

Acquisition Method Info

Method Name Daytam AA method.m
 Method Path D:\MassHunter\Methods\Daytam AA method.m
 Method Description 6460 Triple Quad LC/MS ESI with Agilent Jet Stream Technology
 Positive MRM Reserpine Checkout

Device List

HiP Sampler
 Quat. Pump
 Column Comp.
 QQQ

MS QQQ Mass Spectrometer

Ion Source AJS ESI Tune File D:\MassHunter\Tune\QQQ\G6460C\atunes.TUNE.XML
 Stop Mode No Limit/As Pump Stop Time (min) 2.0
 Time Filter On Time Filter Width (min) 0.1

Time Segments

Index	Start Time (min)	Scan Type	Ion Mode	Div Valve	Delta EMV	Store
1	0	MRM	ESI+Agilent Jet Stream	To Waste	0	No
2	1.5	MRM	ESI+Agilent Jet Stream	To MS	0	Yes
3	5.5	MRM	ESI+Agilent Jet Stream	To Waste	0	No

Time Segment 1

Scan Segments

Cpd Name	ISTD?	Prec Ion	MS1 Res	Prod Ion	MS2 Res	Dwell	Frag (V)	CE (V)	Cell Acc (V)	Polarity
Compound 1	No	350	Unit/Enh (6490)	200	Unit/Enh (6490)	200	135	0	7	Positive

Scan Parameters

Data Stg Centroid Threshold 0

Source Parameters

Parameter	Value (+)	Value (-)
Gas Temp (°C)	150	150
Gas Flow (l/min)	11	11
Nebulizer (psi)	40	40
SheathGasHeater	400	400
SheathGasFlow	11	11
Capillary (V)	2000	0
VCharging	0	0

Time Segment 2

Scan Segments

Cpd Name	ISTD?	Prec Ion	MS1 Res	Prod Ion	MS2 Res	Dwell	Frag (V)	CE (V)	Cell Acc (V)	Polarity
DL-Cystine IS	Yes	244.9	Unit/Enh (6490)	153.9	Unit/Enh (6490)	5	90	8	7	Positive
L-Cystine	No	241	Unit/Enh (6490)	74.2	Unit/Enh (6490)	5	100	24	7	Positive
L-Tryptophan IS	Yes	210.1	Unit/Enh (6490)	192.1	Unit/Enh (6490)	5	90	4	7	Positive
L-Tryptophan	No	205.1	Unit/Enh (6490)	188.1	Unit/Enh (6490)	5	80	1	7	Positive
L-Tyrosine IS	Yes	192.1	Unit/Enh (6490)	145.2	Unit/Enh (6490)	5	80	8	7	Positive
L-Tyrosine	No	182.1	Unit/Enh (6490)	165	Unit/Enh (6490)	5	100	1	7	Positive
L-Citrulline IS	Yes	177.2	Unit/Enh (6490)	160.2	Unit/Enh (6490)	5	90	4	7	Positive
L-Arginine IS	Yes	177.2	Unit/Enh (6490)	70.2	Unit/Enh (6490)	5	110	20	7	Positive
L-Citrulline	No	176.4	Unit/Enh (6490)	159.3	Unit/Enh (6490)	5	80	3	7	Positive

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Cpd Name	ISTD?	Prec Ion	MS1 Res	Prod Ion	MS2 Res	Dwell	Frag (V)	CE (V)	Cell Acc (V)	Polarity
L-Arginine	No	175.2	Unit/Enh (6490)	70.2	Unit/Enh (6490)	5	110	20	7	Positive
L-Phenylalanine IS	Yes	175.1	Unit/Enh (6490)	129.1	Unit/Enh (6490)	5	100	8	7	Positive
3-Methyl-L-Histidine IS	Yes	173.2	Unit/Enh (6490)	127.2	Unit/Enh (6490)	5	80	10	7	Positive
3-Methyl-L-Histidine L-	No	170.1	Unit/Enh (6490)	124.1	Unit/Enh (6490)	5	100	10	7	Positive
L-Phenylalanine DL-5-Hydroxy lysine	No	166.1	Unit/Enh (6490)	120.1	Unit/Enh (6490)	5	80	6	7	Positive
L-Histidine	No	163.1	Unit/Enh (6490)	128.1	Unit/Enh (6490)	5	90	6	7	Positive
L-Methionine IS	Yes	153.1	Unit/Enh (6490)	107.2	Unit/Enh (6490)	5	80	6	7	Positive
L-Lysine IS	Yes	151.1	Unit/Enh (6490)	88.1	Unit/Enh (6490)	5	90	16	7	Positive
L-Methionine L-Glutamic acid IS	No	150.1	Unit/Enh (6490)	104.1	Unit/Enh (6490)	5	80	4	7	Positive
L-Glutamine IS	Yes	150.1	Unit/Enh (6490)	85.2	Unit/Enh (6490)	5	80	12	7	Positive
L-Glutamic acid	Yes	148.1	Unit/Enh (6490)	85.1	Unit/Enh (6490)	5	80	14	7	Positive
L-Lysine	No	148.1	Unit/Enh (6490)	84.2	Unit/Enh (6490)	5	80	12	7	Positive
L-Glutamine L-Leucine IS	No	147.1	Unit/Enh (6490)	84.2	Unit/Enh (6490)	5	80	12	7	Positive
L-Ornithine IS	No	147.1	Unit/Enh (6490)	84.2	Unit/Enh (6490)	5	80	12	7	Positive
L-Aspartic acid IS	Yes	142.2	Unit/Enh (6490)	96.3	Unit/Enh (6490)	5	90	6	7	Positive
L-Asparagine IS	Yes	138.2	Unit/Enh (6490)	74.2	Unit/Enh (6490)	5	80	16	7	Positive
L-Aspartic acid L-ornithine	Yes	137.1	Unit/Enh (6490)	91.2	Unit/Enh (6490)	5	90	5	7	Positive
L-Asparagine	Yes	135	Unit/Enh (6490)	89.1	Unit/Enh (6490)	5	80	4	7	Positive
L-Aspartic acid	No	134.1	Unit/Enh (6490)	74.1	Unit/Enh (6490)	5	90	10	7	Positive
L-ornithine	No	133.2	Unit/Enh (6490)	70.3	Unit/Enh (6490)	5	80	14	7	Positive
L-asparagine	No	133.1	Unit/Enh (6490)	74.2	Unit/Enh (6490)	5	70	10	7	Positive
L-isoleucine	No	132.2	Unit/Enh (6490)	69.2	Unit/Enh (6490)	5	100	14	7	Positive
Trans-4-hydroxy L-proline	No	132.2	Unit/Enh (6490)	68.2	Unit/Enh (6490)	5	90	20	7	Positive
L-Leucine	No	132.2	Unit/Enh (6490)	43.3	Unit/Enh (6490)	5	100	24	7	Positive
DL-Valine IS	Yes	126.1	Unit/Enh (6490)	80.2	Unit/Enh (6490)	5	80	8	7	Positive
Taurine	No	126.1	Unit/Enh (6490)	44.3	Unit/Enh (6490)	5	110	14	7	Positive
L-Proline IS	Yes	122.1	Unit/Enh (6490)	75.2	Unit/Enh (6490)	5	90	14	7	Positive
L-Threonine IS	Yes	121.1	Unit/Enh (6490)	75	Unit/Enh (6490)	5	80	6	7	Positive
L-Threonine	No	120.2	Unit/Enh (6490)	74.2	Unit/Enh (6490)	5	80	4	7	Positive
L-Valine	No	118.2	Unit/Enh (6490)	72.2	Unit/Enh (6490)	5	80	4	7	Positive
L-Norvaline	No	118.1	Unit/Enh (6490)	72.1	Unit/Enh (6490)	5	90	5	7	Positive
L-Proline	No	116.2	Unit/Enh (6490)	70.2	Unit/Enh (6490)	5	90	12	7	Positive
L-Serine IS	Yes	109.1	Unit/Enh (6490)	63	Unit/Enh (6490)	5	90	8	7	Positive
L-Serine	No	106.2	Unit/Enh (6490)	60.2	Unit/Enh (6490)	5	80	4	7	Positive
L-2-aminobutyric acid	No	104.2	Unit/Enh (6490)	58.3	Unit/Enh (6490)	5	80	4	7	Positive
3-Amino isobutyric acid	No	104.1	Unit/Enh (6490)	86.2	Unit/Enh (6490)	5	100	2	7	Positive

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Cpd Name	ISTD?	Prec Ion	MS1 Res	Prod Ion	MS2 Res	Dwell	Frag (V)	CE (V)	Cell Acc (V)	Polarity
Gamma-aminobutyric acid	No	104	Unit/Enh (6490)	87.1	Unit/Enh (6490)	5	80	6	7	Positive
L-Alanine IS	Yes	94.1	Unit/Enh (6490)	48.2	Unit/Enh (6490)	5	90	6	7	Positive
L-Alanine	No	90.2	Unit/Enh (6490)	44.2	Unit/Enh (6490)	5	80	4	7	Positive
Beta-Alanine	No	90.1	Unit/Enh (6490)	72.1	Unit/Enh (6490)	5	80	2	7	Positive
Sarcosine	No	90.1	Unit/Enh (6490)	44.2	Unit/Enh (6490)	5	90	8	7	Positive
L-Glycine IS	Yes	78.2	Unit/Enh (6490)	31.3	Unit/Enh (6490)	5	90	4	7	Positive
L-Glycin	No	76.2	Unit/Enh (6490)	30.1	Unit/Enh (6490)	5	80	1	7	Positive
Ethanolamine	No	62.1	Unit/Enh (6490)	44.2	Unit/Enh (6490)	5	80	4	7	Positive

Scan Parameters

Data Stg Threshold
Centroid 0

Source Parameters

Parameter	Value (+)	Value (-)
Gas Temp (°C)	150	150
Gas Flow (l/min)	11	11
Nebulizer (psi)	40	40
SheathGasHeater	400	400
SheathGasFlow	11	11
Capillary (V)	2000	0
VCharging	0	0

Time Segment 3

Scan Segments

Cpd Name	ISTD?	Prec Ion	MS1 Res	Prod Ion	MS2 Res	Dwell	Frag (V)	CE (V)	Cell Acc (V)	Polarity
Compound 1	No	350	Unit/Enh (6490)	200	Unit/Enh (6490)	200	135	0	7	Positive

Scan Parameters

Data Stg Threshold
Centroid 0

Source Parameters

Parameter	Value (+)	Value (-)
Gas Temp (°C)	150	150
Gas Flow (l/min)	11	11
Nebulizer (psi)	40	40
SheathGasHeater	400	400
SheathGasFlow	11	11
Capillary (V)	2000	0
VCharging	0	0

Chromatograms

Chrom Type Label Offset Y-Range
TIC TIC 0 10000000

Instrument Curves

Actual

Name: **HIP Sampler** Model: **G4226A**

Auxiliary

Draw Speed	200.0 µL/min
Eject Speed	200.0 µL/min
Draw Position Offset	0.0 mm
Wait Time After Drawing	2.0 s
Sample Flush Out Factor	5.0
Vial/Well bottom sensing	Yes

Injection

Injection Mode	Injection with needle wash
Injection Volume	3.00 µL
Needle Wash	
Needle Wash Location	Flush Port
Wash Time	2.0 s

High throughput

Automatic Delay Volume Reduction	No
Overlapped Injection	
Enable Overlapped Injection	No

Valve Switching

Valve Movements	0
Valve Switch Time 1	
Switch Time 1 Enabled	No
Valve Switch Time 2	
Switch Time 2 Enabled	No
Valve Switch Time 3	
Switch Time 3 Enabled	No
Valve Switch Time 4	
Switch Time 4 Enabled	No

Stop Time

Stoptime Mode	As pump/No limit
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Post Time

Posttime Mode	Off
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Name: **Quat. Pump** Model: **G4204A**

Flow 0.700 mL/min
 Use Solvent Types No
 Low Pressure Limit 0.00 bar
 High Pressure Limit 600.00 bar
 Max. Flow Ramp Up 100.000 mL/min²
 Primary Channel Automatic
 Max. Flow Ramp Down 100.000 mL/min²

Stroke

Automatic Stroke Calculation Yes

Compress

Compressibility Mode No Compensation

Mixer Selection

Selected Mixer Use Mixer if installed

Blend Assist

Enabled No

Stop Time

Stoptime Mode Time set

Stoptime 10.00 min

Post Time

Posttime Mode Off

Solvent Composition

	Channel	Ch. 1 Solv.	Name 1	Used	Percent
1	A	100.0 % Water V.03		Yes	22.00 %
2	B	100.0 % Acetonitrile V.03		Yes	78.00 %
3	C			No	
4	D			No	

Timetable

	Time	A	B	C	D	Flow	Pressure
1	1.00 min	22.00 %	78.00 %	0.00 %	0.00 %	0.700 mL/min	--- bar
2	4.00 min	70.00 %	30.00 %	0.00 %	0.00 %	0.700 mL/min	--- bar
3	5.00 min	70.00 %	30.00 %	0.00 %	0.00 %	0.700 mL/min	--- bar
4	5.10 min	22.00 %	78.00 %	0.00 %	0.00 %	0.700 mL/min	--- bar
5	9.00 min	22.00 %	78.00 %	0.00 %	0.00 %	0.700 mL/min	--- bar

Name: **Column Comp.** Model: **G1316A**

Left Temperature Control

Temperature Control Mode Temperature Set

Temperature 30.0 °C

Enable Analysis Left Temperature

Enable Analysis Left Temperature On Yes

Enable Analysis Left Temperature Value 0.8 °C

Right Temperature Control

Right temperature Control Mode Combined

Enable Analysis Right Temperature

Enable Analysis Right Temperature On Yes

Enable Analysis Right Temperature Value 0.8 °C

Stop Time

Stoptime Mode As pump/injector

Post Time

Posttime Mode Off