Instrument Method: Exploris_Aurora_25cm_max4uL_NormDIA_16mzst_90min_EndWash_BCS.meth

Thermo EASY-LC method print for Thermo

Sample pickup:

Volume [μ l] : 4.00 Flow [μ l / min] : 20.00 Volume [µl]

Sample loading:_

Volume [μ l] : 10.00 Flow [μ l / min] : (unspecified) Max. pressure [Bar] : 750.00

Gradient:

Time [mm:ss]	Duration [mm:ss]	Flow [nl/min]	Mixture [%B]
00:00	00:00	250.00	2.00
05:00	05:00	250.00	2.00
80:00	75:00	250.00	28.00
90:00	10:00	250.00	44.00
95:00	05:00	250.00	100.00
98:00	03:00	300.00	100.00
101:00	03:00	300.00	2.00
104:00	03:00	300.00	2.00
107:00	03:00	300.00	100.00
110:00	03:00	300.00	100.00

Pre-column equilibration:

Volume [μ l] : 0.00 Flow [μ l / min] : (unspecified)

Max. pressure [Bar] : 750.00

Analytical column equilibration:

Volume [µl] : 3.00 Flow [µl / min] : (unspecified) Max. pressure [Bar] : 750.00

Autosampler wash:

Flush volume [μ l] : 100.00

Method Summary

Method Settings

Application Mode: **Peptide**Method Duration (min): **110**

Global Parameters

Ion Source

lon Source Type: NSI
Spray Voltage: Static
Positive Ion (V): 2000
Negative Ion (V): 600
Sweep Gas (Arb): 0

Ion Transfer Tube Temp (°C): 280
Use Ion Source Settings from Tune: False

FAIMS Mode: Not Installed

MS Global Settings

Infusion Mode: Liquid Chromatography

Expected LC Peak Width (s): 25
Advanced Peak Determination: True

Default Charge State: 3

Internal Mass Calibration: EASY-IC™

Mode: Run Start

Experiment#1 [tSIM]

Start Time (min): 0 End Time (min): 110

Master Scan:

tSIM

Multiplex lons: False
Isolation Window (m/z): 620
Isolation Offset: Off

Orbitrap Resolution: 60000

Instrument Method: Exploris_Aurora_25cm_max4uL_NormDIA_16mzst_90min_EndWash_BCS.meth

RF Lens (%): 40 AGC Target: Custom

Normalized AGC Target (%): 250

Maximum Injection Time Mode: Custom Maximum Injection Time (ms): 60

Microscans: 1
Data Type: Centroid
Polarity: Positive

Source Fragmentation: Disabled

Loop Control: All Scan Description:

Time Mode: Unscheduled

Mass List Table

Mass List Table				
Compound	Formula	Adduct	m/z	z
		(no adduct)	700	1

Experiment#2 [tMS2]

Start Time (min): 0 End Time (min): 110

Master Scan:

tMS2

Multiplex lons: False Isolation Window (m/z): 16 Isolation Offset: Off

Collision Energy Type: **Normalized** HCD Collision Energy (%): **27** Orbitrap Resolution: **30000**

TurboTMT: Off

Scan Range Mode: Auto

RF Lens (%): 40 AGC Target: Custom

Normalized AGC Target (%): 1000

Maximum Injection Time Mode: Custom

Maximum Injection Time (ms): 60

Microscans: 1
Data Type: Centroid
Polarity: Positive

Source Fragmentation: Disabled

Loop Control: N

N (Number of Spectra): 38

Dynamic RT: Off

Time Mode: Unscheduled

Scan Description:

Mass List Table

Mass List Table				
Compound	Formula	Adduct	m/z	z
		(no adduct)	408.4356	3
		(no adduct)	424.4428	3
		(no adduct)	440.4501	3
		(no adduct)	456.4574	3
		(no adduct)	472.4647	3
		(no adduct)	488.4719	3
		(no adduct)	504.4792	3
		(no adduct)	520.4865	3
		(no adduct)	536.4938	3
		(no adduct)	552.501	3
		(no adduct)	568.5083	3
		(no adduct)	584.5156	3
		(no adduct)	600.5228	3
		(no adduct)	616.5302	3
		(no adduct)	632.5374	3
		(no adduct)	648.5447	3
		(no adduct)	664.552	3
		(no adduct)	680.5592	3
		(no adduct)	696.5665	3
		(no adduct)	712.5737	3
		(no adduct)	728.5811	3
		(no adduct)	744.5884	3

(no adduct)	760.5956	3
(no adduct)	776.6029	3
(no adduct)	792.6101	3
(no adduct)	808.6174	3
(no adduct)	824.6248	3
(no adduct)	840.632	3
(no adduct)	856.6393	3
(no adduct)	872.6466	3
(no adduct)	888.6538	3
(no adduct)	904.6611	3
(no adduct)	920.6683	3
(no adduct)	936.6757	3
(no adduct)	952.6829	3
(no adduct)	968.6902	3
(no adduct)	984.6975	3
(no adduct)	1000.7047	3
(no adduct)	400.4319	3
(no adduct)	416.4392	3
(no adduct)	432.4465	3
(no adduct)	448.4537	3
(no adduct)	464.461	3
(no adduct)	480.4683	3
(no adduct)	496.4756	3
(no adduct)	512.4828	3
(no adduct)	528.4901	3
(no adduct)	544.4974	3
(no adduct)	560.5046	3
(no adduct)	576.512	3
(no adduct)	592.5192	3
(no adduct)	608.5265	3

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