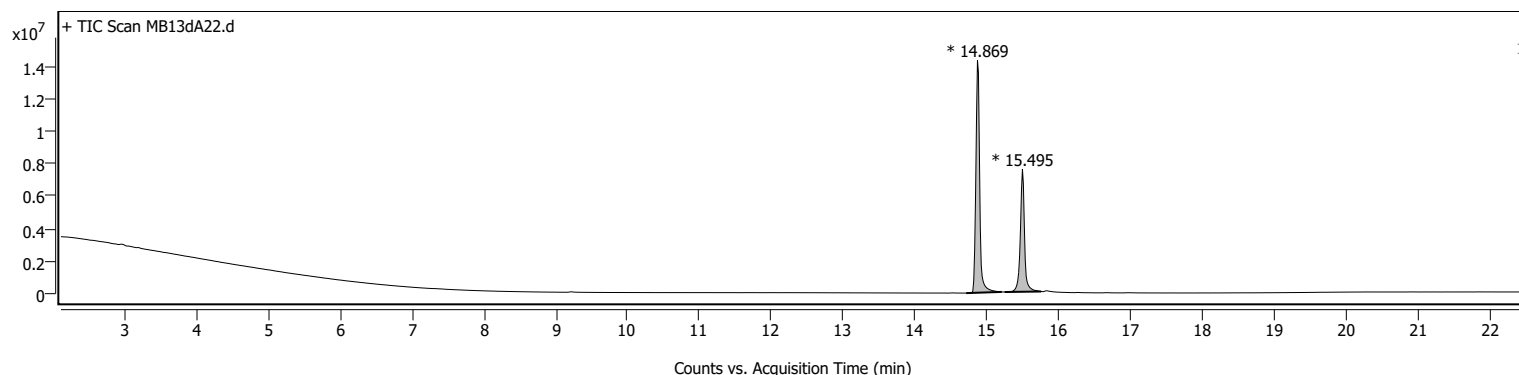
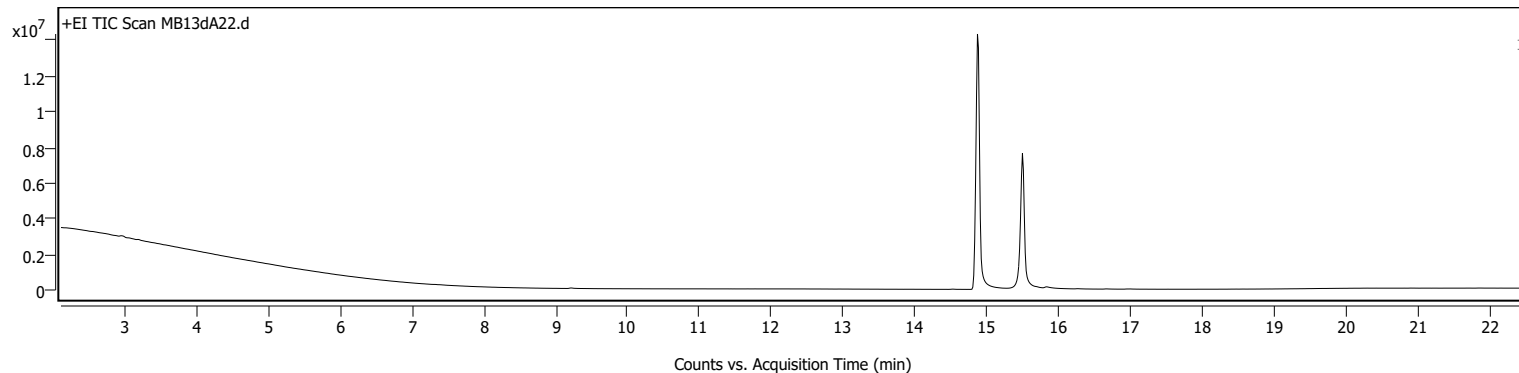


## Sample Information

<b>Name</b>	MB13dA22	<b>Data File Path</b>	D:\MassHunter\GCMS\1\data\MB\MB13dA22.D
<b>Sample ID</b>		<b>Acq. Time (Local)</b>	6/10/2022 6:49:44 PM (UTC+02:00)
<b>Instrument</b>	GCMS	<b>Method Path (Acq)</b>	D:\MassHunter\GCMS\1\methods\Standard HP 5 MS Temp 40 -320C_solvent front 2 m.M
<b>MS Type</b>	Q	<b>Version (Acq SW)</b>	MassHunter GC/MS Acquisition 10.0.384.1 14-Feb-2019 Copyright © 1989-2018 Agilent Technologies, Inc.
<b>Inj. Vol. (ul)</b>	0.5	<b>IRM Status</b>	
<b>Position</b>	121	<b>Method Path (DA)</b>	D:\MassHunter\GCMS\1\data\MB\MB13dA22.D\Results\Qual\Version4\default.m
<b>Plate Pos.</b>		<b>Target Source Path</b>	
<b>Operator</b>		<b>Result Summary</b>	

## Sample Chromatograms



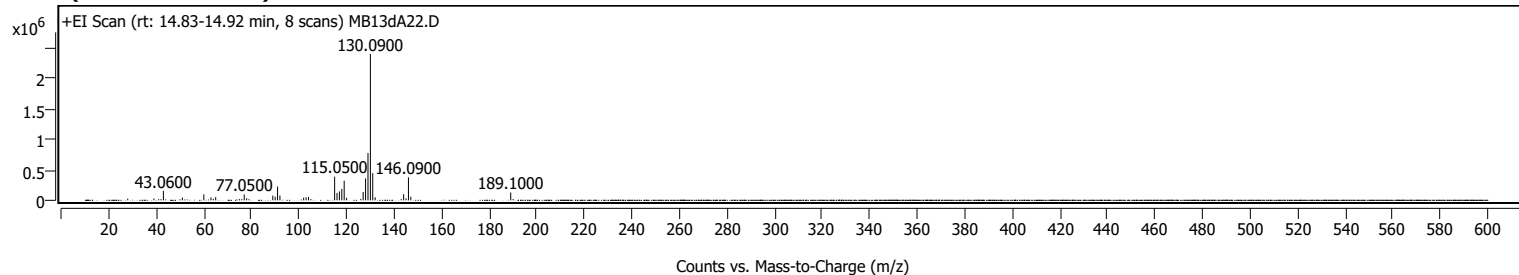
### Chromatogram Peaks

Peak	Start	RT	End	Height	Area	Area %	SNR
1	14.713	14.869	15.208	14310389	52735784	100.00	
2	15.247	15.495	15.756	7553811	30579322	57.99	

## Sample Spectra

### + Scan (rt: 14.83-14.92 min)

### Peak 1 from + TIC Scan



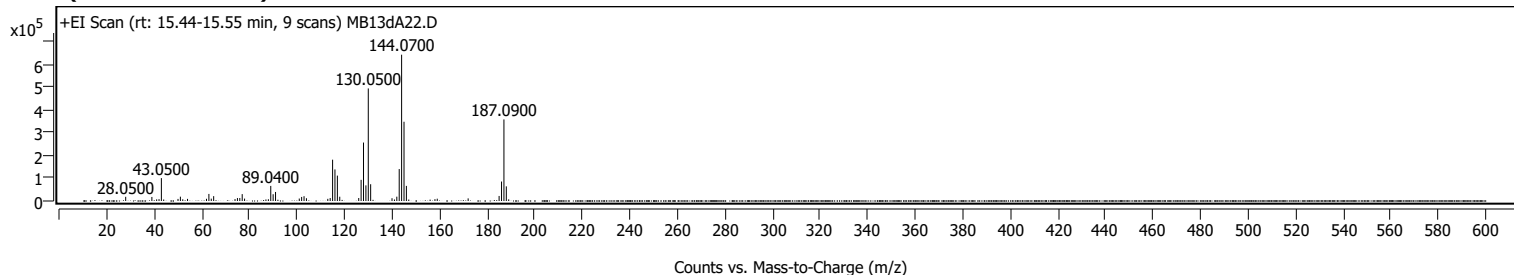
# Analysis Report

## Spectrum Peaks

m/z	Z	Abund	Abund %	m/z (Calc)	Diff (ppm)	Ion Species	Formula	Ion Type
28.0600		26728	1.11					
39.0600		29926	1.24					
43.0600		156588	6.50					
51.0500		45954	1.91					
60.0500		98031	4.07					
63.0500		47419	1.97					
64.0400		24483	1.02					
65.0400		53627	2.23					
77.0500		97303	4.04					
78.0600		33701	1.40					
89.0500		77752	3.23					
90.0500		60552	2.52					
91.0600		226879	9.42					
92.0600		79325	3.29					
102.0400		44426	1.85					
103.0500		48538	2.02					
104.0500		56783	2.36					
115.0500		389675	16.19					
116.0600		121087	5.03					
117.0600		146941	6.10					
118.0700		187651	7.79					
119.0700	1	322865	13.41					
120.0800	1	42316	1.76					
127.0500		138674	5.76					
128.0600		360006	14.95					
129.0800		778540	32.34					
130.0900		2407487	100.00					
131.0800	1	447159	18.57					
132.0800	1	50643	2.10					
144.0700		102283	4.25					
145.0800		31919	1.33					
146.0900	1	378341	15.72					
147.0900	1	59024	2.45					
189.1000		128595	5.34					

## + Scan (rt: 15.44-15.55 min)

## Peak 2 from + TIC Scan



# Analysis Report

## Spectrum Peaks

m/z	Z	Abund	Abund %	m/z (Calc)	Diff (ppm)	Ion Species	Formula	Ion Type
28.0500		18458	2.86					
39.0600		17884	2.77					
41.0600		7163	1.11					
42.0600		8200	1.27					
43.0500		100855	15.63					
44.0200		6563	1.02					
50.0400		9216	1.43					
51.0400		19012	2.95					
52.0400		7557	1.17					
54.0500		9293	1.44					
62.0200		9746	1.51					
63.0300		31133	4.83					
64.0300		10340	1.60					
65.0400		21442	3.32					
74.0200		8486	1.32					
75.0300		13058	2.02					
76.0300		13177	2.04					
77.0400		30384	4.71					
78.0300		10584	1.64					
87.0200		6838	1.06					
88.0300		7739	1.20					
89.0400		66594	10.32					
90.0400		29831	4.62					
91.0500		39871	6.18					
101.0300		10798	1.67					
102.0300		17916	2.78					
103.0400		21183	3.28					
104.0500		12245	1.90					
113.0300		9496	1.47					
114.0300		12963	2.01					
115.0500		182160	28.23					
116.0400		139265	21.58					
117.0500		111942	17.35					
118.0500		18799	2.91					
126.0400		13115	2.03					
127.0400		93285	14.46					
128.0500		257817	39.96					
129.0500		68877	10.68					
130.0500	1	496400	76.94					
131.0500	1	73995	11.47					
140.0300		12259	1.90					
141.0500		6746	1.05					
142.0500		19071	2.96					
143.0600		140856	21.83					
144.0700		645210	100.00					
145.0800		349478	54.16					
146.0700		66707	10.34					
158.0800		8152	1.26					
159.0500		10161	1.57					
172.0500		11572	1.79					
185.0600		22190	3.44					
186.0800		85812	13.30					
187.0900	1	359762	55.76					
188.0900	1	64550	10.00					
189.1000	1	7334	1.14					

MassHunter Qual 10.0  
(End of Report)