$CFU \text{ per gram} = \frac{CFU_{avg}}{m_s} * DF$$

CFU_{avg} = Average CFU counted from n inoculations

m_s = mass of sample (mg)

DF = Dilution factor

If negative control sample is evaluated as positive, process have to be repeated due to possible contamination in the process of inoculation or incubation.

Responsibles



Mr. Ján Galbavý
CEO

Analysis results relate only to the samples tested.

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