

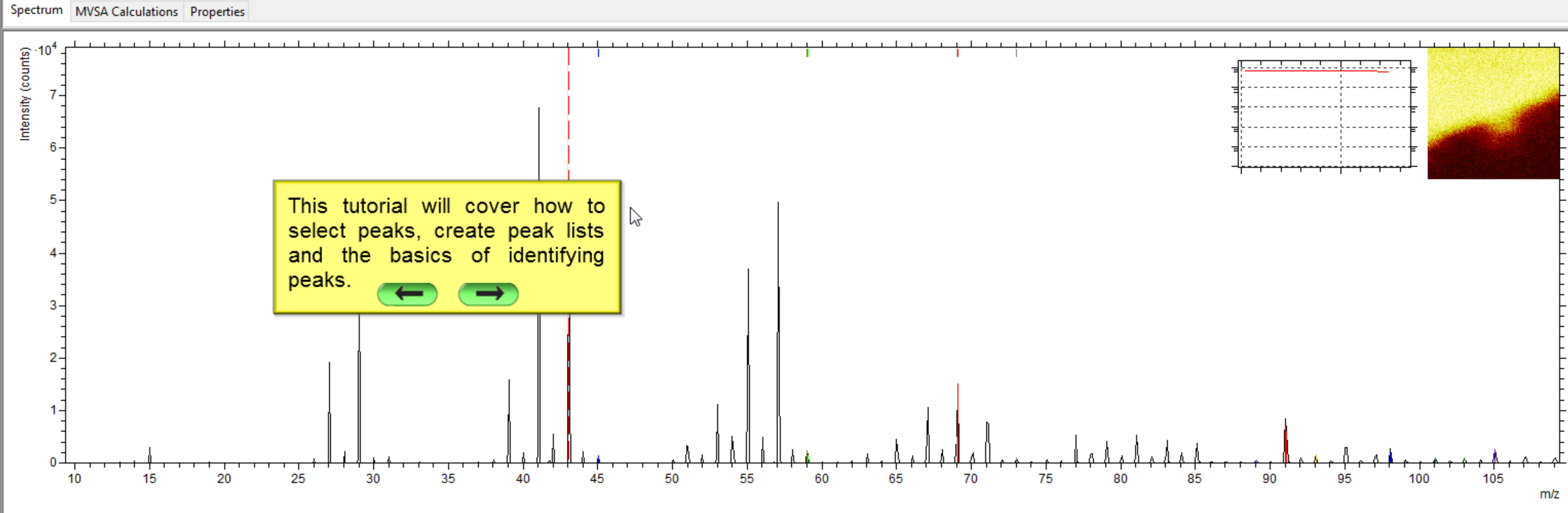
C_3H_7+	m/z	43.0572	Area (cts)	1,153,550	Explained (%)	100.0	Resolution	5,856
Check...	Dev. (ppm)	68.6	Counts / Shot	0.9779	Peak Difference...		Width (ns)	1.90

Compilations

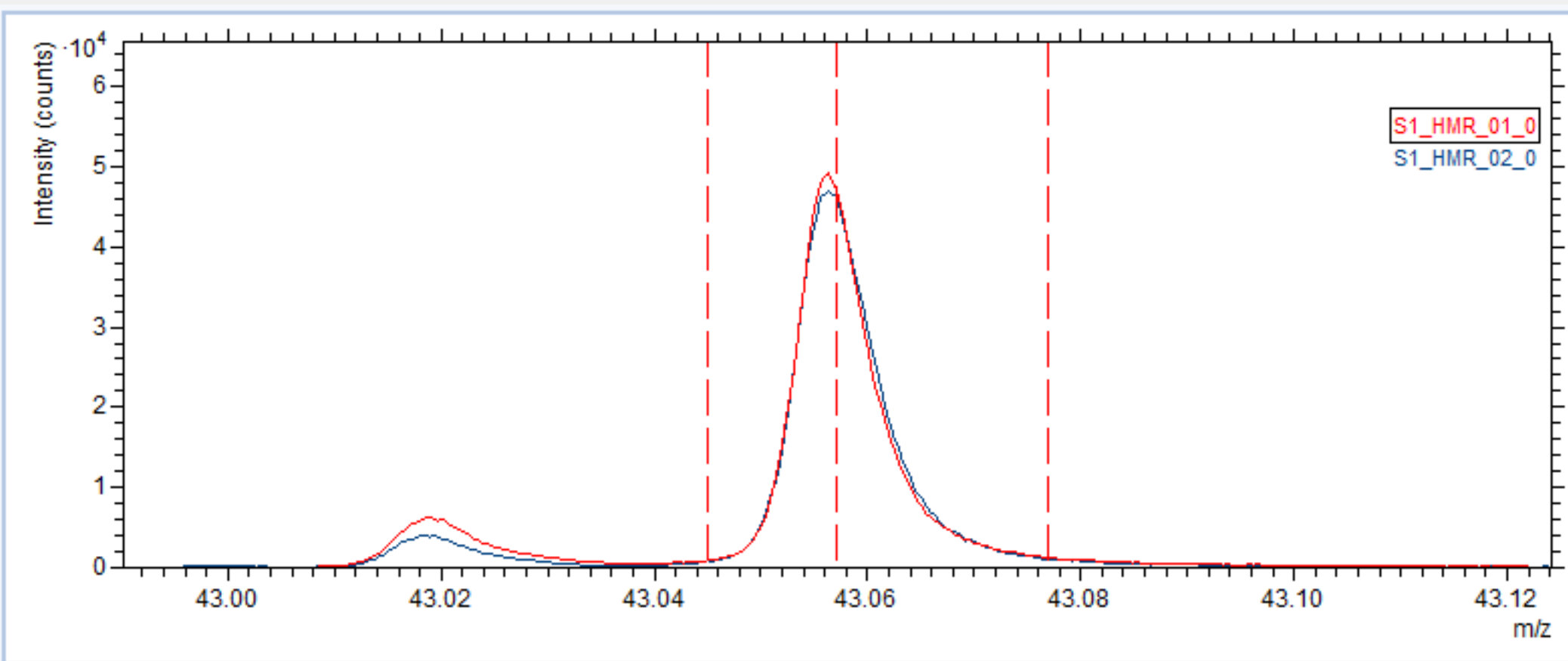
Spectra

S1\_HMR\_01\_0  
<No Sample Name> (S1\_HMR\_01)

S1\_HMR\_02\_0  
<No Sample Name> (S1\_HMR\_02)



V	No.	m / z	Area / cts	Color	Peak Label	Description
<b>Mass Intervals</b>						
<input checked="" type="checkbox"/>	2	45.0352	32713	Blue	45.03 u	PEG
<input checked="" type="checkbox"/>	3	59.0136	51836	Olive	59.01 u	PMMA
<input checked="" type="checkbox"/>	4	59.0511	37293	Green	59.05 u	PEG
<input checked="" type="checkbox"/>	5	69.0376	118182	Grey	69.04 u	PMMA
<input checked="" type="checkbox"/>	6	69.0762	360824	Red	69.08 u	
<input checked="" type="checkbox"/>	7	73.0342	16483	Dark Grey	73.03 u	PEG
<input checked="" type="checkbox"/>	8	87.0478	6134	Orange	87.05 u	PEG
<input checked="" type="checkbox"/>	9	89.0617	3870	Light Blue	89.06 u	PEG
<input checked="" type="checkbox"/>	10	91.0566	238681	Red	91.05 u	PS
<input checked="" type="checkbox"/>	11	93.0730	55170	Yellow	93.07 u	PMMA
<input checked="" type="checkbox"/>	12	98.0763	70210	Blue	98.07 u	GTP
<input checked="" type="checkbox"/>	13	101.0605	26893	Green	101.06 u	PMMA
<input checked="" type="checkbox"/>	14	103.0490	34262	Light Green	103.05 u	PEG
<input checked="" type="checkbox"/>	15	105.0688	63931	Purple	105.07 u	PS



<No Sample Name> (S1\_HMR\_01) \* - Measurement Explorer - Spectra

File Edit Spectrum Mass Interval List Peak List View Help

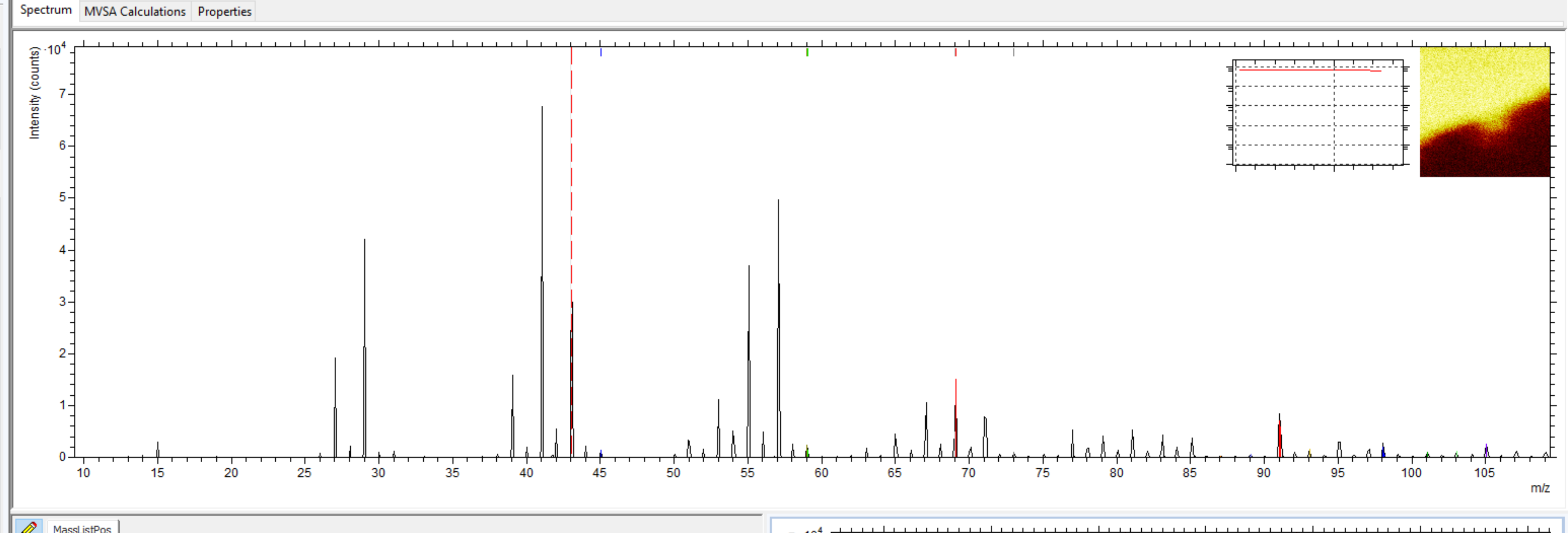
Total Range Total Area All Intervals Find: C3H3

m/z	43.0572	Area (cts)	1,153,550	Explained (%)	100.0	Resolution	5,856
Dev. (ppm)	68.6	Counts / Shot	0.9779	Peak Difference...		Width (ns)	1.90

Compilations

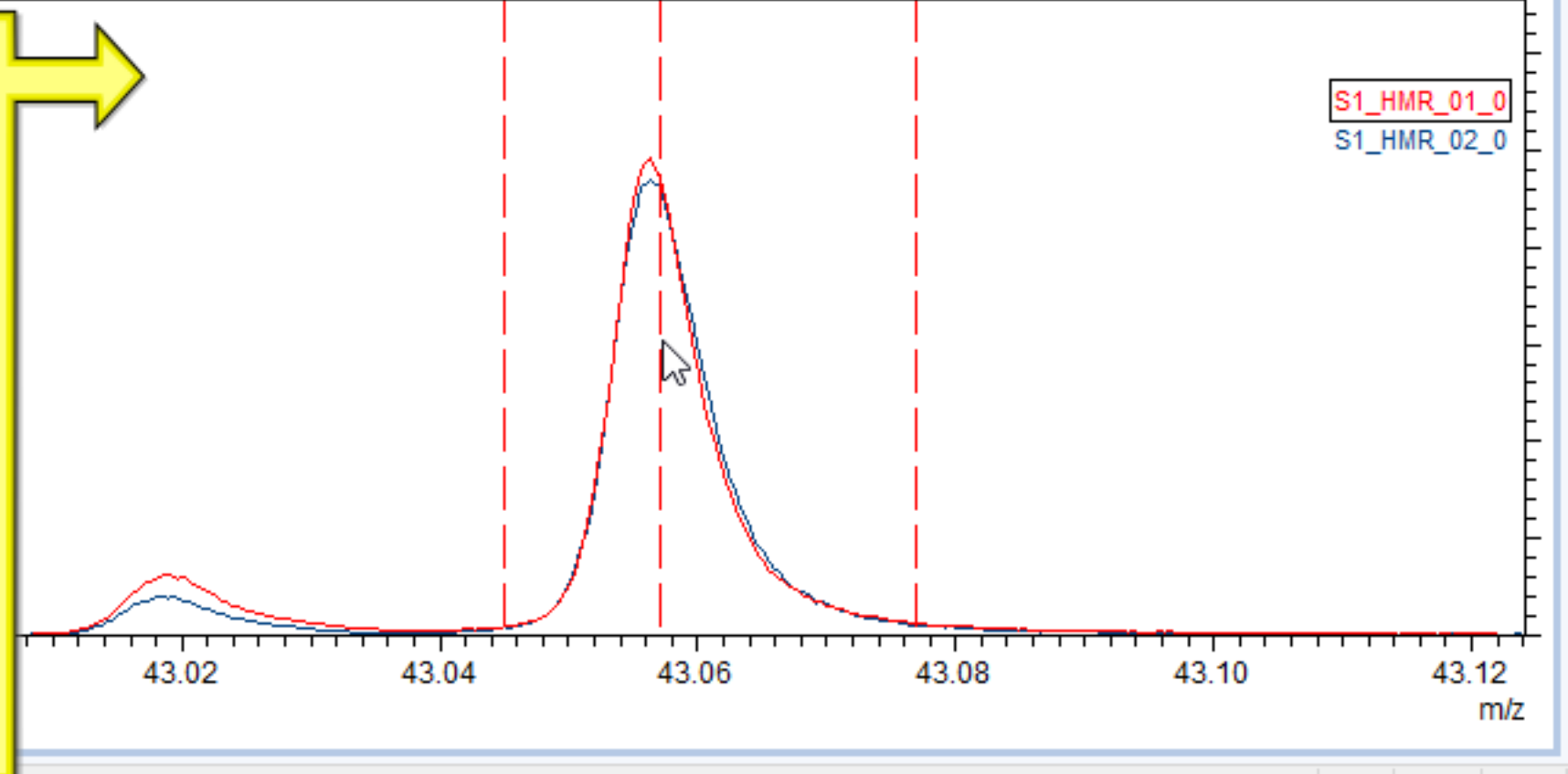
Spectra

- S1\_HMR\_01\_0
- <No Sample Name> (S1\_HMR\_01)
- S1\_HMR\_02\_0
- <No Sample Name> (S1\_HMR\_02)



Left clicking on a peak will automatically create three dotted lines. The two outside lines mark the peak integration limits and the center line defines the centroid of the peak. The position of all of the lines can be modified. You will need to adjust the positions of the lines before trying to identify the peak or looking at peak areas.

NOTE: I disagree how IONTOF defines their peak limit and centroid lines. They place the lines weighted by any assymetry in the selected peak. This will place the centroid to the side of the true peak center and the peak integration limits too far out. The centroid should be placed at the highest point of the peak and the integration limits should be placed closer to the full width at half maximum of the peak.



File Edit Spectrum Mass Interval List Peak List View Help

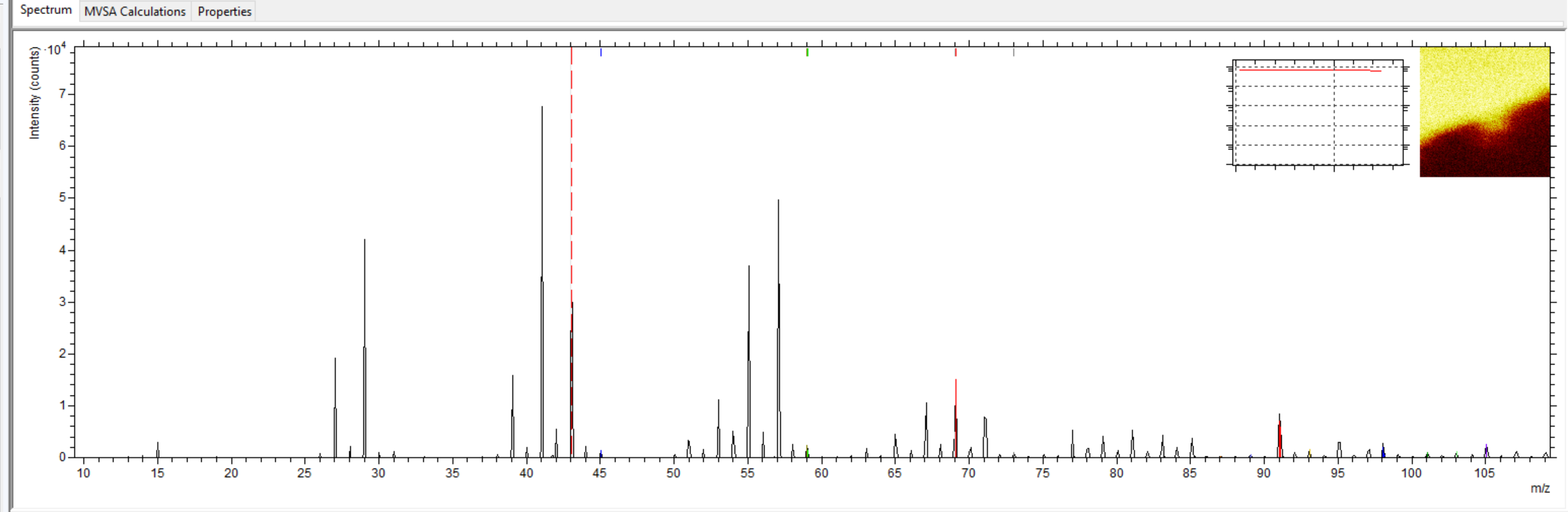
Total Range Total Area All Intervals Find: C3H3

C\_3H\_7+ m/z 43.0572 Area (cts) 1,153,550 Explained (%) 100.0 Resolution 5,856  
 Dev. (ppm) 68.6 Counts / Shot 0.9779 Peak Difference... Width (ns) 1.90

Compilations

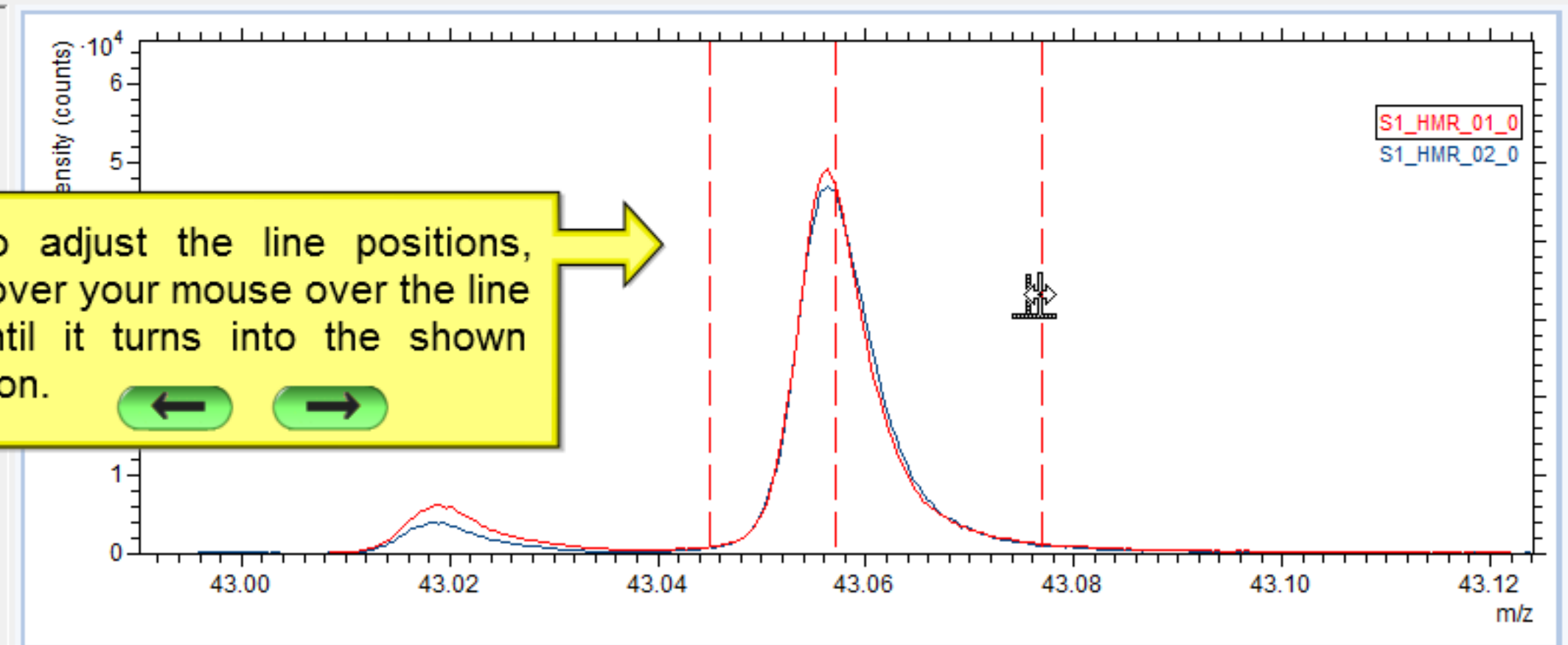
Spectra

S1\_HMR\_01\_0  
 <No Sample Name> (S1\_HMR\_01)  
 S1\_HMR\_02\_0  
 <No Sample Name> (S1\_HMR\_02)



MassListPos

V	No.	m / z	Area / cts	Color	Peak Label	Description
<b>Mass Intervals</b>						
<input checked="" type="checkbox"/>	2	45.0352	32713	Blue	45.03 u	PEG
<input checked="" type="checkbox"/>	3	59.0136	51836	Olive	59.01 u	PMMA
<input checked="" type="checkbox"/>	4	59.0511	37293	Green	59.05 u	PEG
<input checked="" type="checkbox"/>	5	69.0376	118182	Grey	69.04 u	PMMA
<input checked="" type="checkbox"/>	6	69.0762	360824	Red	69.08 u	
<input checked="" type="checkbox"/>	7	73.0342	16483	Dark Grey	73.03 u	PEG
<input checked="" type="checkbox"/>	8	87.0478	6134	Orange	87.05 u	PEG
<input checked="" type="checkbox"/>	9	89.0617	3870	Light Blue	89.06 u	PEG
<input checked="" type="checkbox"/>	10	91.0566	238681	Red	91.05 u	PS
<input checked="" type="checkbox"/>	11	93.0730	55170	Yellow	93.07 u	PMMA
<input checked="" type="checkbox"/>	12	98.0763	70210	Blue	98.07 u	GTP
<input checked="" type="checkbox"/>	13	101.0605	26893	Green	101.06 u	PMMA
<input checked="" type="checkbox"/>	14	103.0490	34262	Light Green	103.05 u	PEG
<input checked="" type="checkbox"/>	15	105.0688	63931	Purple	105.07 u	PS



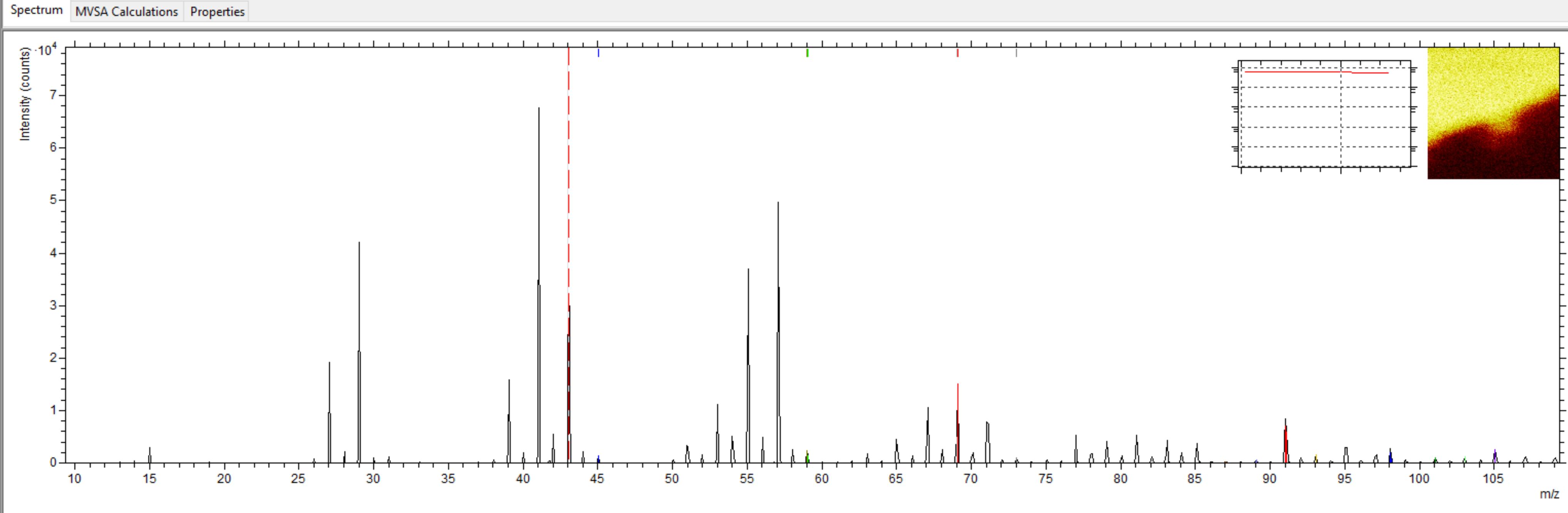
C_3H_7+	m/z	43.0566	Area (cts)	1,015,984	Explained (%)	100.0	Resolution	6,408
Check...	Dev. (ppm)	55.7	Counts / Shot	0.8613	Peak Difference...		Width (ns)	1.74

Compilations

Spectra

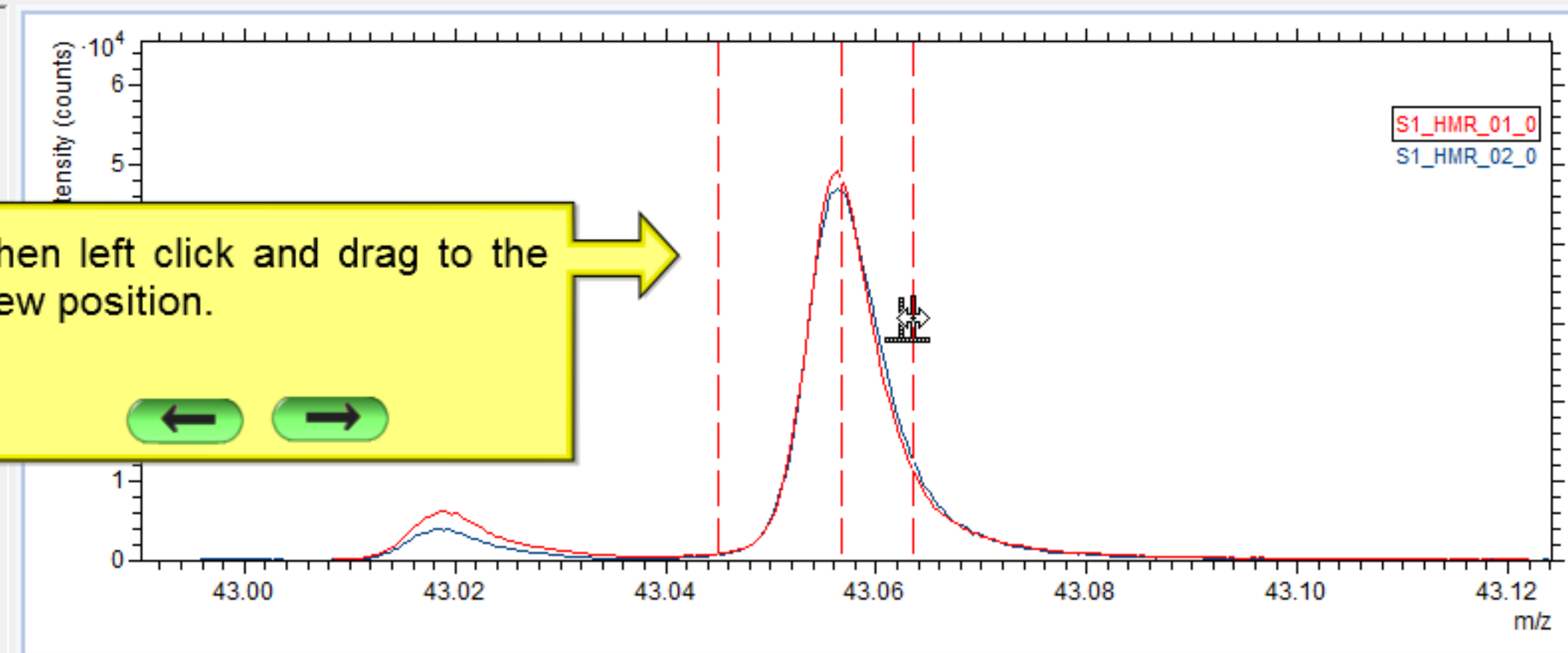
S1\_HMR\_01\_0  
<No Sample Name> (S1\_HMR\_01)

S1\_HMR\_02\_0  
<No Sample Name> (S1\_HMR\_02)



MassListPos

V	No.	m / z	Area / cts	Color	Peak Label	Description
<b>Mass Intervals</b>						
<input checked="" type="checkbox"/>	2	45.0352	32713	Blue	45.03 u	PEG
<input checked="" type="checkbox"/>	3	59.0136	51836	Olive	59.01 u	PMMA
<input checked="" type="checkbox"/>	4	59.0511	37293	Green	59.05 u	PEG
<input checked="" type="checkbox"/>	5	69.0376	118182	Grey	69.04 u	PMMA
<input checked="" type="checkbox"/>	6	69.0762	360824	Red	69.08 u	
<input checked="" type="checkbox"/>	7	73.0342	16483	Dark Grey	73.03 u	PEG
<input checked="" type="checkbox"/>	8	87.0478	6134	Orange	87.05 u	PEG
<input checked="" type="checkbox"/>	9	89.0617	3870	Light Blue	89.06 u	PEG
<input checked="" type="checkbox"/>	10	91.0566	238681	Red	91.05 u	PS
<input checked="" type="checkbox"/>	11	93.0730	55170	Yellow	93.07 u	PMMA
<input checked="" type="checkbox"/>	12	98.0763	70210	Blue	98.07 u	GTP
<input checked="" type="checkbox"/>	13	101.0605	26893	Green	101.06 u	PMMA
<input checked="" type="checkbox"/>	14	103.0490	34262	Light Green	103.05 u	PEG
<input checked="" type="checkbox"/>	15	105.0688	63931	Purple	105.07 u	PS



<No Sample Name> (S1\_HMR\_01) \* - Measurement Explorer - Spectra

File Edit Spectrum Mass Interval List Peak List View Help

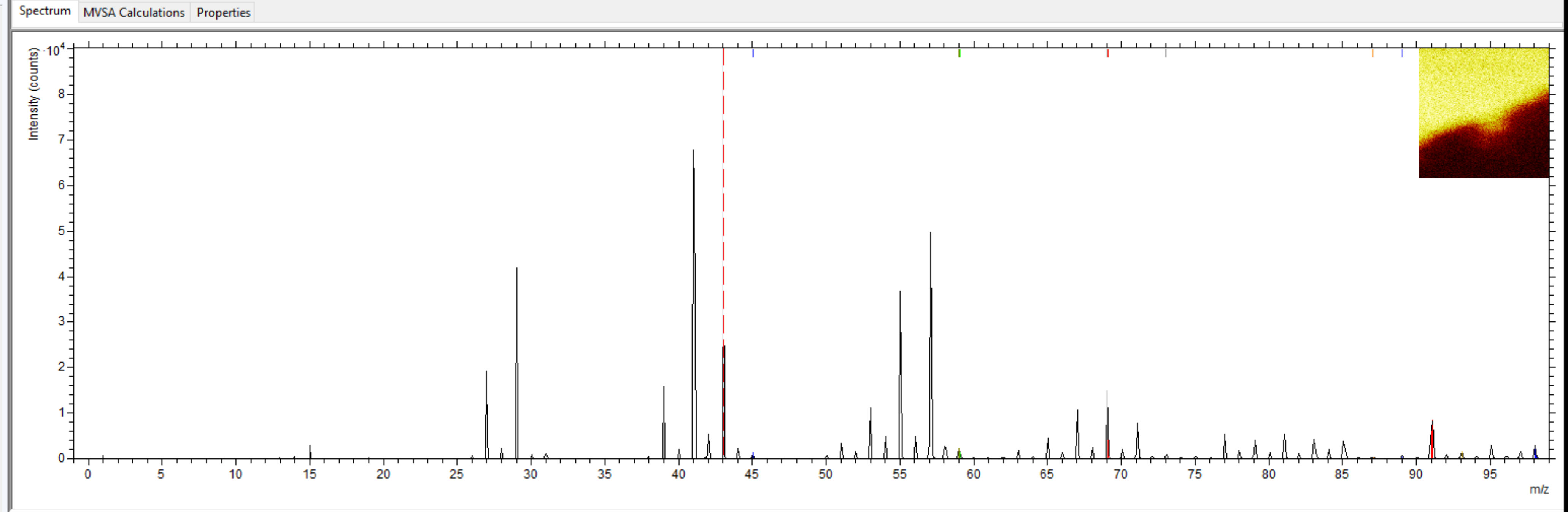
Total Range Total Area All Intervals Find: C3H3

C\_3H\_7+ m/z 43.0564 Area (cts) 960,738 Explained (%) 100.0 Resolution 6,770  
 Dev. (ppm) 50.6 Counts / Shot 0.8144 Peak Difference... Width (ns) 1.65

Compilations

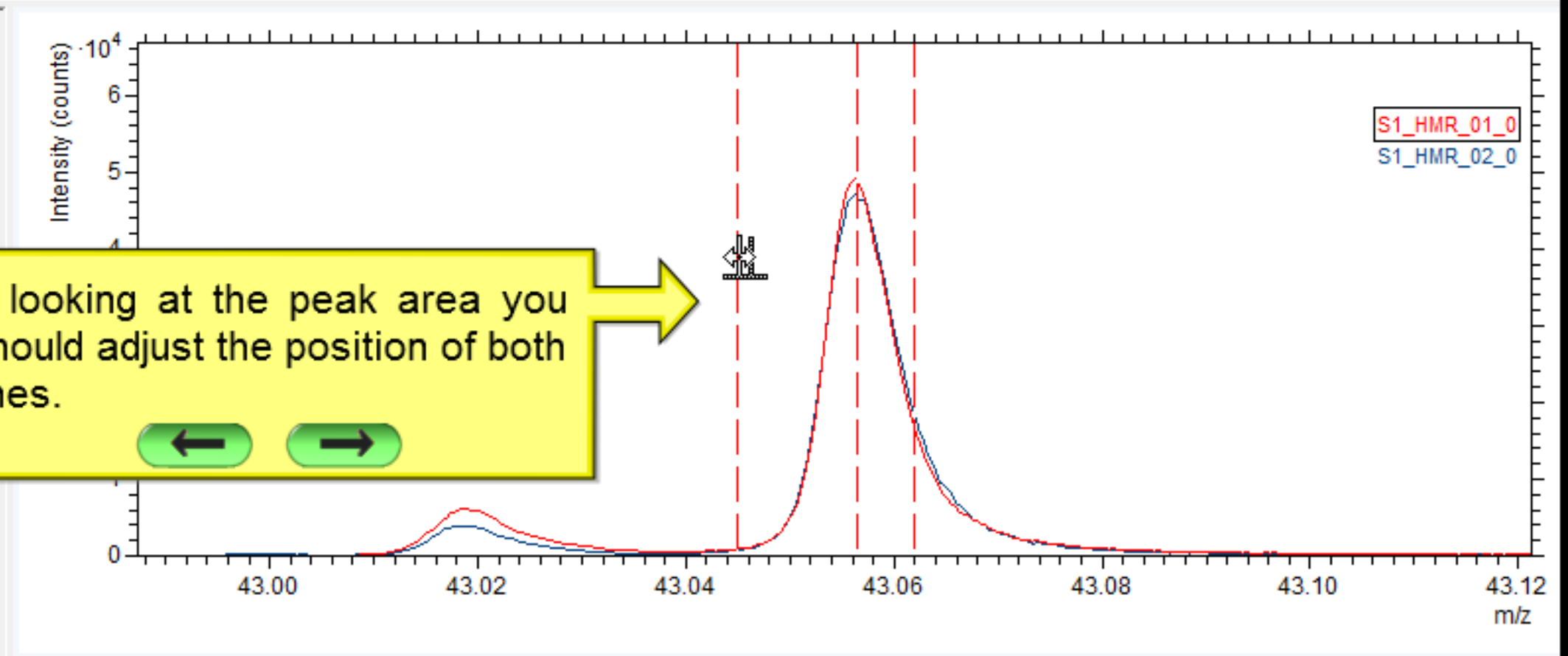
Spectra

S1\_HMR\_01\_0 \*  
 <No Sample Name> (S1\_HMR\_01) \*  
 S1\_HMR\_02\_0  
 <No Sample Name> (S1\_HMR\_02)



MassListPos

V	No.	m/z	Area / cts	Color	Peak Label	Description
<b>Mass Intervals</b>						
<input checked="" type="checkbox"/>	2	45.0352	32713	Blue	45.03 u	PEG
<input checked="" type="checkbox"/>	3	59.0136	51836	Olive	59.01 u	PMMA
<input checked="" type="checkbox"/>	4	59.0511	37293	Green	59.05 u	PEG
<input checked="" type="checkbox"/>	5	69.0376	118182	Grey	69.04 u	PMMA
<input checked="" type="checkbox"/>	6	69.0762	360824	Red	69.08 u	
<input checked="" type="checkbox"/>	7	73.0342	16483	Dark Grey	73.03 u	PEG
<input checked="" type="checkbox"/>	8	87.0478	6134	Orange	87.05 u	PEG
<input checked="" type="checkbox"/>	9	89.0617	3870	Light Blue	89.06 u	PEG
<input checked="" type="checkbox"/>	10	91.0566	238681	Red	91.05 u	PS
<input checked="" type="checkbox"/>	11	93.0730	55170	Yellow	93.07 u	PMMA
<input checked="" type="checkbox"/>	12	98.0763	70210	Blue	98.07 u	GTP
<input checked="" type="checkbox"/>	13	101.0605	26893	Green	101.06 u	PMMA
<input checked="" type="checkbox"/>	14	103.0490	34262	Light Green	103.05 u	PEG
<input checked="" type="checkbox"/>	15	105.0688	63931	Purple	105.07 u	PS



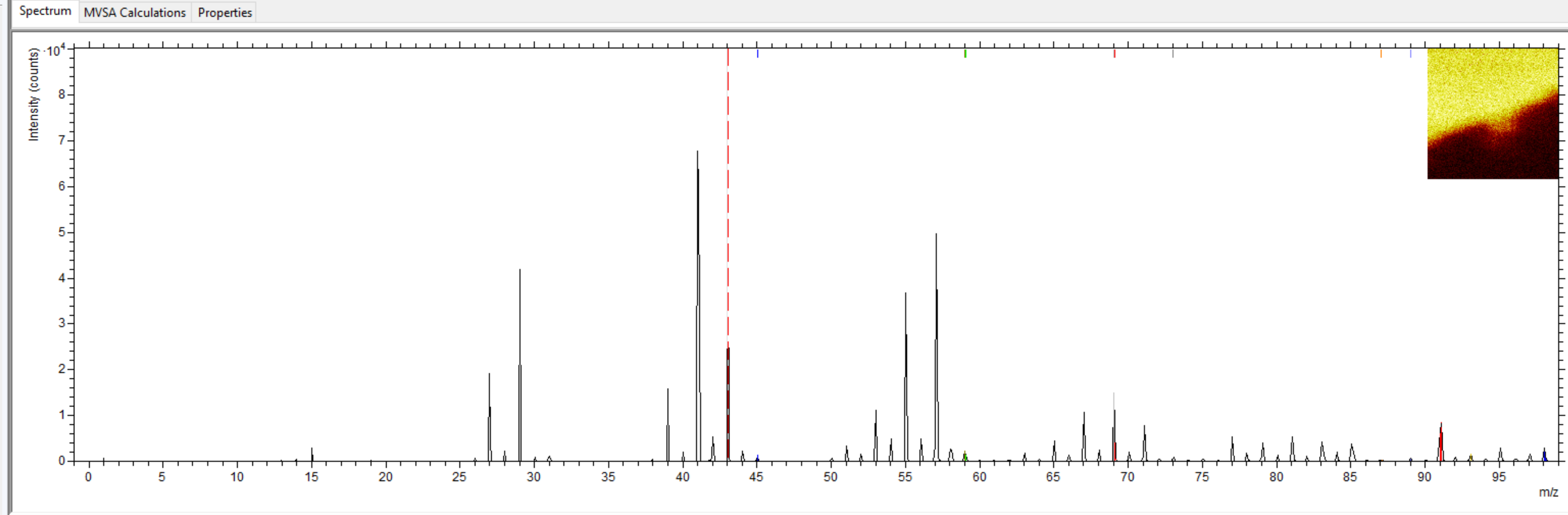
C_3H_7+	m/z	43.0567	Area (cts)	876,855	Explained (%)	100.0	Resolution	8,083
Check...	Dev. (ppm)	58.4	Counts / Shot	0.7433	Peak Difference...		Width (ns)	1.38

Compilations

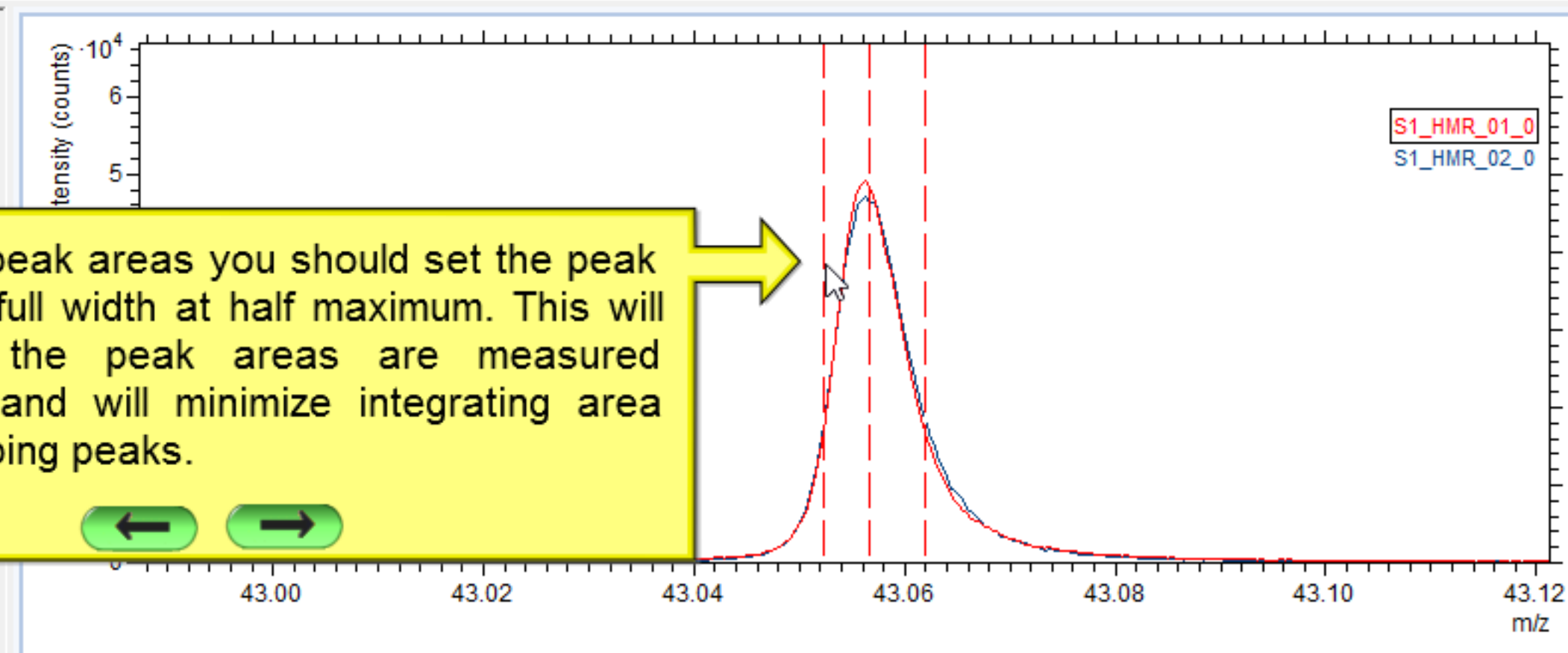
Spectra

S1\_HMR\_01\_0 \*  
<No Sample Name> (S1\_HMR\_01) \*

S1\_HMR\_02\_0  
<No Sample Name> (S1\_HMR\_02)



V	No.	m/z	Area / cts	Color	Peak Label	Description
<b>Mass Intervals</b>						
<input checked="" type="checkbox"/>	2	45.0352	32713	Blue	45.03 u	PEG
<input checked="" type="checkbox"/>	3	59.0136	51836	Olive	59.01 u	PMMA
<input checked="" type="checkbox"/>	4	59.0511	37293	Green	59.05 u	PEG
<input checked="" type="checkbox"/>	5	69.0376	118182	Grey	69.04 u	PMMA
<input checked="" type="checkbox"/>	6	69.0762	360824	Red	69.08 u	
<input checked="" type="checkbox"/>	7	73.0342	16483	Dark Grey	73.03 u	PEG
<input checked="" type="checkbox"/>	8	87.0478	6134	Orange	87.05 u	PEG
<input checked="" type="checkbox"/>	9	89.0617	3870	Light Blue	89.06 u	PEG
<input checked="" type="checkbox"/>	10	91.0566	238681	Red	91.05 u	PS
<input checked="" type="checkbox"/>	11	93.0730	55170	Yellow	93.07 u	PMMA
<input checked="" type="checkbox"/>	12	98.0763	70210	Blue	98.07 u	GTP
<input checked="" type="checkbox"/>	13	101.0605	26893	Green	101.06 u	PMMA
<input checked="" type="checkbox"/>	14	103.0490	34262	Light Green	103.05 u	PEG
<input checked="" type="checkbox"/>	15	105.0688	63931	Purple	105.07 u	PS



If looking at peak areas you should set the peak limits at the full width at half maximum. This will make sure the peak areas are measured consistently and will minimize integrating area from overlapping peaks.

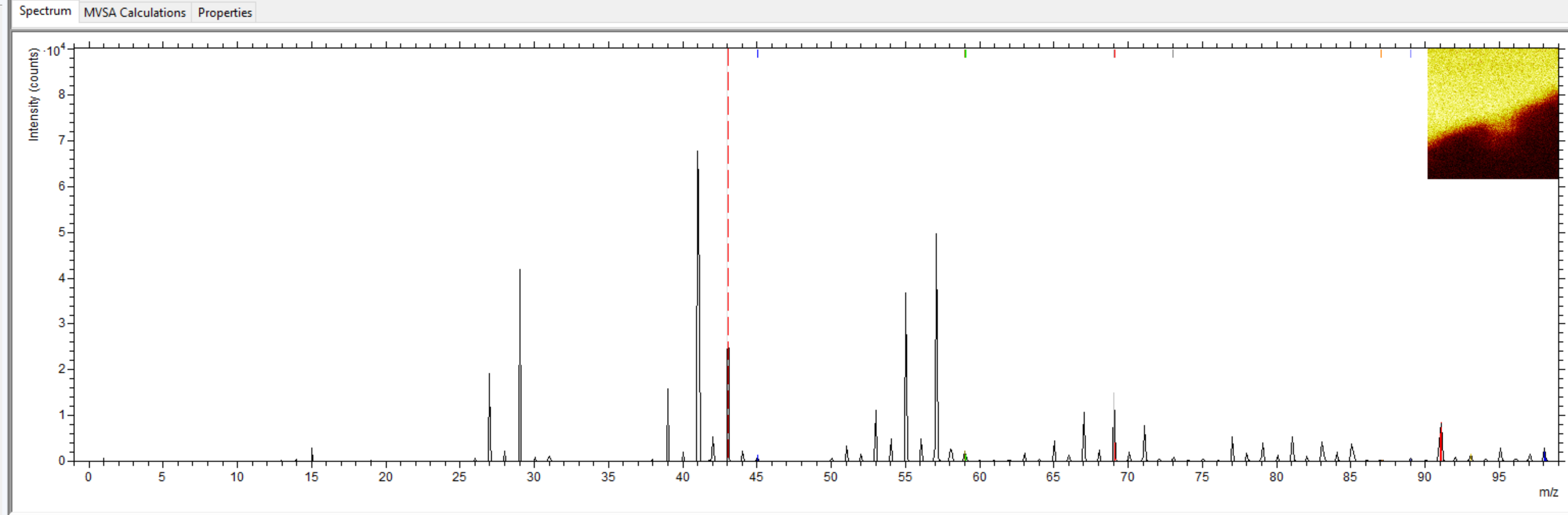
C_3H_7+	m/z	43.0564	Area (cts)	845,732	Explained (%)	100.0	Resolution	8,116
Check...	Dev. (ppm)	50.3	Counts / Shot	0.7169	Peak Difference...		Width (ns)	1.37

Compilations

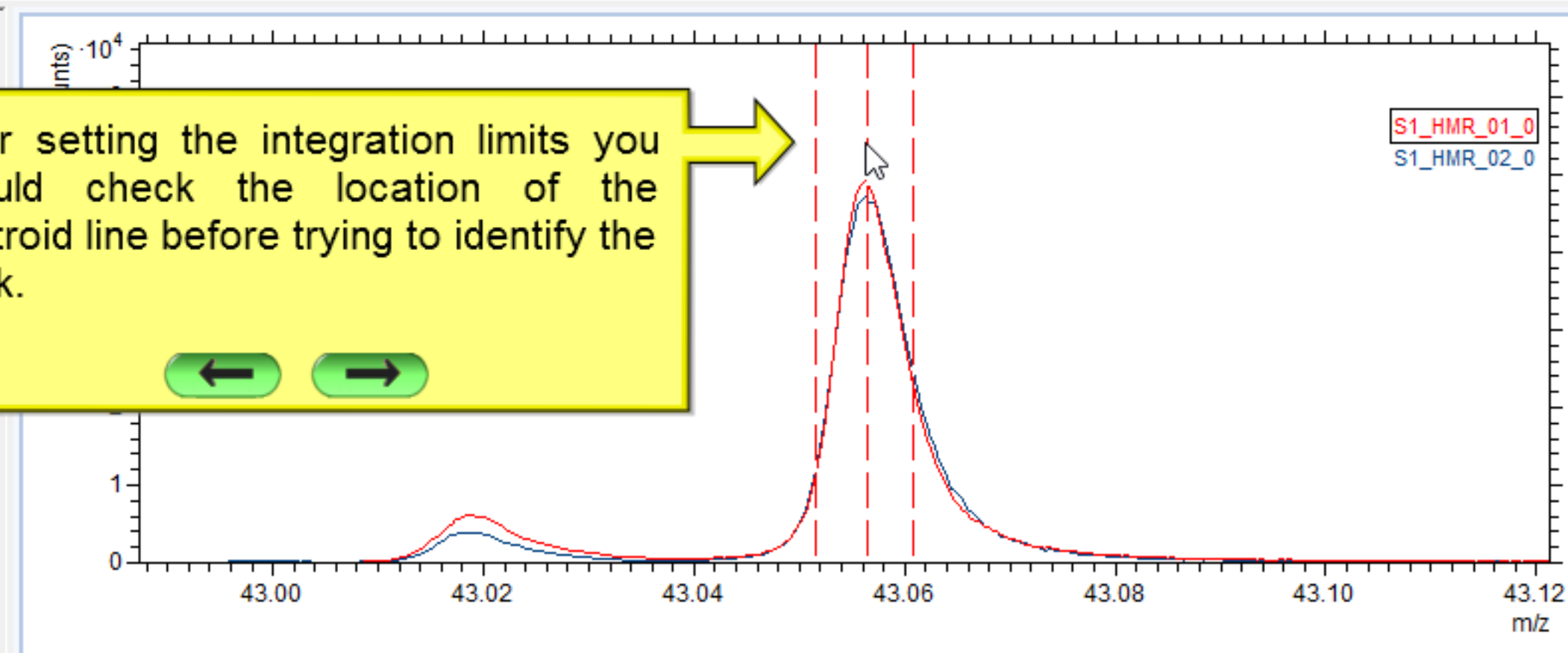
Spectra

S1\_HMR\_01\_0 \*  
<No Sample Name> (S1\_HMR\_01) \*

S1\_HMR\_02\_0  
<No Sample Name> (S1\_HMR\_02)



V	No.	m / z	Area / cts	Color	Peak Label	Description
<b>Mass Intervals</b>						
<input checked="" type="checkbox"/>	2	45.0352	32713	Blue	45.03 u	PEG
<input checked="" type="checkbox"/>	3	59.0136	51836	Olive	59.01 u	PMMA
<input checked="" type="checkbox"/>	4	59.0511	37293	Green	59.05 u	PEG
<input checked="" type="checkbox"/>	5	69.0376	118182	Grey	69.04 u	PMMA
<input checked="" type="checkbox"/>	6	69.0762	360824	Red	69.08 u	
<input checked="" type="checkbox"/>	7	73.0342	16483	Dark Grey	73.03 u	PEG
<input checked="" type="checkbox"/>	8	87.0478	6134	Orange	87.05 u	PEG
<input checked="" type="checkbox"/>	9	89.0617	3870	Light Blue	89.06 u	PEG
<input checked="" type="checkbox"/>	10	91.0566	238681	Red	91.05 u	PS
<input checked="" type="checkbox"/>	11	93.0730	55170	Yellow	93.07 u	PMMA
<input checked="" type="checkbox"/>	12	98.0763	70210	Blue	98.07 u	GTP
<input checked="" type="checkbox"/>	13	101.0605	26893	Green	101.06 u	PMMA
<input checked="" type="checkbox"/>	14	103.0490	34262	Light Green	103.05 u	PEG
<input checked="" type="checkbox"/>	15	105.0688	63931	Purple	105.07 u	PS



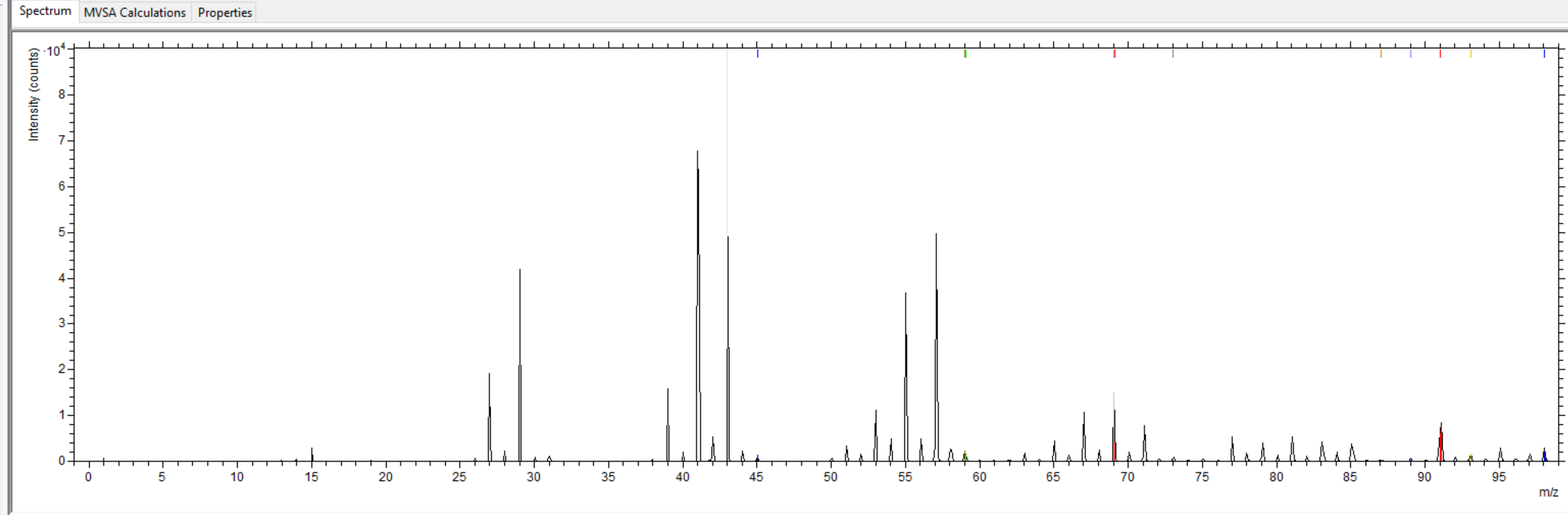
After setting the integration limits you should check the location of the centroid line before trying to identify the peak.

m/z: 43.0205 Area (cts): 173,776 Explained (%): 80.6 Resolution: 5,219  
 Dev. (ppm): -44.7 Counts / Shot: 0.1473 Peak Difference... Width (ns): 2.13

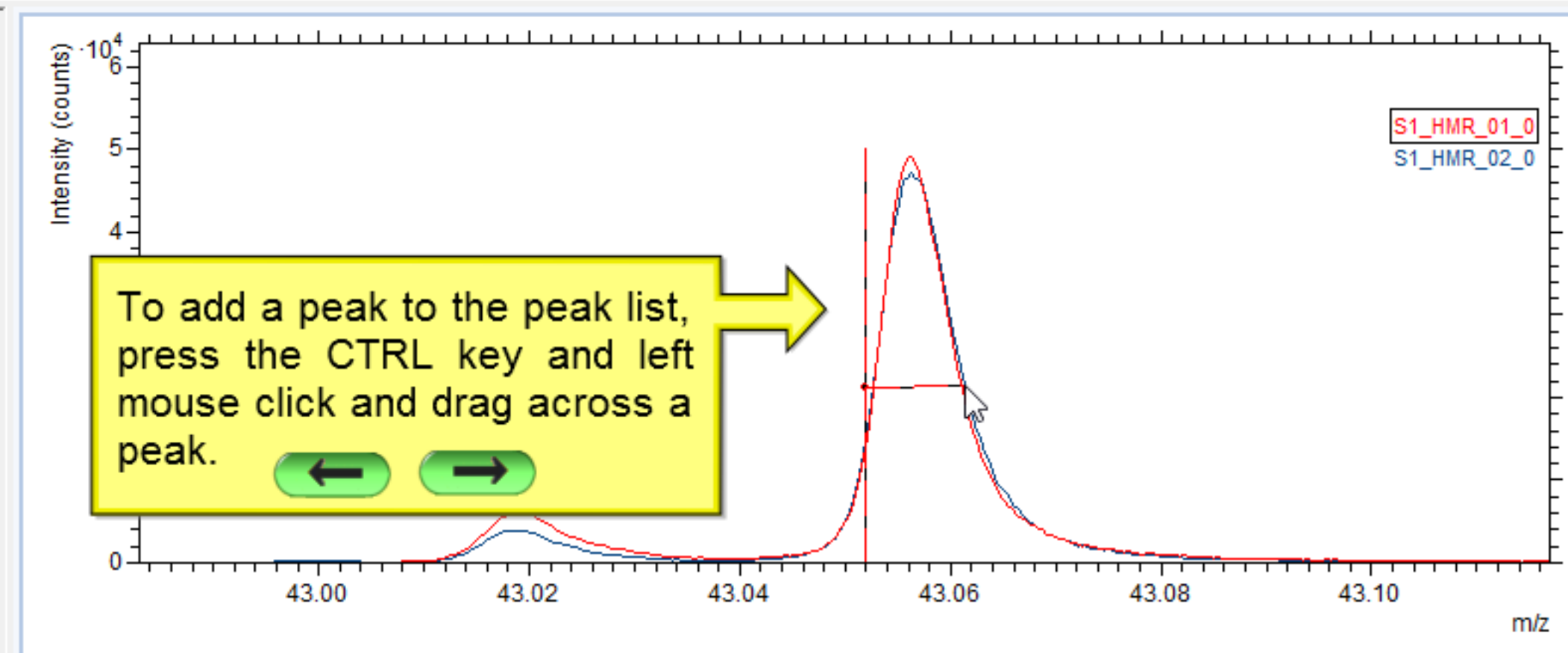
Compilations

Spectra

- Spectra
- S1\_HMR\_01\_0 \*
    - <No Sample Name> (S1\_HMR\_01) \*
  - S1\_HMR\_02\_0
    - <No Sample Name> (S1\_HMR\_02)



V	No.	m / z	Area / cts	Color	Peak Label	Description
<b>Mass Intervals</b>						
<input checked="" type="checkbox"/>	2	45.0352	32713	Blue	45.03 u	PEG
<input checked="" type="checkbox"/>	3	59.0136	51836	Olive	59.01 u	PMMA
<input checked="" type="checkbox"/>	4	59.0511	37293	Green	59.05 u	PEG
<input checked="" type="checkbox"/>	5	69.0376	118182	Grey	69.04 u	PMMA
<input checked="" type="checkbox"/>	6	69.0762	360824	Red	69.08 u	
<input checked="" type="checkbox"/>	7	73.0342	16483	Dark Grey	73.03 u	PEG
<input checked="" type="checkbox"/>	8	87.0478	6134	Orange	87.05 u	PEG
<input checked="" type="checkbox"/>	9	89.0617	3870	Light Blue	89.06 u	PEG
<input checked="" type="checkbox"/>	10	91.0566	238681	Red	91.05 u	PS
<input checked="" type="checkbox"/>	11	93.0730	55170	Yellow	93.07 u	PMMA
<input checked="" type="checkbox"/>	12	98.0763	70210	Blue	98.07 u	GTP
<input checked="" type="checkbox"/>	13	101.0605	26893	Green	101.06 u	PMMA
<input checked="" type="checkbox"/>	14	103.0490	34262	Light Green	103.05 u	PEG
<input checked="" type="checkbox"/>	15	105.0688	63931	Purple	105.07 u	PS



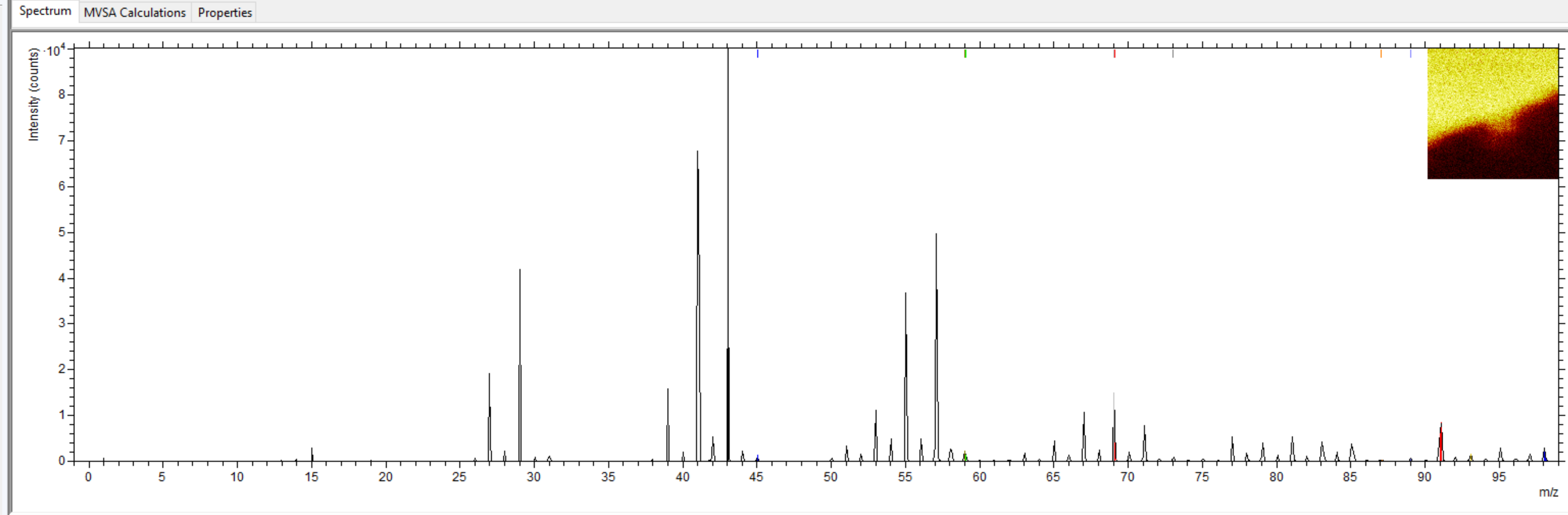
m/z  Area (cts)  Explained (%)  Resolution   
 Dev. (ppm)  Counts / Shot  Peak Difference...  Width (ns)

Compilations

Spectra

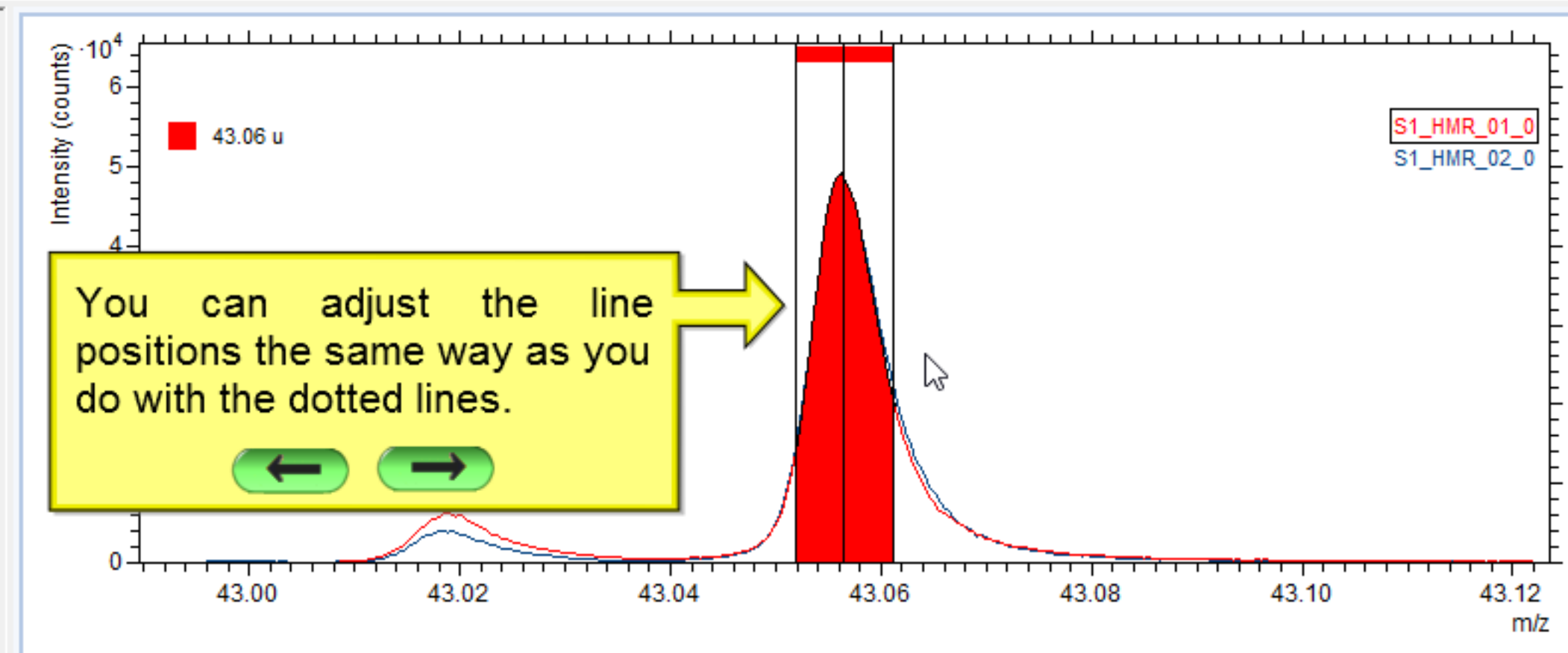
S1\_HMR\_01\_0 \*  
<No Sample Name> (S1\_HMR\_01) \*

S1\_HMR\_02\_0  
<No Sample Name> (S1\_HMR\_02)



MassListPos

V	No.	m / z	Area / cts	Color	Peak Label	Description
<input checked="" type="checkbox"/>	22	139.0715	17478	Blue	139.07 u	PMMA
<input checked="" type="checkbox"/>	23	151.0745	4167	Purple	151.07 u	PEG
<input checked="" type="checkbox"/>	24	165.0590	18022	Orange	165.05 u	PS
<input checked="" type="checkbox"/>	25	171.0820	31193	Red	171.07 u	GTP
<input checked="" type="checkbox"/>	26	175.1104	15764	Olive	175.10 u	PEG
<input checked="" type="checkbox"/>	27	186.1037	8941	Pink	186.09 u	PMMA
<input checked="" type="checkbox"/>	28	193.1071	8863	Light Blue	193.10 u	PS
<input checked="" type="checkbox"/>	29	202.0650	7833	Light Green	202.06 u	PS
<input checked="" type="checkbox"/>	30	215.0729	7554	Dark Green	215.07 u	PS
<input checked="" type="checkbox"/>	31	239.2506	24382	Brown	239.24 u	GTP
<input checked="" type="checkbox"/>	32	287.1483	526	Grey	287.13 u	PEG
<input checked="" type="checkbox"/>	33	313.2827	38824	Yellow	313.26 u	GTP
<input checked="" type="checkbox"/>	34	331.1772	331	Green	331.18 u	PEG
<input checked="" type="checkbox"/>	35	367.2733	10864	Magenta	367.25 u	GTP
<input checked="" type="checkbox"/>	36	551.5105	170106	Blue	551.47 u	GTP



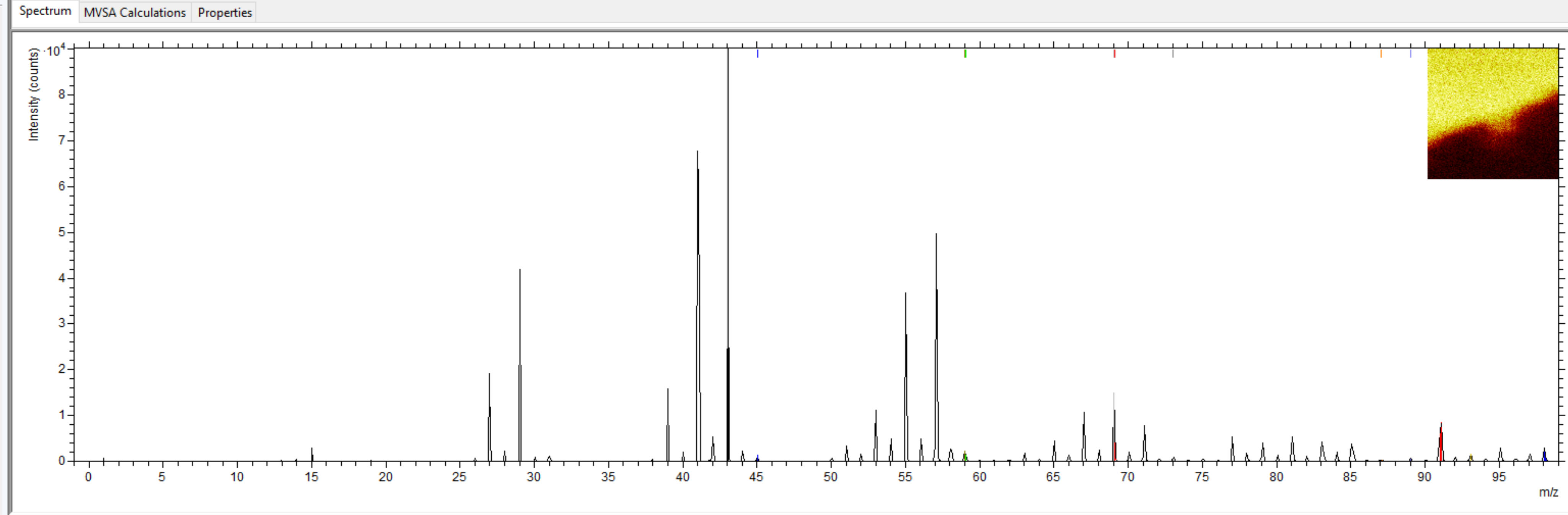
m/z  ?  
 Area (cts)  Explained (%)  
 Counts / Shot  Peak Difference...  
 Resolution   
 Width (ns)

Compilations

Spectra

S1\_HMR\_01\_0 \*  
<No Sample Name> (S1\_HMR\_01) \*

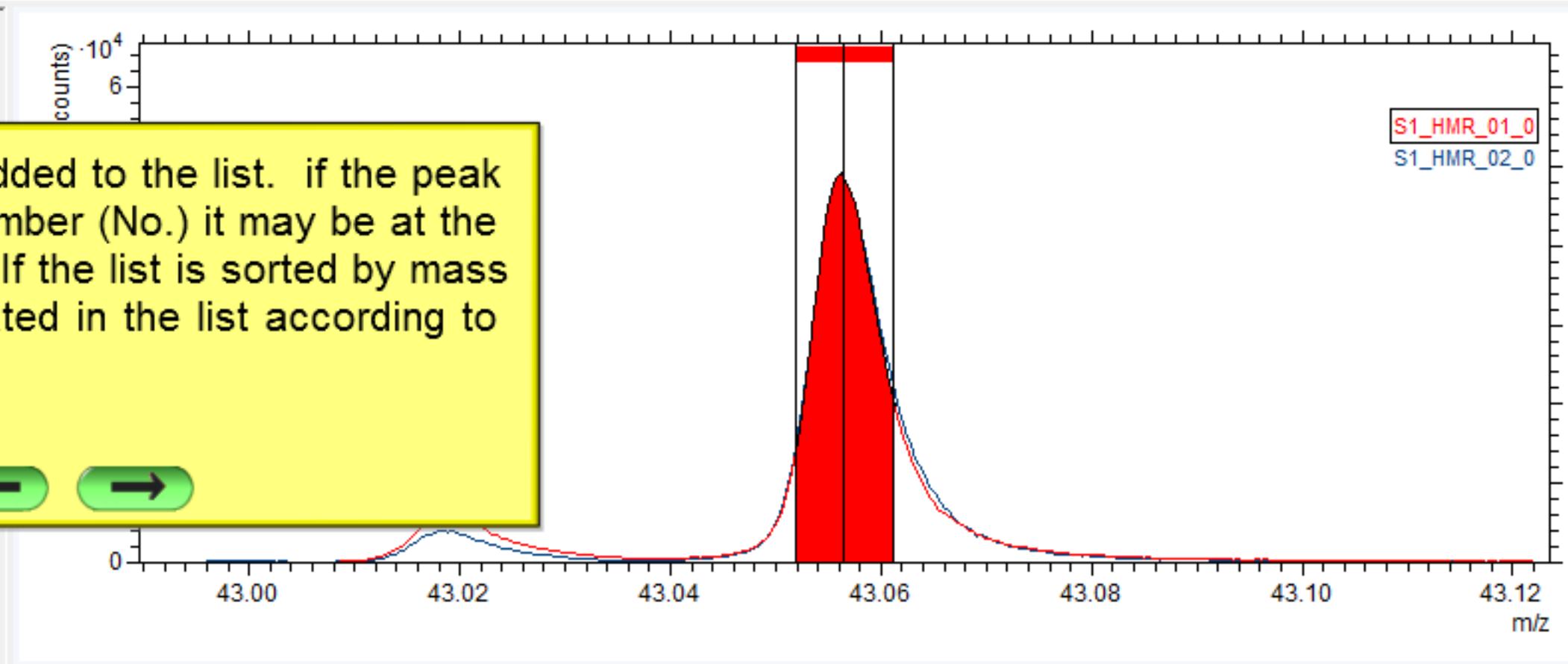
S1\_HMR\_02\_0  
<No Sample Name> (S1\_HMR\_02)



MassListPos

V	No.	m/z	Area / cts	Color	Peak Label	Description
<b>Mass Intervals</b>						
<input checked="" type="checkbox"/>	37	43.0565	854832	Red	43.06 u	
<input checked="" type="checkbox"/>	2	45.0352	32713	Blue	45.03 u	PEG
<input checked="" type="checkbox"/>	3	59.0136	51836	Olive	59.01 u	PMMA
<input checked="" type="checkbox"/>	4	59.0511	37293	Green	59.05 u	PEG
<input checked="" type="checkbox"/>	5	69.0376	118182	Grey	69.04 u	PMMA
<input checked="" type="checkbox"/>	6	69.0762	360824	Red	69.08 u	
<input checked="" type="checkbox"/>	7	73.0342	16483	Grey	73.03 u	PEG
<input checked="" type="checkbox"/>	8	87.0478	6134	Orange	87.05 u	PEG
<input checked="" type="checkbox"/>	9	89.0617	3870	Blue	89.06 u	PEG
<input checked="" type="checkbox"/>	10	91.0566	238681	Red	91.05 u	PS
<input checked="" type="checkbox"/>	11	93.0730	55170	Yellow	93.07 u	PMMA
<input checked="" type="checkbox"/>	12	98.0763	70210	Blue	98.07 u	GTP
<input checked="" type="checkbox"/>	13	101.0605	26893	Green	101.06 u	PMMA
<input checked="" type="checkbox"/>	14	103.0490	34262	Light Green	103.05 u	PEG

The peak will be added to the list. If the peak list is sorted by number (No.) it may be at the bottom of the list. If the list is sorted by mass (m/z) it will be located in the list according to its mass.



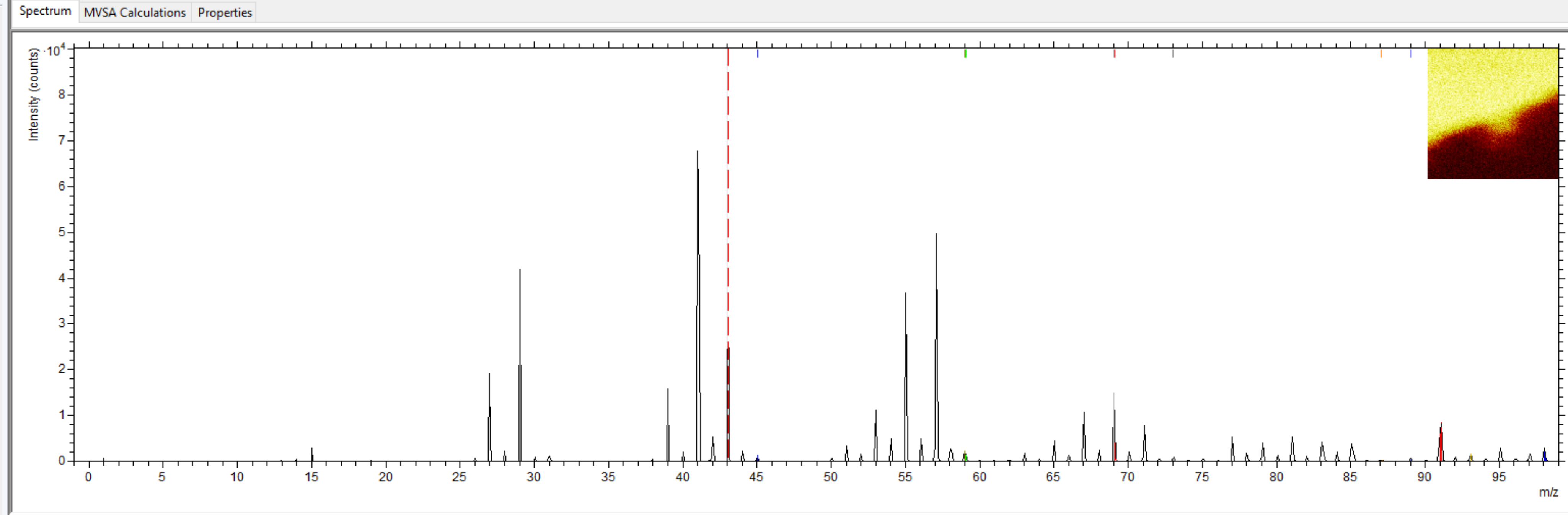
C_3H_7+	m/z	43.0572	Area (cts)	1,153,550	Explained (%)	100.0	Resolution	5,856
Check...	Dev. (ppm)	68.6	Counts / Shot	0.9779	Peak Difference...		Width (ns)	1.90

Compilations

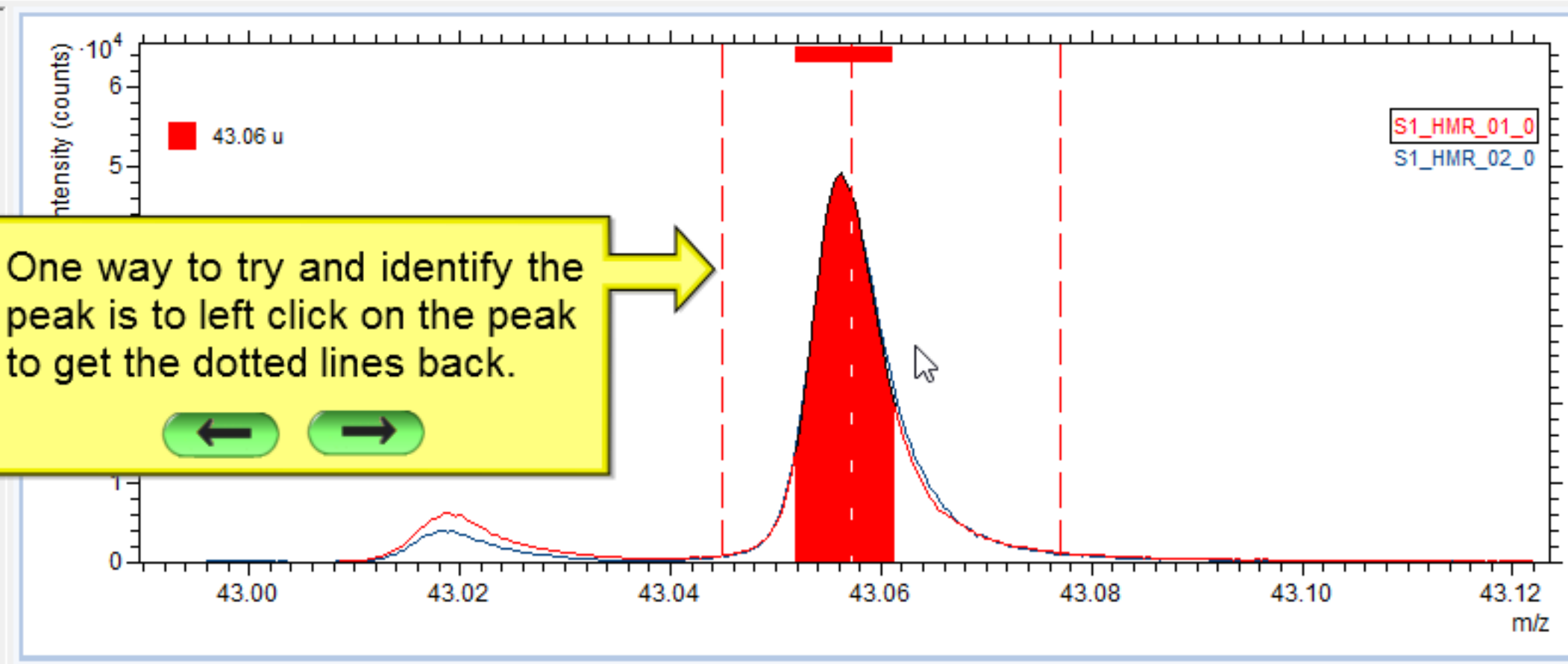
Spectra

S1\_HMR\_01\_0 \*  
<No Sample Name> (S1\_HMR\_01) \*

S1\_HMR\_02\_0  
<No Sample Name> (S1\_HMR\_02)



V	No.	m/z	Area / cts	Color	Peak Label	Description
<input checked="" type="checkbox"/>	37	43.0565	854832	Red	43.06 u	
<input checked="" type="checkbox"/>	2	45.0352	32713	Blue	45.03 u	PEG
<input checked="" type="checkbox"/>	3	59.0136	51836	Olive	59.01 u	PMMA
<input checked="" type="checkbox"/>	4	59.0511	37293	Green	59.05 u	PEG
<input checked="" type="checkbox"/>	5	69.0376	118182	Grey	69.04 u	PMMA
<input checked="" type="checkbox"/>	6	69.0762	360824	Red	69.08 u	
<input checked="" type="checkbox"/>	7	73.0342	16483	Grey	73.03 u	PEG
<input checked="" type="checkbox"/>	8	87.0478	6134	Orange	87.05 u	PEG
<input checked="" type="checkbox"/>	9	89.0617	3870	Blue	89.06 u	PEG
<input checked="" type="checkbox"/>	10	91.0566	238681	Red	91.05 u	PS
<input checked="" type="checkbox"/>	11	93.0730	55170	Yellow	93.07 u	PMMA
<input checked="" type="checkbox"/>	12	98.0763	70210	Blue	98.07 u	GTP
<input checked="" type="checkbox"/>	13	101.0605	26893	Green	101.06 u	PMMA
<input checked="" type="checkbox"/>	14	103.0490	34262	Light Green	103.05 u	PEG



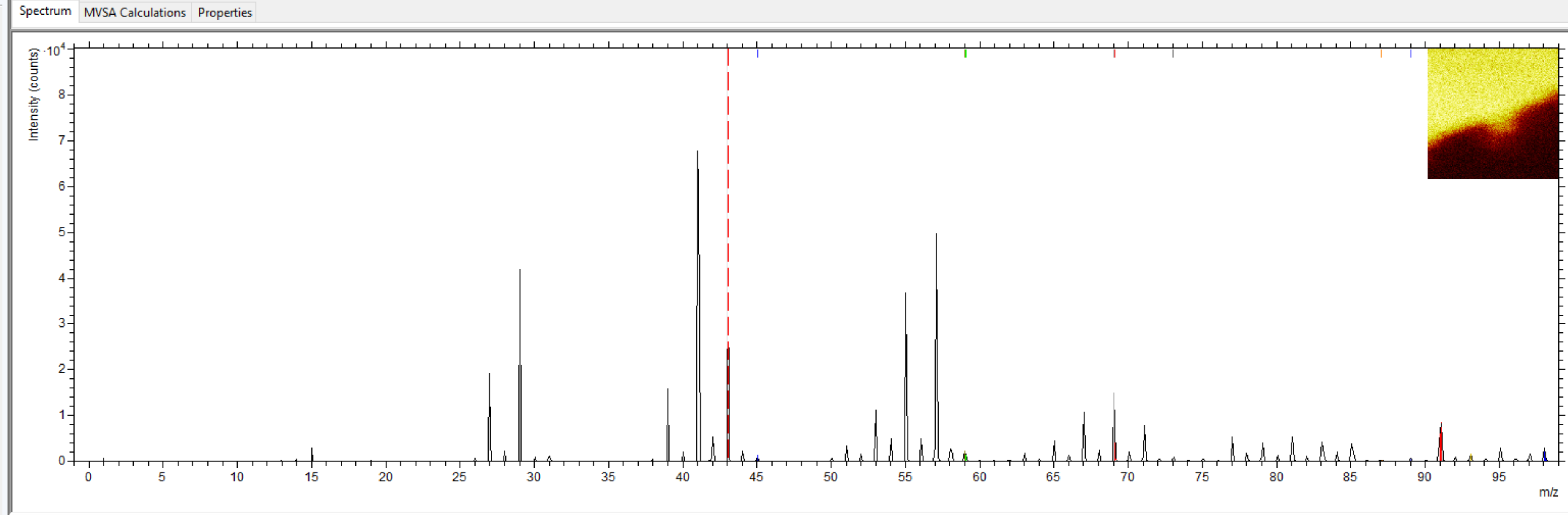
C_3H_7+	m/z	43.0563	Area (cts)	923,468	Explained (%)	100.0	Resolution	7,028
Check...	Dev. (ppm)	47.2	Counts / Shot	0.7828	Peak Difference...		Width (ns)	1.59

Compilations

Spectra

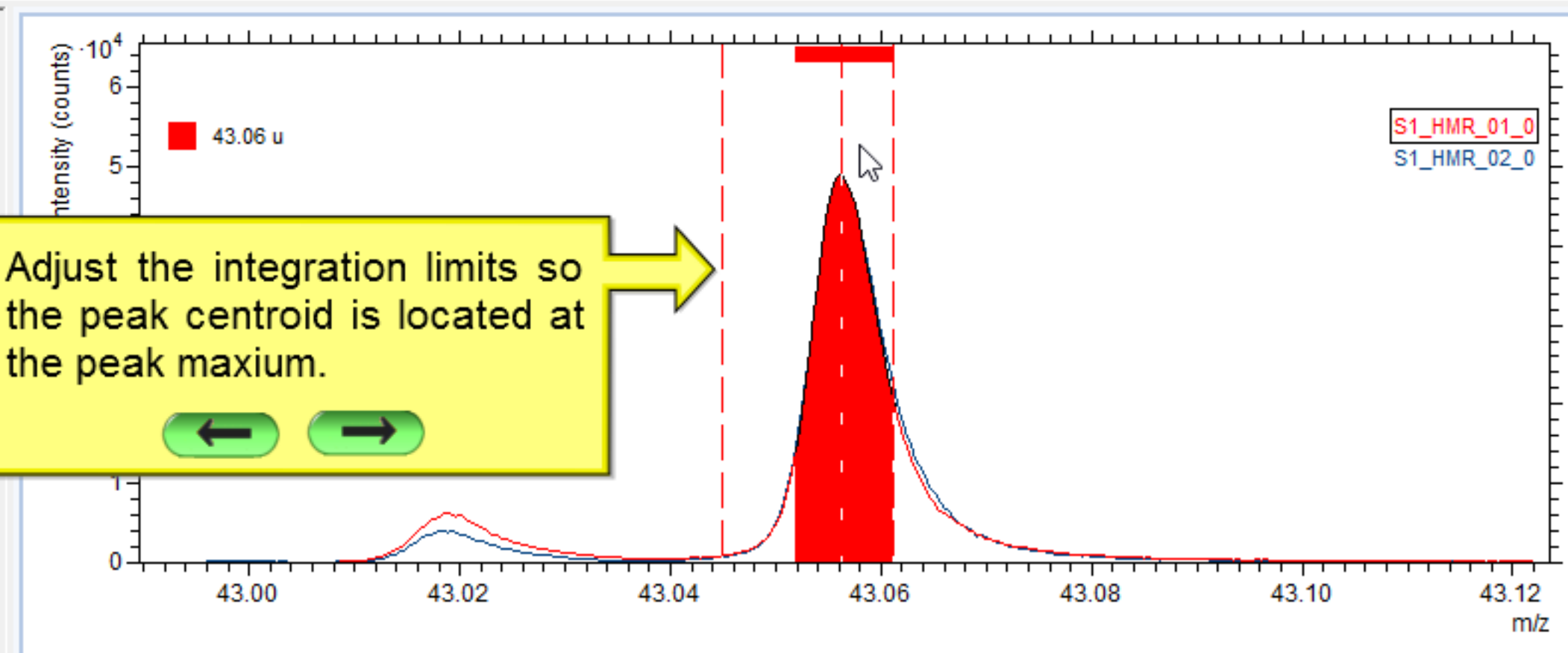
S1\_HMR\_01\_0 \*  
<No Sample Name> (S1\_HMR\_01) \*

S1\_HMR\_02\_0  
<No Sample Name> (S1\_HMR\_02)



MassListPos

V	No.	m/z	Area / cts	Color	Peak Label	Description
<b>Mass Intervals</b>						
<input checked="" type="checkbox"/>	37	43.0565	854832	Red	43.06 u	
<input checked="" type="checkbox"/>	2	45.0352	32713	Blue	45.03 u	PEG
<input checked="" type="checkbox"/>	3	59.0136	51836	Olive	59.01 u	PMMA
<input checked="" type="checkbox"/>	4	59.0511	37293	Green	59.05 u	PEG
<input checked="" type="checkbox"/>	5	69.0376	118182	Grey	69.04 u	PMMA
<input checked="" type="checkbox"/>	6	69.0762	360824	Red	69.08 u	
<input checked="" type="checkbox"/>	7	73.0342	16483	Grey	73.03 u	PEG
<input checked="" type="checkbox"/>	8	87.0478	6134	Orange	87.05 u	PEG
<input checked="" type="checkbox"/>	9	89.0617	3870	Blue	89.06 u	PEG
<input checked="" type="checkbox"/>	10	91.0566	238681	Red	91.05 u	PS
<input checked="" type="checkbox"/>	11	93.0730	55170	Yellow	93.07 u	PMMA
<input checked="" type="checkbox"/>	12	98.0763	70210	Blue	98.07 u	GTP
<input checked="" type="checkbox"/>	13	101.0605	26893	Green	101.06 u	PMMA
<input checked="" type="checkbox"/>	14	103.0490	34262	Light Green	103.05 u	PEG



C\_3H\_7+ Check... Add Peak

← →

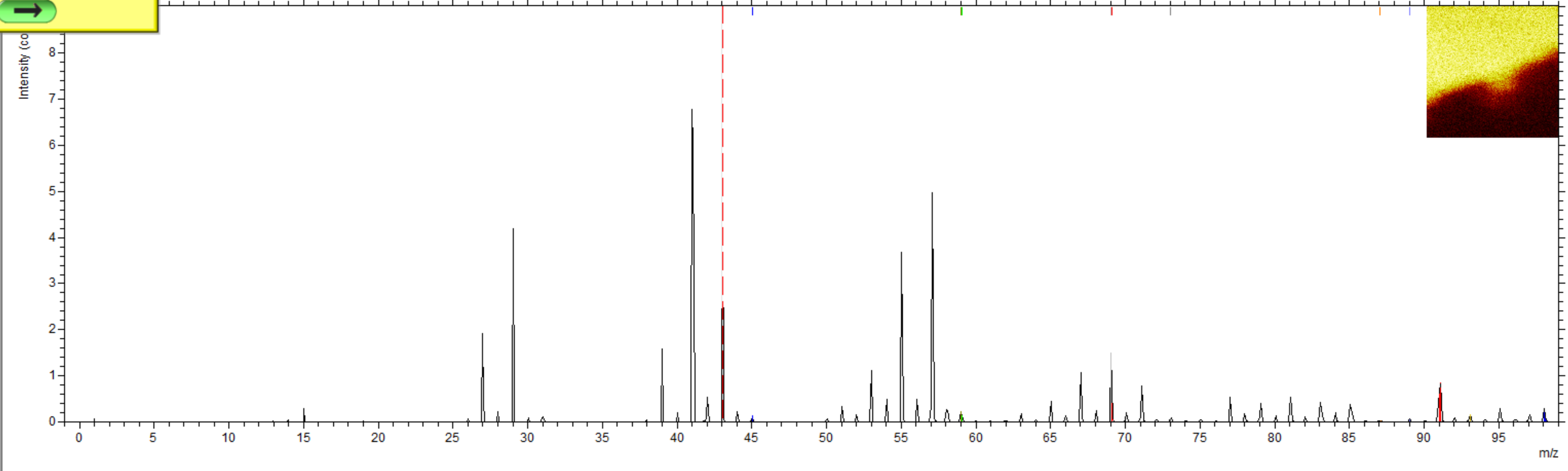
With the peak selected and the centroid at the peak maximum, press the 'Check...' button.

<input type="checkbox"/> Explained (%)	100.0	Resolution	7,028
Peak Difference...		Width (ns)	1.59

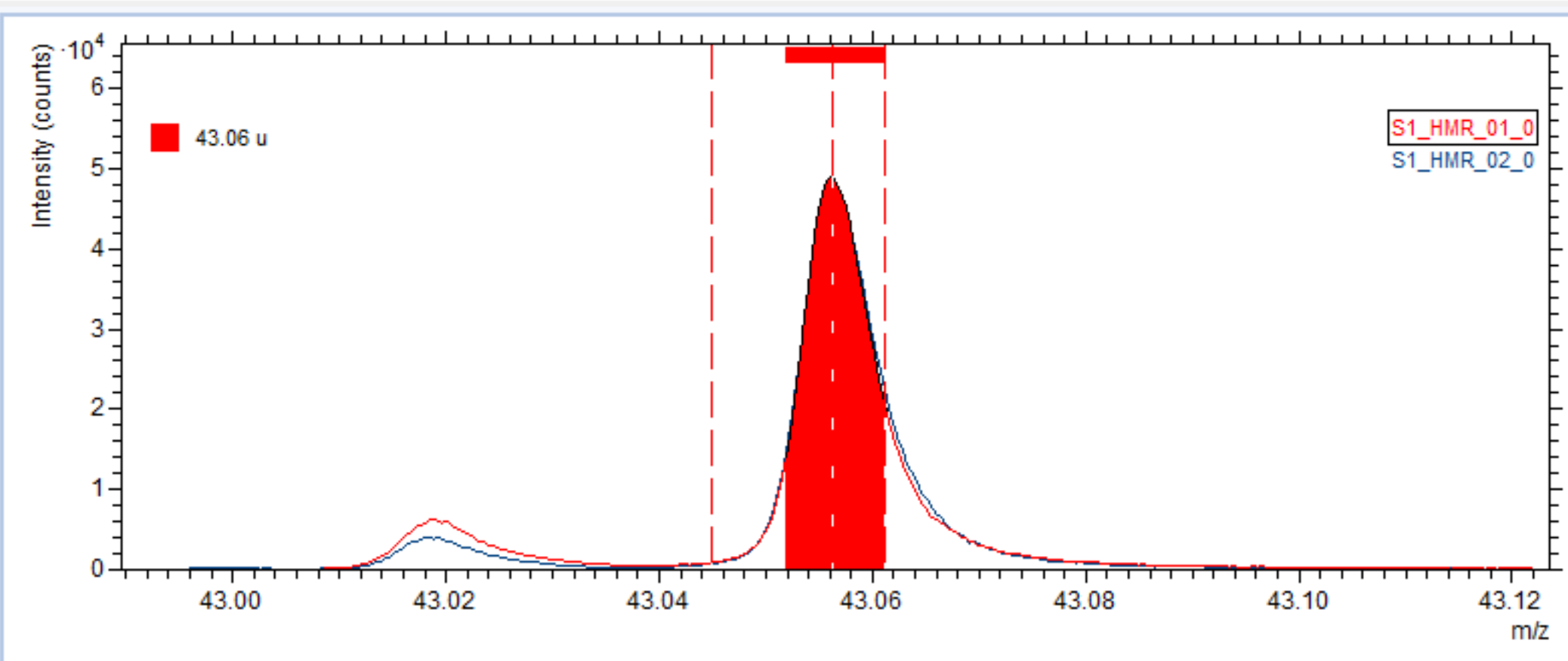
Spectra

S1\_HMR\_01\_0 \*  
<No Sample Name> (S1\_HMR\_01) \*

S1\_HMR\_02\_0  
<No Sample Name> (S1\_HMR\_02)



V	No.	m/z	Area / cts	Color	Peak Label	Description
<b>Mass Intervals</b>						
<input checked="" type="checkbox"/>	37	43.0565	854832	Red	43.06 u	
<input checked="" type="checkbox"/>	2	45.0352	32713	Blue	45.03 u	PEG
<input checked="" type="checkbox"/>	3	59.0136	51836	Olive	59.01 u	PMMA
<input checked="" type="checkbox"/>	4	59.0511	37293	Green	59.05 u	PEG
<input checked="" type="checkbox"/>	5	69.0376	118182	Grey	69.04 u	PMMA
<input checked="" type="checkbox"/>	6	69.0762	360824	Red	69.08 u	
<input checked="" type="checkbox"/>	7	73.0342	16483	Grey	73.03 u	PEG
<input checked="" type="checkbox"/>	8	87.0478	6134	Orange	87.05 u	PEG
<input checked="" type="checkbox"/>	9	89.0617	3870	Blue	89.06 u	PEG
<input checked="" type="checkbox"/>	10	91.0566	238681	Red	91.05 u	PS
<input checked="" type="checkbox"/>	11	93.0730	55170	Yellow	93.07 u	PMMA
<input checked="" type="checkbox"/>	12	98.0763	70210	Blue	98.07 u	GTP
<input checked="" type="checkbox"/>	13	101.0605	26893	Green	101.06 u	PMMA
<input checked="" type="checkbox"/>	14	103.0490	34262	Light Green	103.05 u	PEG



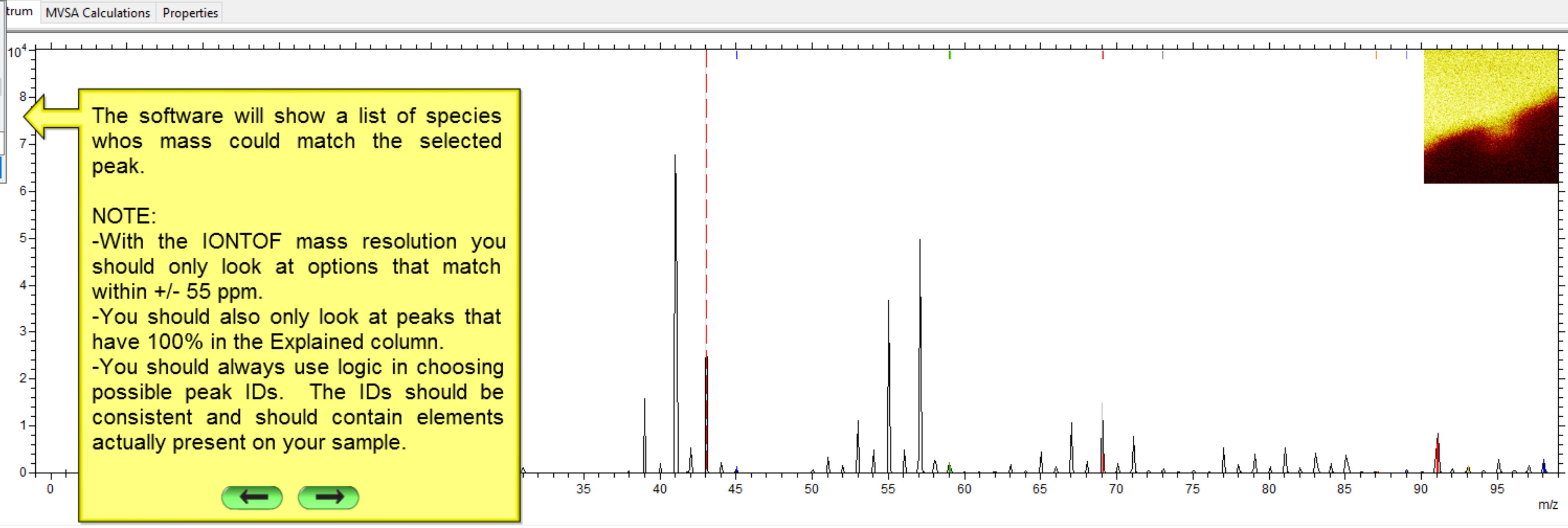
C <sub>3</sub> H <sub>7</sub> <sup>+</sup>	m/z	43.0563	Area (cts)	923,468	Explained (%)	100.0	Resolution	7,028
	Dev. (ppm)	47.2	Counts / Shot	0.7828	Peak Difference...		Width (ns)	1.59

Assignments	m/z	Dev. (ppm)	Explained (%)	Library
C <sub>2</sub> H <sub>5</sub> N <sup>+</sup>	43.0417	339.4	100.0	
C <sub>2</sub> <sup>13</sup> CH <sub>6</sub> <sup>+</sup>	43.0498	151.1	9.6	
C <sub>2</sub> H <sub>4</sub> CH <sub>3</sub> <sup>+</sup>	43.0542	47.2	100.0	
C <sub>3</sub> H <sub>7</sub> <sup>+</sup>	43.0542	47.2	100.0	

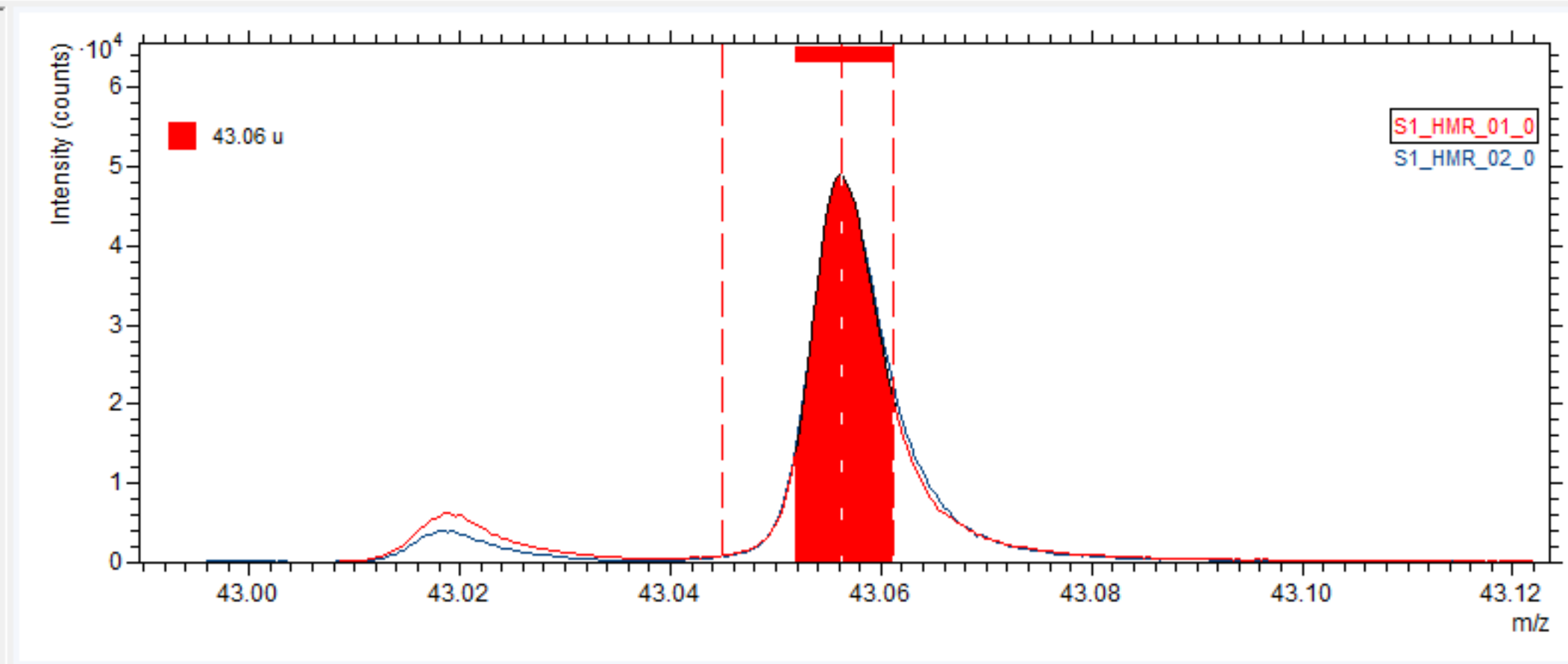
Buttons: Recalculate, Add Compound, Show Isotopes, Cancel, OK

S1\_HMR\_01\_0 \*  
 <No Sample Name> (S1\_HMR\_01) \*  
 S1\_HMR\_02\_0  
 <No Sample Name> (S1\_HMR\_02)

Mass Interval Lists  
 MassListPos (+)



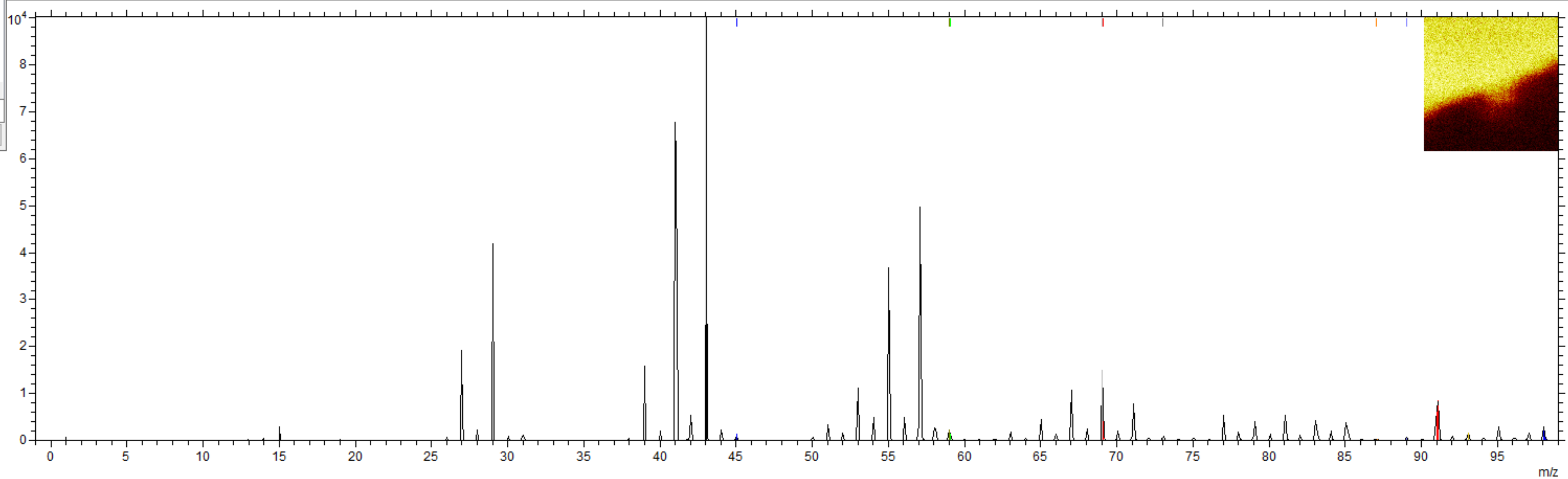
V	No.	m/z	Area / cts	Color	Peak Label	Description
✓	37	43.0565	854832	Red	43.06 u	
✓	2	45.0352	32713	Blue	45.03 u	PEG
✓	3	59.0136	51836	Olive	59.01 u	PMMA
✓	4	59.0511	37293	Green	59.05 u	PEG
✓	5	69.0376	118182	Grey	69.04 u	PMMA
✓	6	69.0762	360824	Red	69.08 u	
✓	7	73.0342	16483	Grey	73.03 u	PEG
✓	8	87.0478	6134	Orange	87.05 u	PEG
✓	9	89.0617	3870	Blue	89.06 u	PEG
✓	10	91.0566	238681	Red	91.05 u	PS
✓	11	93.0730	55170	Yellow	93.07 u	PMMA
✓	12	98.0763	70210	Blue	98.07 u	GTP
✓	13	101.0605	26893	Green	101.06 u	PMMA
✓	14	103.0490	34262	Light Green	103.05 u	PEG



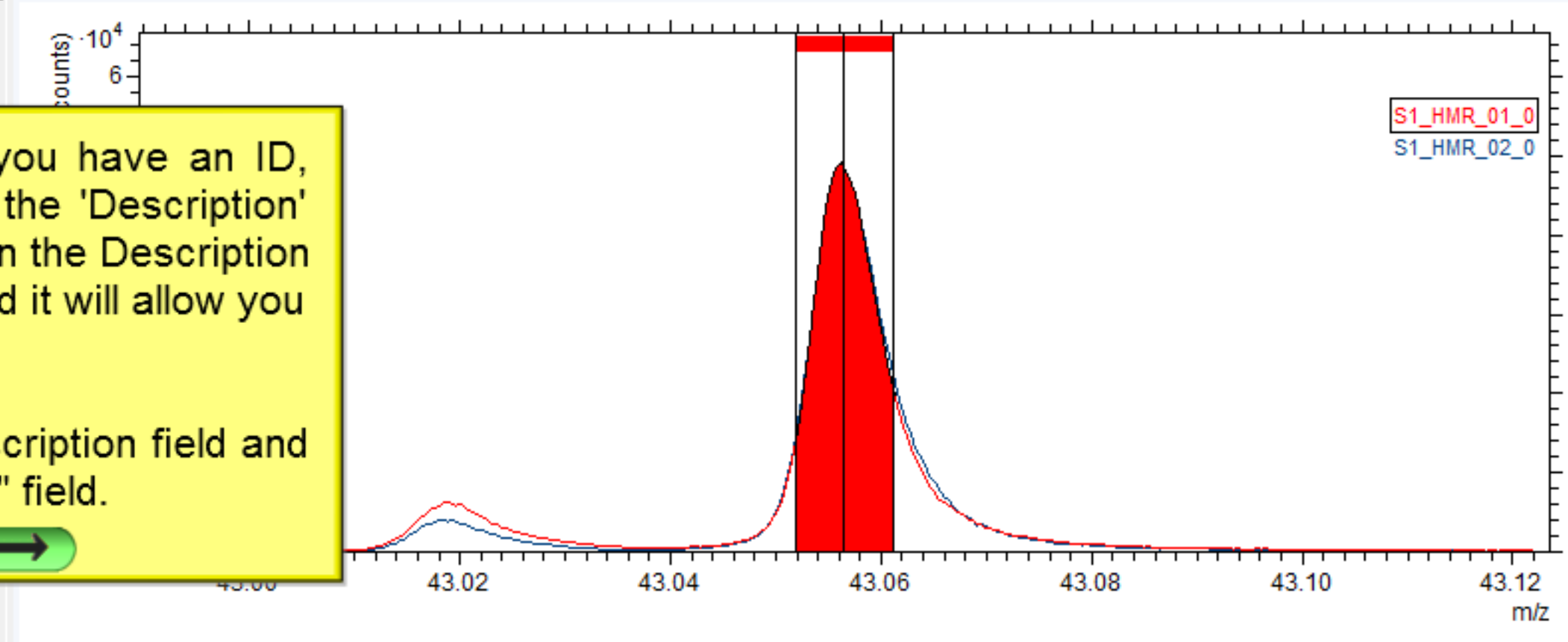
Explained (%)

Assignments m/z Dev. (ppm) Explained (%) Library Occurrence

- S1\_HMR\_01\_0 \*
  - <No Sample Name> (S1\_HMR\_01) \* +
  - S1\_HMR\_02\_0 +



V	No.	m / z	Area / cts	Color	Peak Label	Description
<b>Mass Intervals</b>						
<input checked="" type="checkbox"/>	37	43.0565	854832		43.06 u	C3H7
<input checked="" type="checkbox"/>	2	45.0352	32713		45.03 u	PEG
<input checked="" type="checkbox"/>	3	59.0136	51836		59.01 u	PMMA
<input checked="" type="checkbox"/>	4	59.0511	37293		59.05 u	PEG
<input checked="" type="checkbox"/>	5	69.0376	118182		69.04 u	PMMA
<input checked="" type="checkbox"/>	6	69.0762	360824		69.08 u	
<input checked="" type="checkbox"/>	7	73.0342	16483		73.03 u	PEG
<input checked="" type="checkbox"/>	8	87.0478	6134		87.05 u	PEG
<input checked="" type="checkbox"/>	9	89.0617	3870		89.06 u	PEG
<input checked="" type="checkbox"/>	10	91.0566	238681		91.05 u	PS
<input checked="" type="checkbox"/>	11	93.0730	55170		93.07 u	PMMA
<input checked="" type="checkbox"/>	12	98.0763	70210		98.07 u	GTP
<input checked="" type="checkbox"/>	13	101.0605	26893		101.06 u	PMMA
<input checked="" type="checkbox"/>	14	103.0490	34262		103.05 u	PEG



Once you believe you have an ID, you can enter it in the 'Description' column. Left click on the Description field for the peak and it will allow you to type.

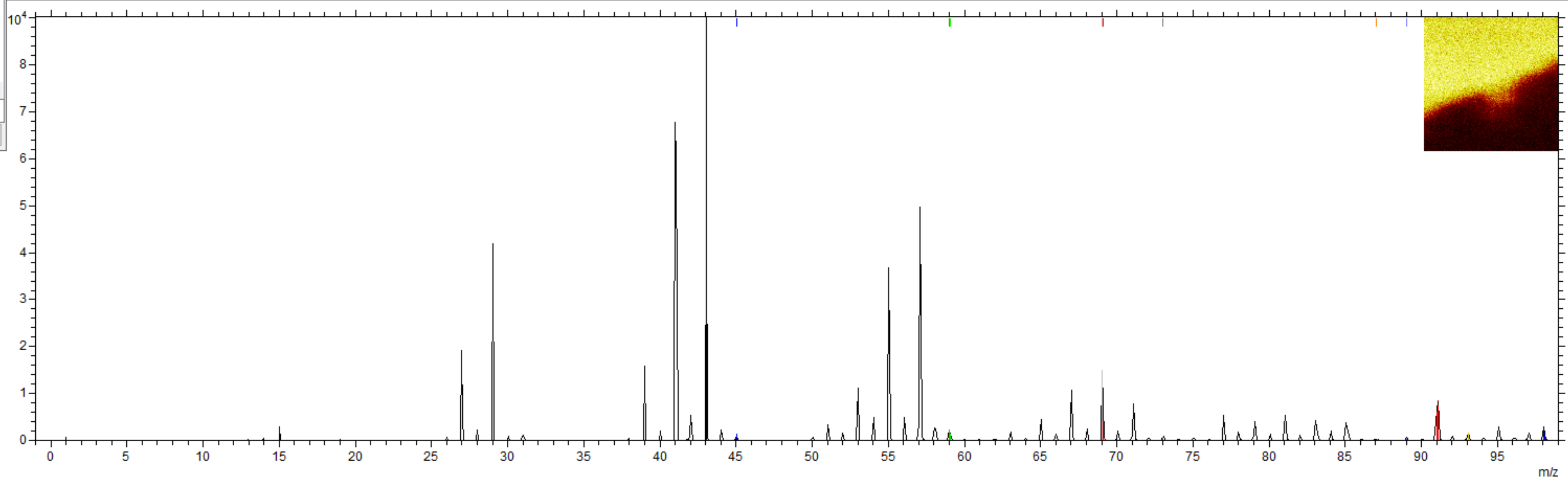
NOTE: Use the Description field and not the "Assginment" field.

← →

Explained (%)

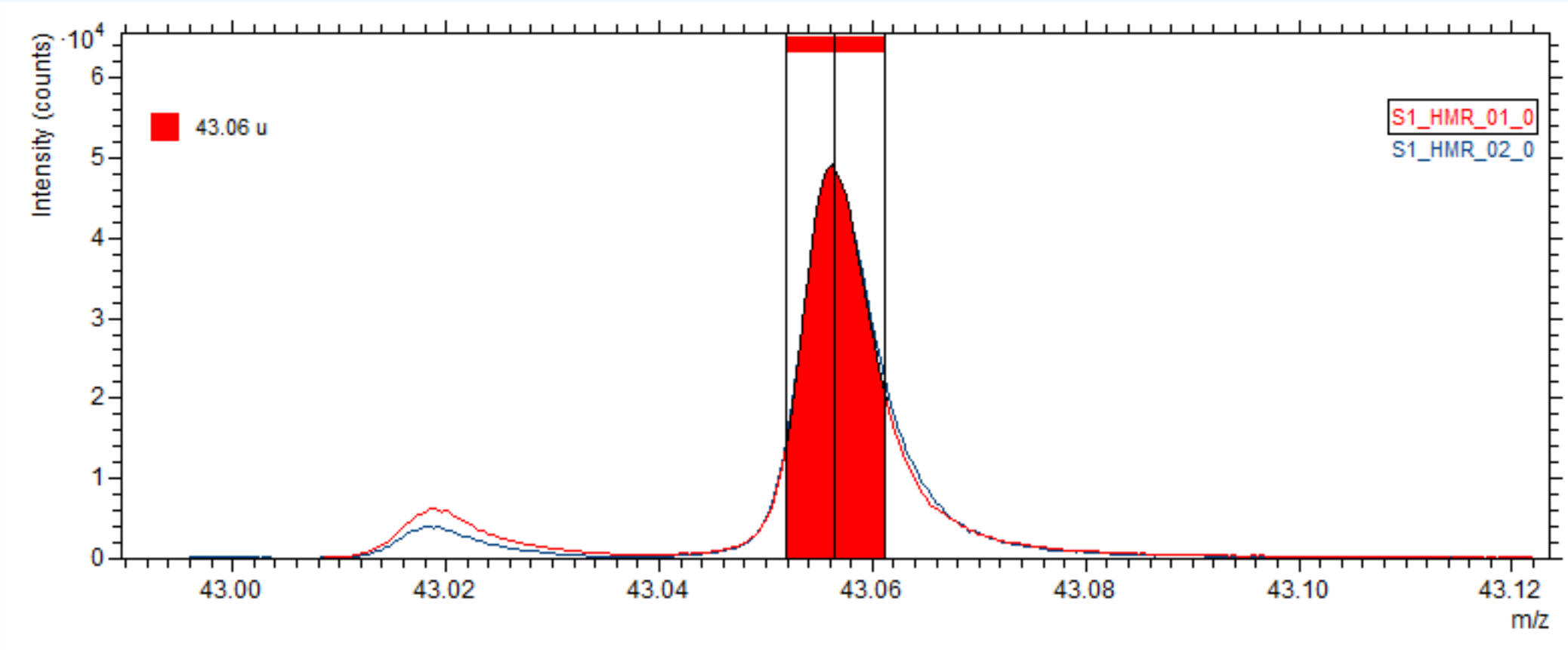
Assignments m/z Dev. (ppm) Explained (%) Library Occurrence

- S1\_HMR\_01\_0 \*
  - <No Sample Name> (S1\_HMR\_01) \* +
  - S1\_HMR\_02\_0 +



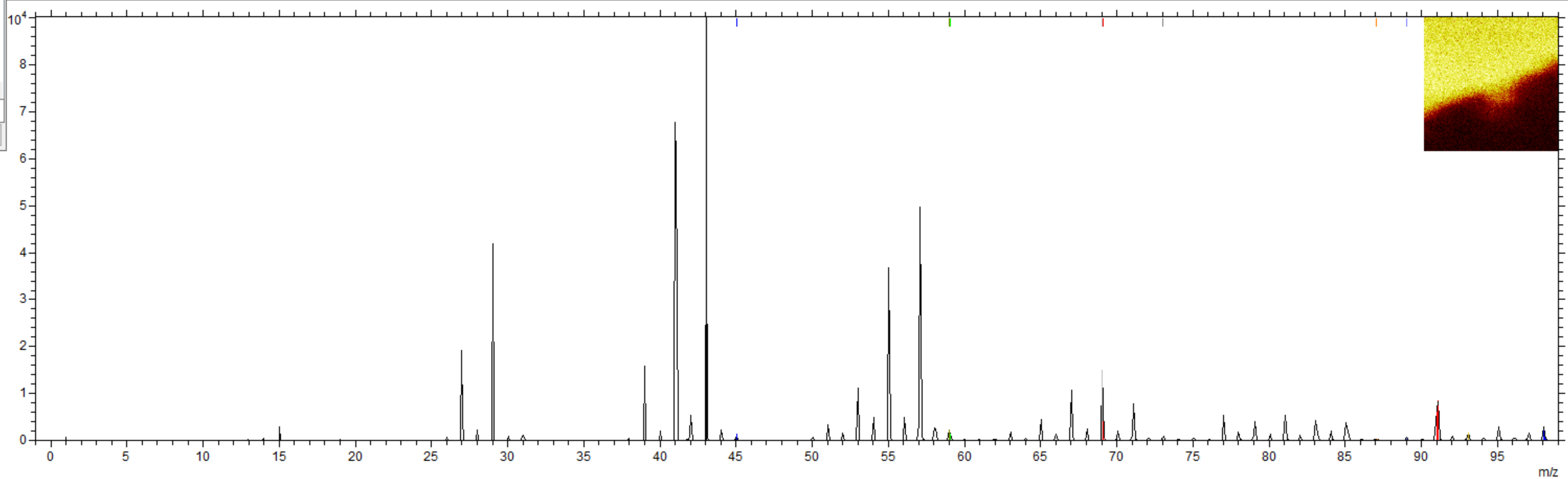
V	No.	m/z	Area / cts	Color	Peak Label	Description
<b>Mass Intervals</b>						
<input checked="" type="checkbox"/>	37	43.06				
<input checked="" type="checkbox"/>	2	45.0452				
<input checked="" type="checkbox"/>	3	59.0136				
<input checked="" type="checkbox"/>	4	59.0511				
<input checked="" type="checkbox"/>	5	69.0376				
<input checked="" type="checkbox"/>	6	69.0762				
<input checked="" type="checkbox"/>	7	73.0342				
<input checked="" type="checkbox"/>	8	87.0478				
<input checked="" type="checkbox"/>	9	89.0617	3870		89.06 u	PEG
<input checked="" type="checkbox"/>	10	91.0566	238681		91.05 u	PS
<input checked="" type="checkbox"/>	11	93.0730	55170		93.07 u	PMMA
<input checked="" type="checkbox"/>	12	98.0763	70210		98.07 u	GTP
<input checked="" type="checkbox"/>	13	101.0605	26893		101.06 u	PMMA
<input checked="" type="checkbox"/>	14	103.0490	34262		103.05 u	PEG

This list is sorted by mass. Note that the peak numbers are not in order. You need to change this before exporting any data or the masses will not be in order.



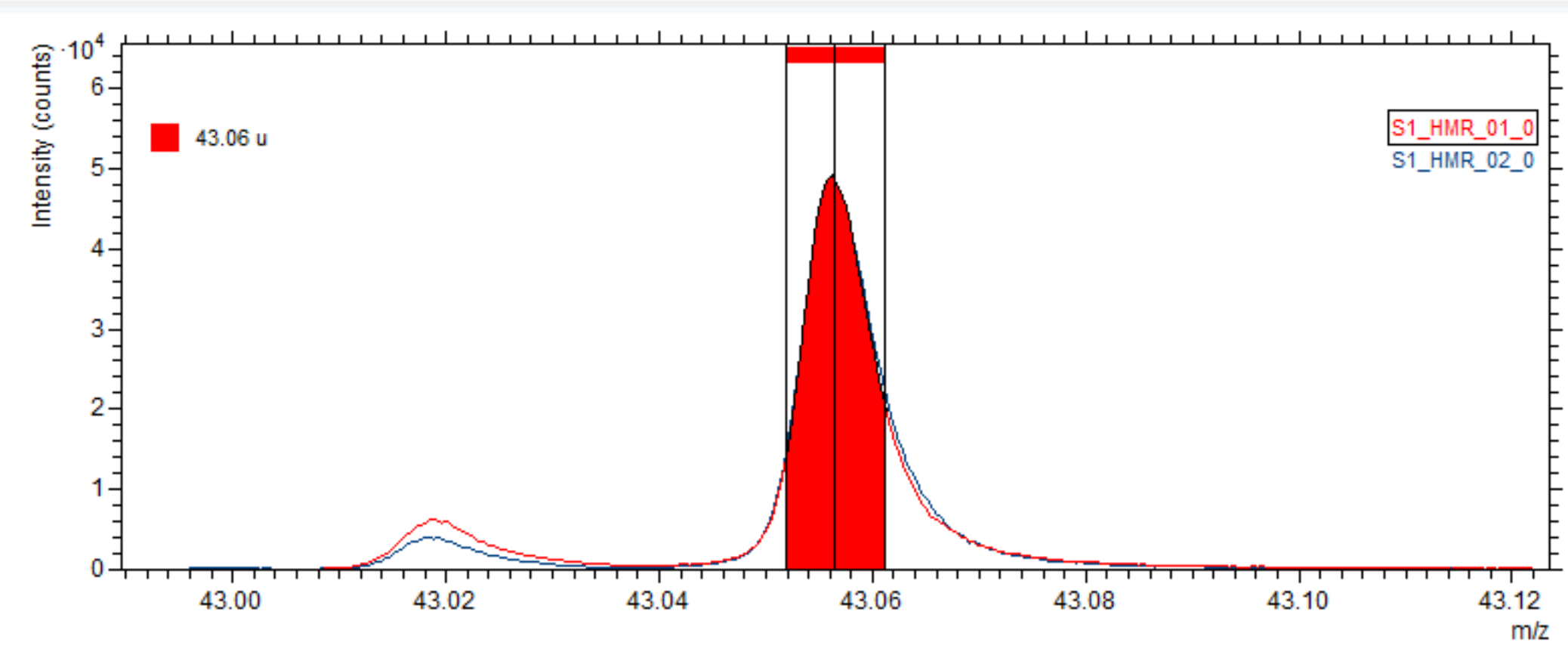
Explained (%)

- S1\_HMR\_01\_0 \*
  - <No Sample Name> (S1\_HMR\_01) \*
  - S1\_HMR\_02\_0
  - <No Sample Name> (S1\_HMR\_02)

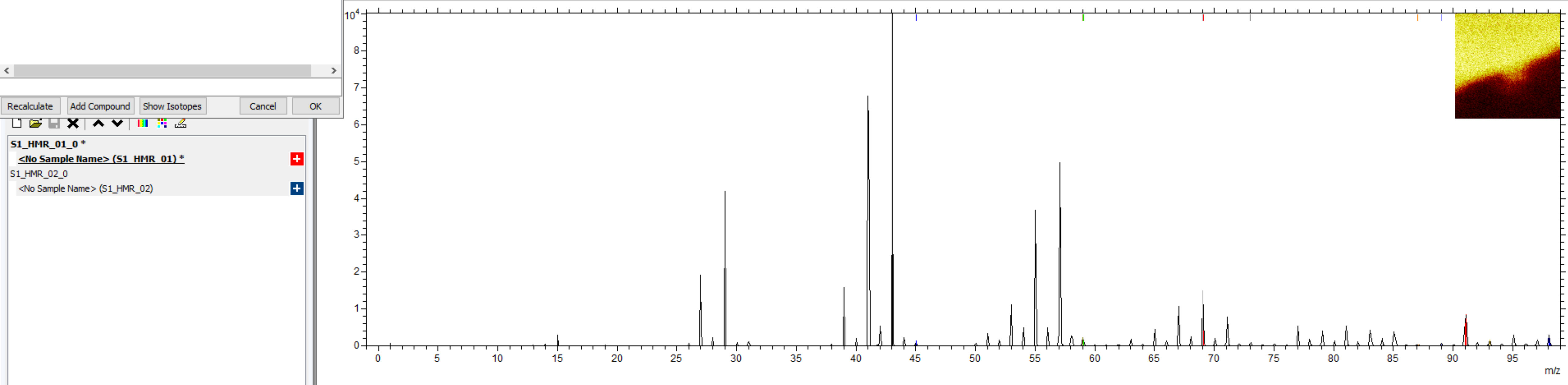


V	No.	m / z	Area / cts	Color	Peak Label	Description
<b>Mass Intervals</b>						
<input checked="" type="checkbox"/>	37	43.0565	854832	Red	43.06 u	C3H7
<input checked="" type="checkbox"/>	2	45.0352	32713	Blue	45.03 u	PEG
<input checked="" type="checkbox"/>	3	59.0136	51836	Olive	59.01 u	PMMA
<input checked="" type="checkbox"/>	4	59.0511	37293	Green	59.05 u	PEG
<input checked="" type="checkbox"/>	5	69.0376	118182	Grey	69.04 u	PMMA
<input checked="" type="checkbox"/>	6	69.0762	360824	Red	69.08 u	
<input checked="" type="checkbox"/>	7					
<input checked="" type="checkbox"/>	9					
<input checked="" type="checkbox"/>	10					
<input checked="" type="checkbox"/>	11					
<input checked="" type="checkbox"/>	12					
<input checked="" type="checkbox"/>	13					

To do this, press the Reorder IDs button found near the bottom of this toolbar. It may be off screen depending on the size of the lower windows.



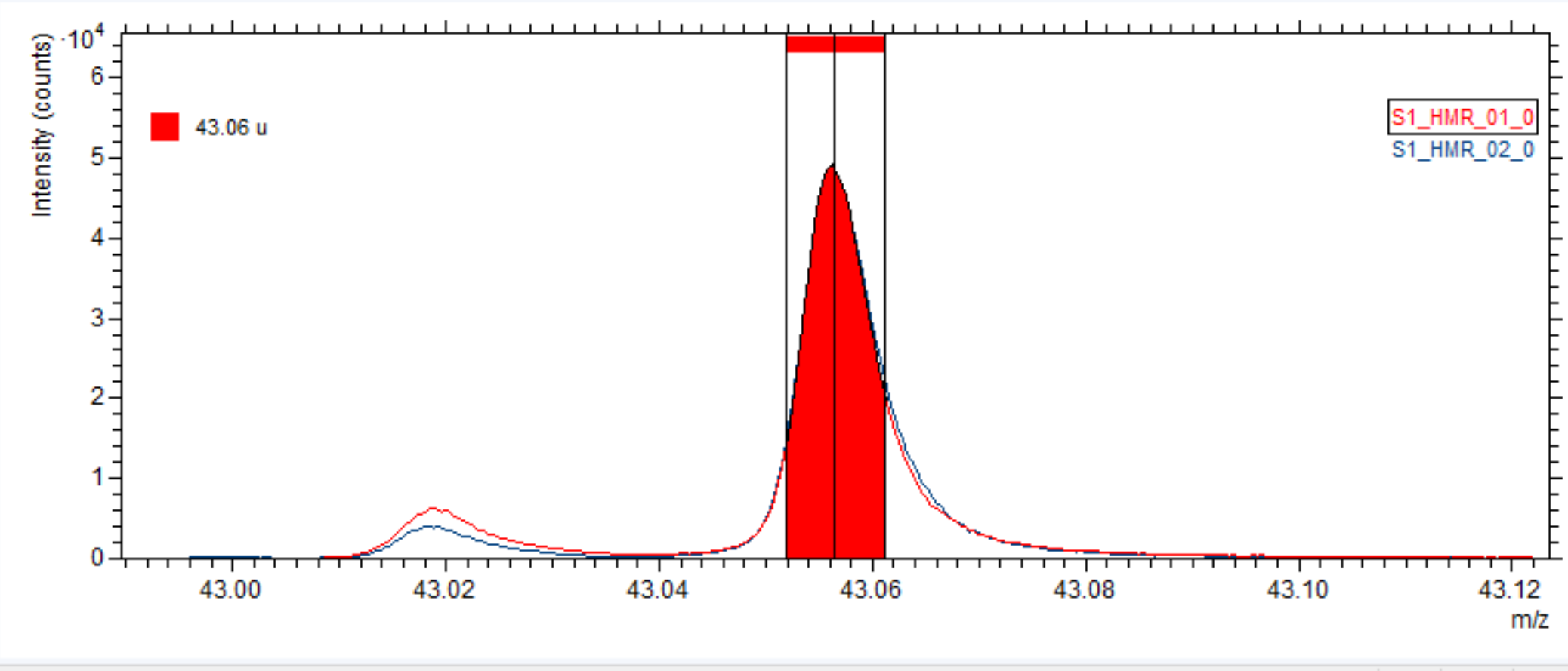
m/z  Area (cts)  Explained (%)  Resolution   
 Dev. (ppm)  Counts / Shot  Peak Difference...  Width (ns)



- S1\_HMR\_01\_0 \*  
 <No Sample Name> (S1\_HMR\_01) \*  
 S1\_HMR\_02\_0  
 <No Sample Name> (S1\_HMR\_02)

V	No.	m/z	Area / cts	Color	Peak Label	Description
<b>Mass Intervals</b>						
<input checked="" type="checkbox"/>	2	43.0565				
<input checked="" type="checkbox"/>	3	45.0352	327			
<input checked="" type="checkbox"/>	4	59.0136	518			
<input checked="" type="checkbox"/>	5	59.0511	372			
<input checked="" type="checkbox"/>	6	69.0376	118			
<input checked="" type="checkbox"/>	7	69.0762	360			
<input checked="" type="checkbox"/>	8	73.0342	164			
<input checked="" type="checkbox"/>	9	87.0478	61			
<input checked="" type="checkbox"/>	10	89.0617	3870		89.06 u	PEG
<input checked="" type="checkbox"/>	11	91.0566	238681		91.05 u	PS
<input checked="" type="checkbox"/>	12	93.0730	55170		93.07 u	PMMA
<input checked="" type="checkbox"/>	13	98.0763	70210		98.07 u	GTP
<input checked="" type="checkbox"/>	14	101.0605	26893		101.06 u	PMMA
<input checked="" type="checkbox"/>	15	103.0490	34262		103.05 u	PEG

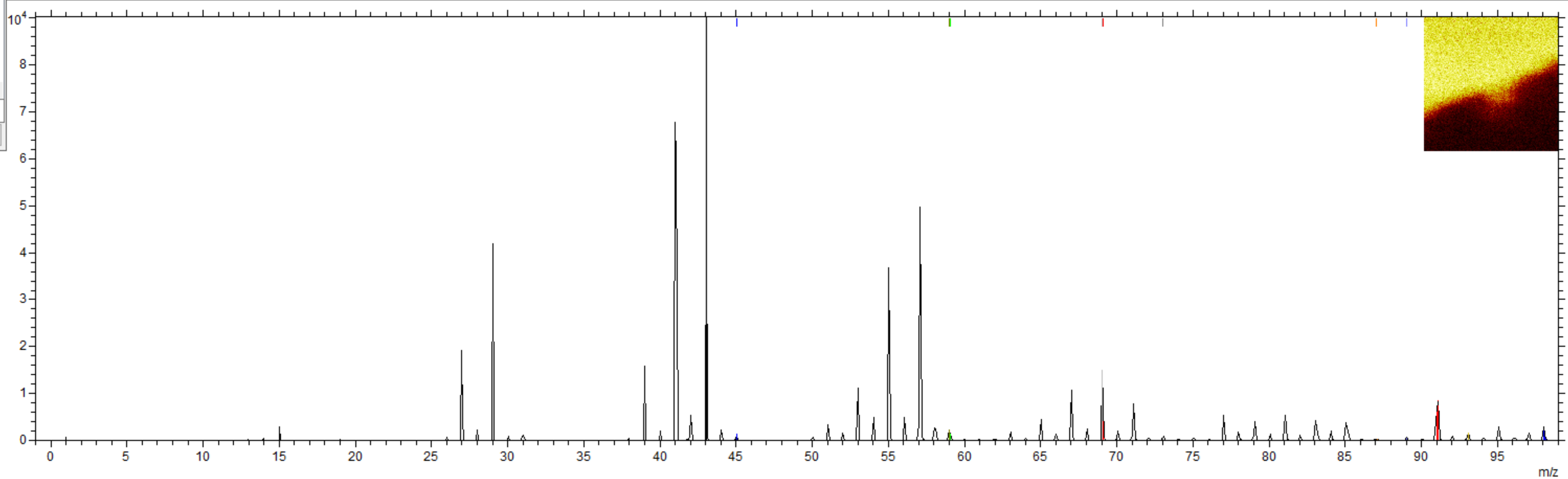
Now the peaks are ordered sequentially by mass and number. The list always starts at 2 because the Total counts is labelled as 1.



m/z  Area (cts)  Explained (%)  Resolution   
 Dev. (ppm)  Counts / Shot  Peak Difference...  Width (ns)

Assignments m/z Dev. (ppm) Explained (%) Library Occurrence

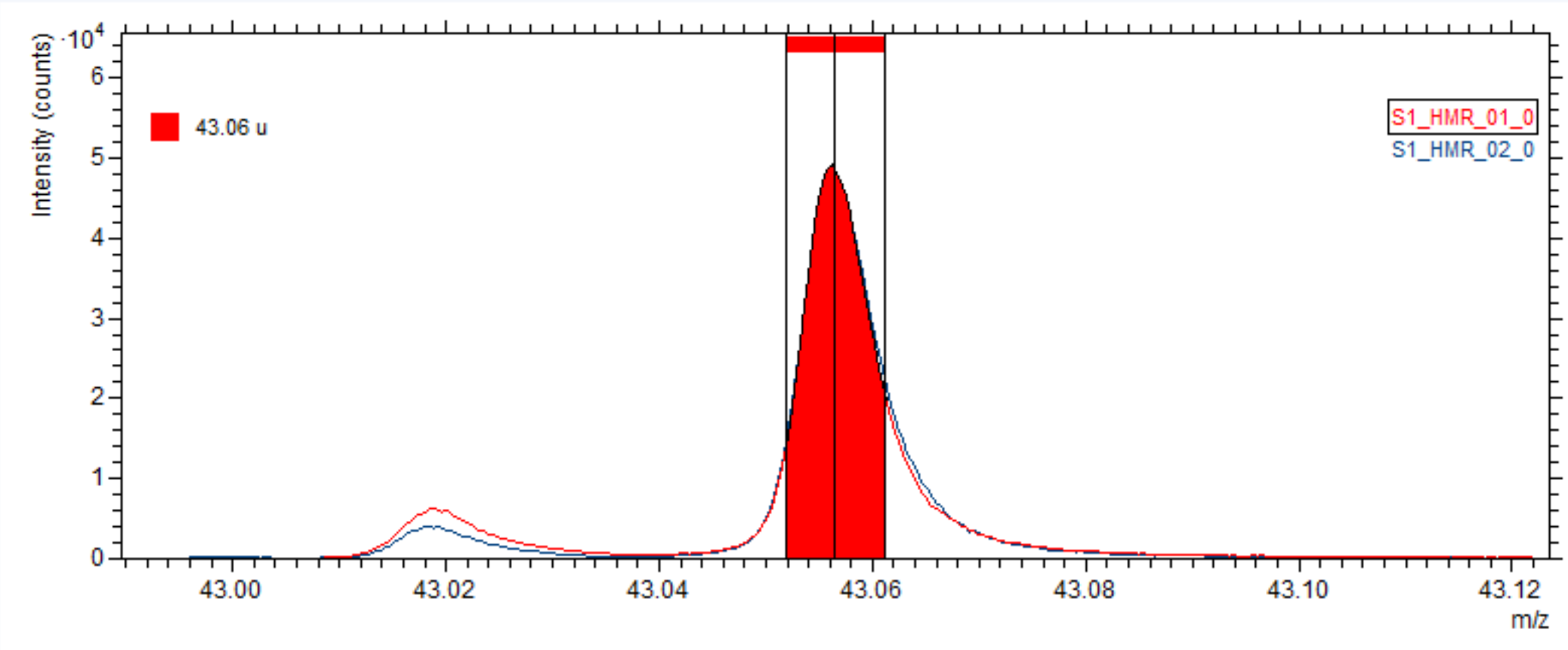
- S1\_HMR\_01\_0 \*  
 <No Sample Name> (S1\_HMR\_01) \*  
 S1\_HMR\_02\_0  
 <No Sample Name> (S1\_HMR\_02)



MassList

No	Mass	Area	Color	Formula	Library
12	93.0730	55170	Yellow	93.07 u	PMMA
13	98.0763	70210	Blue	98.07 u	GTP
14	101.0605	26893	Green	101.06 u	PMMA
15	103.0490	34262	Light Green	103.05 u	PEG

Once you have selected all the peaks you want, right click on this tab...

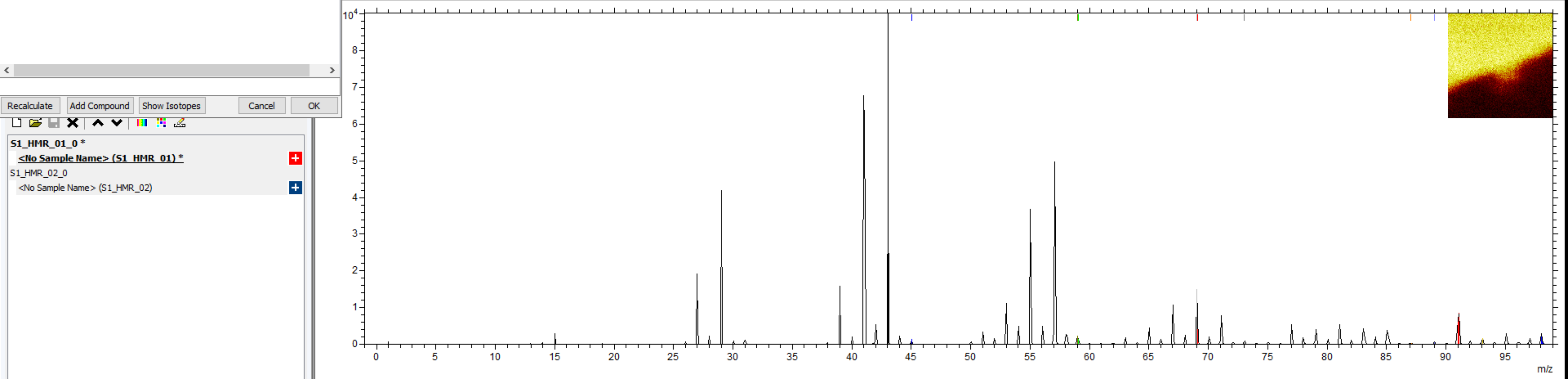


File Edit Spectrum Mass Interval List Peak List View Help

Total Range Total Area All Intervals Find: C3H3

m/z Dev. (ppm) Area (cts) Counts / Shot Explained (%) Resolution Width (ns)

Assignments m/z Dev. (ppm) Explained (%) Library Occurrence



Recalculate Add Compound Show Isotopes Cancel OK

- S1\_HMR\_01\_0 \*
  - <No Sample Name> (S1\_HMR\_01) \* +
  - S1\_HMR\_02\_0
  - <No Sample Name> (S1\_HMR\_02) +

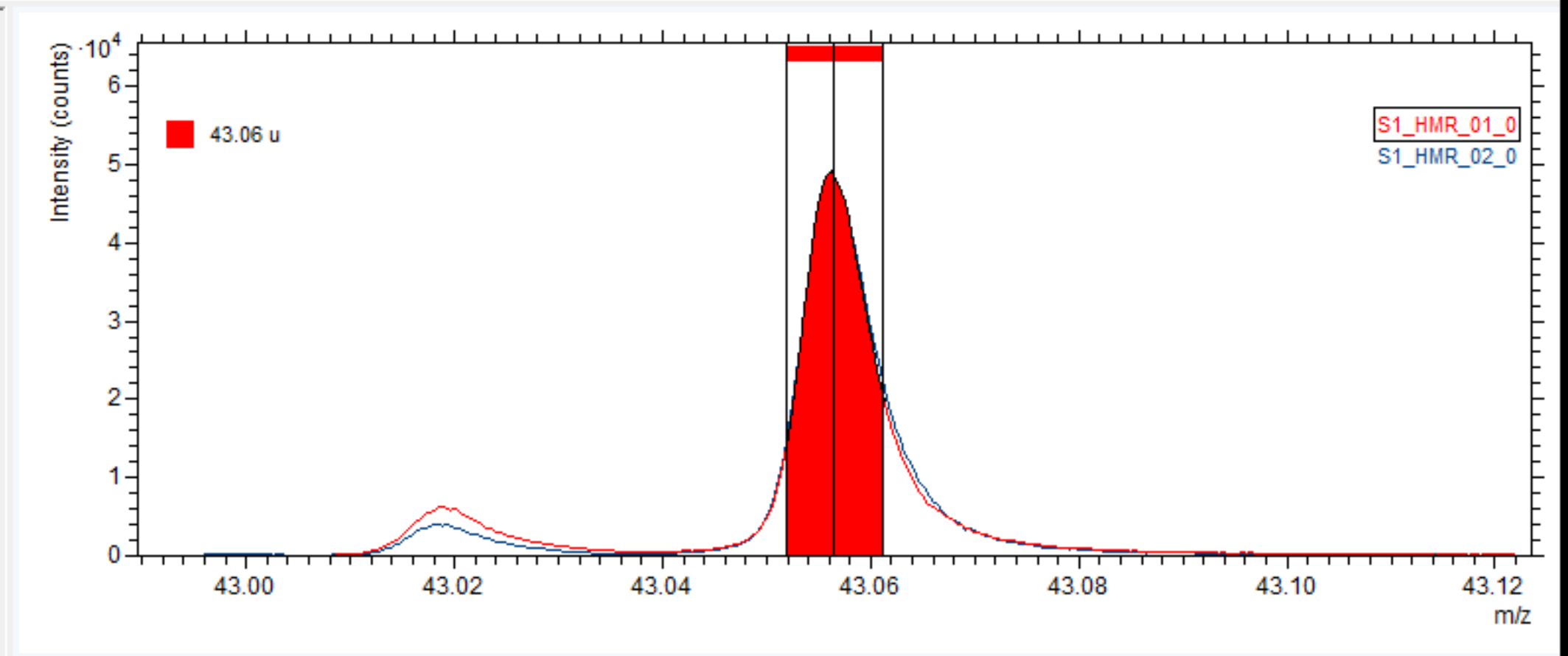
MassListPos (+)

V	No	Mass	Area	Color	Formula	Library
12	93.0730	55170		Yellow	93.07 u	PMMA
13	98.0763	70210		Blue	98.07 u	GTP
14	101.0605	26893		Green	101.06 u	PMMA
15	103.0490	34262		Light Green	103.05 u	PEG

Context menu options: Save As Mass Interval List..., Rename..., Add To, Replace All Existing Peak Lists Of, Copy to Clipboard (Ctrl+C), Copy to Edit, Delete!, Tools, Color..., Properties...

...and select 'Save As Mass Interval List...'

← →



File Edit Spectrum Mass Interval List Peak List View Help

Save As

This PC > Dr Dan Data (E:) > TempSurfaceLabTutorials

Search TempSurfaceLabTutori...

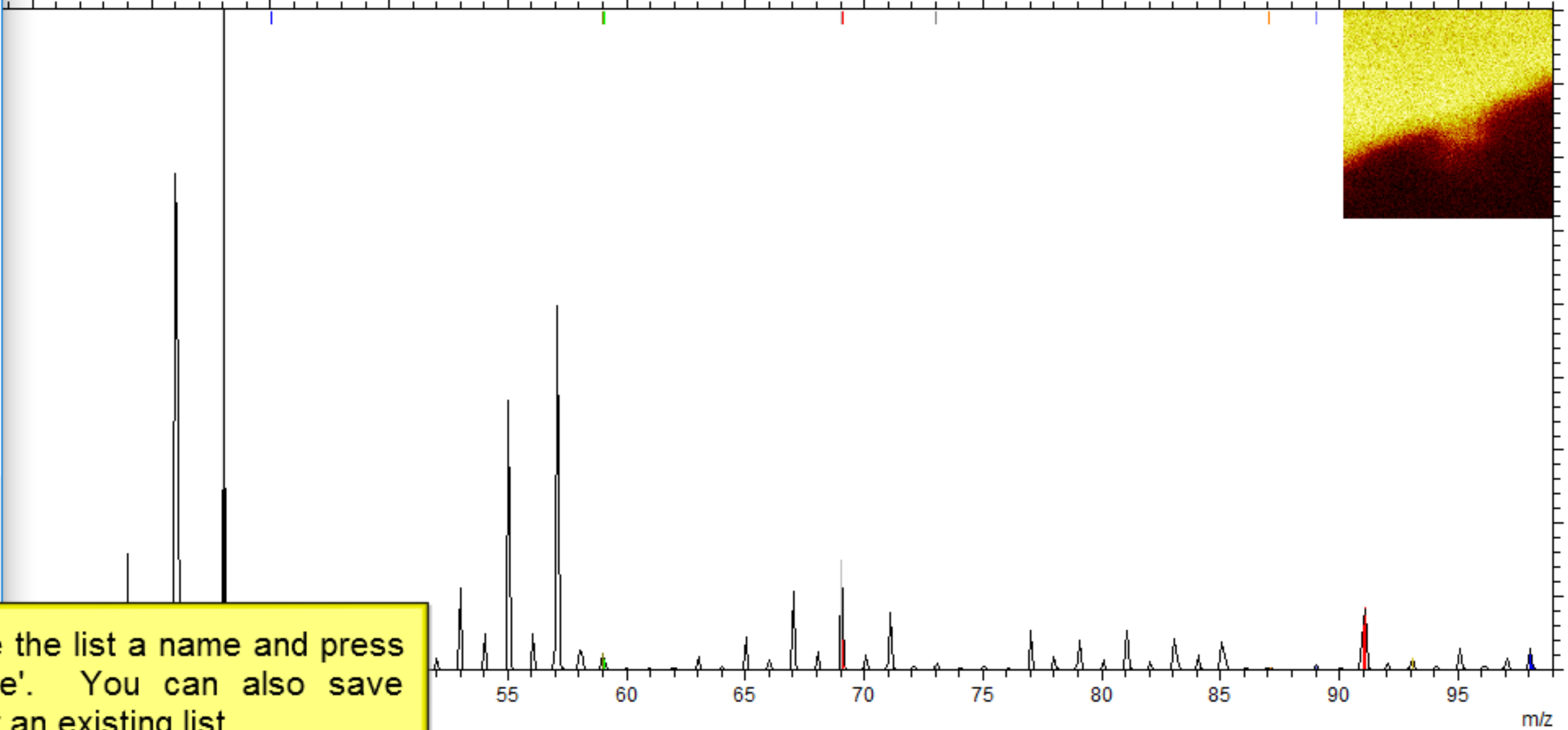
Name	Date modified	Type	Size
MassListPos.itmil	9/23/2021 3:50 PM	ITMIL File	83 KB

File name: MassListPos.itmil

Save as type: TOF-SIMS Mass Interval List Files (\*.itmil)

Save Cancel

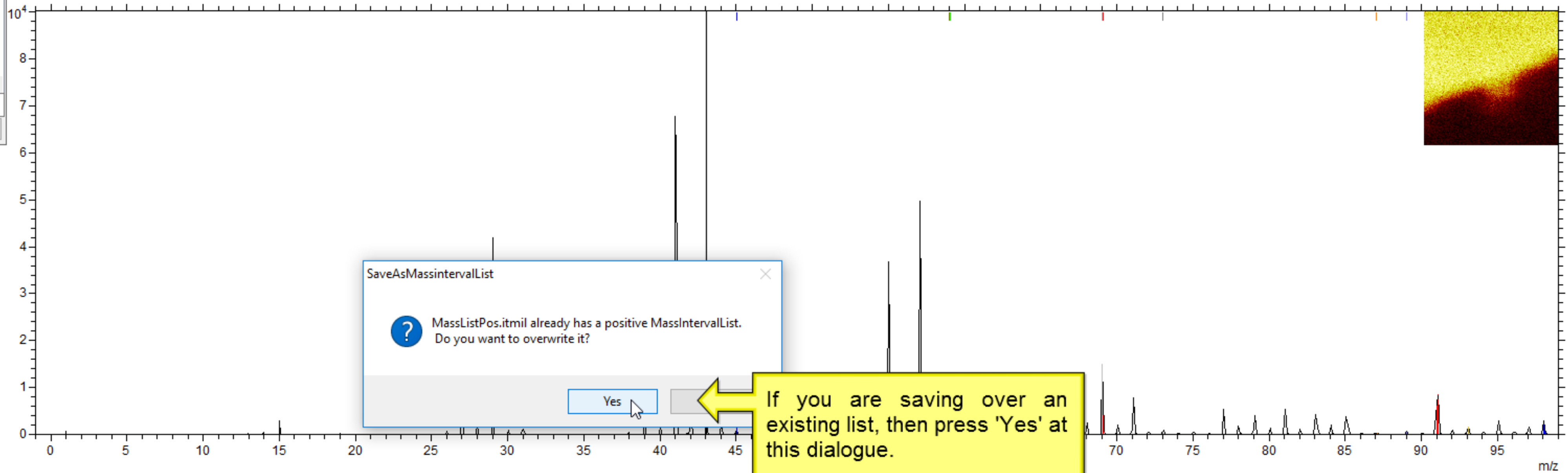
Give the list a name and press 'Save'. You can also save over an existing list.



V	No.	m/z	Area / cts	Color	Peak Label	Description
<b>Mass Intervals</b>						
<input checked="" type="checkbox"/>	2	43.0565	854832	Red	43.06 u	C3H7
<input checked="" type="checkbox"/>	3	45.0352	32713	Blue	45.03 u	PEG
<input checked="" type="checkbox"/>	4	59.0136	51836	Olive	59.01 u	PMMA
<input checked="" type="checkbox"/>	5	59.0511	37293	Green	59.05 u	PEG
<input checked="" type="checkbox"/>	6	69.0376	118182	Grey	69.04 u	PMMA
<input checked="" type="checkbox"/>	7	69.0762	360824	Red	69.08 u	
<input checked="" type="checkbox"/>	8	73.0342	16483	Grey	73.03 u	PEG
<input checked="" type="checkbox"/>	9	87.0478	6134	Orange	87.05 u	PEG
<input checked="" type="checkbox"/>	10	89.0617	3870	Blue	89.06 u	PEG
<input checked="" type="checkbox"/>	11	91.0566	238681	Red	91.05 u	PS
<input checked="" type="checkbox"/>	12	93.0730	55170	Yellow	93.07 u	PMMA
<input checked="" type="checkbox"/>	13	98.0763	70210	Blue	98.07 u	GTP
<input checked="" type="checkbox"/>	14	101.0605	26893	Green	101.06 u	PMMA
<input checked="" type="checkbox"/>	15	103.0490	34262	Light Green	103.05 u	PEG

m/z  Area (cts)  Explained (%)  Resolution   
 Dev. (ppm)  Counts / Shot  Peak Difference...  Width (ns)

- S1\_HMR\_01\_0 \*
  - <No Sample Name> (S1\_HMR\_01) \*
  - S1\_HMR\_02\_0
  - <No Sample Name> (S1\_HMR\_02)

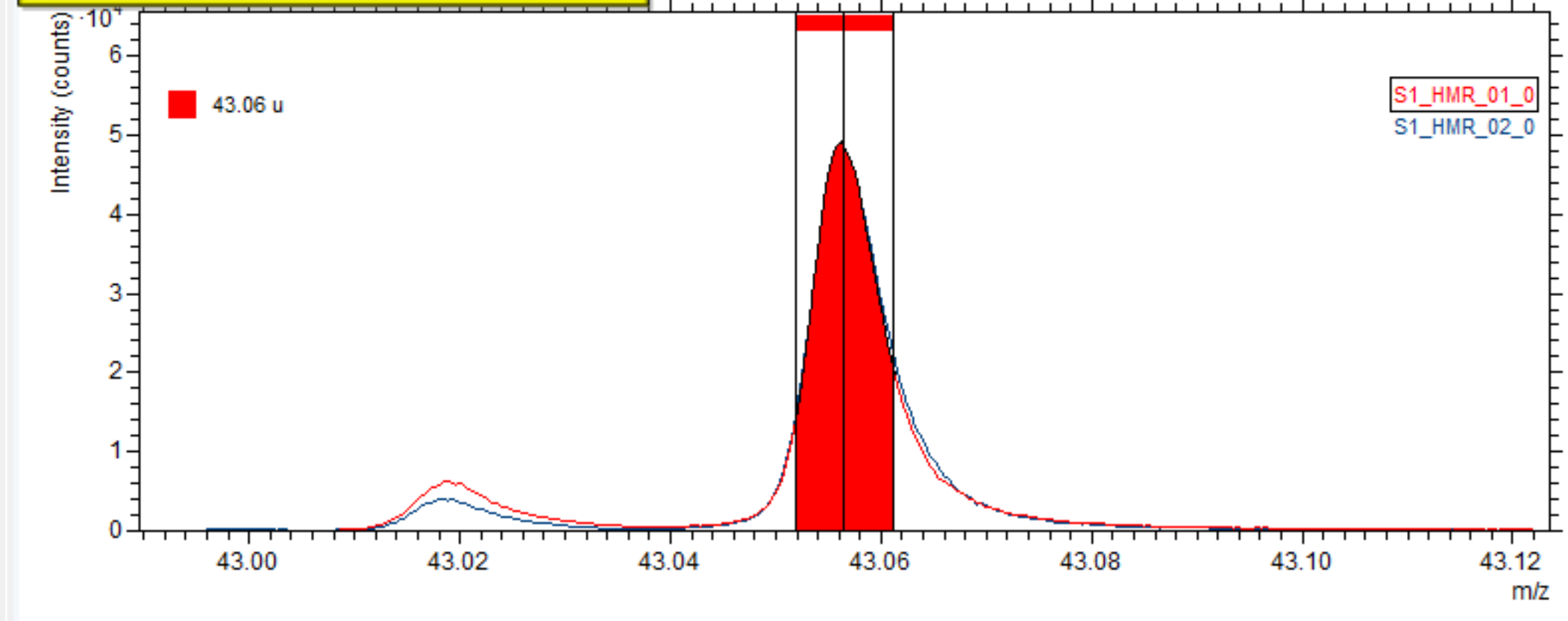


SaveAsMassintervalList

MassListPos.itmil already has a positive MassIntervalList. Do you want to overwrite it?

If you are saving over an existing list, then press 'Yes' at this dialogue.

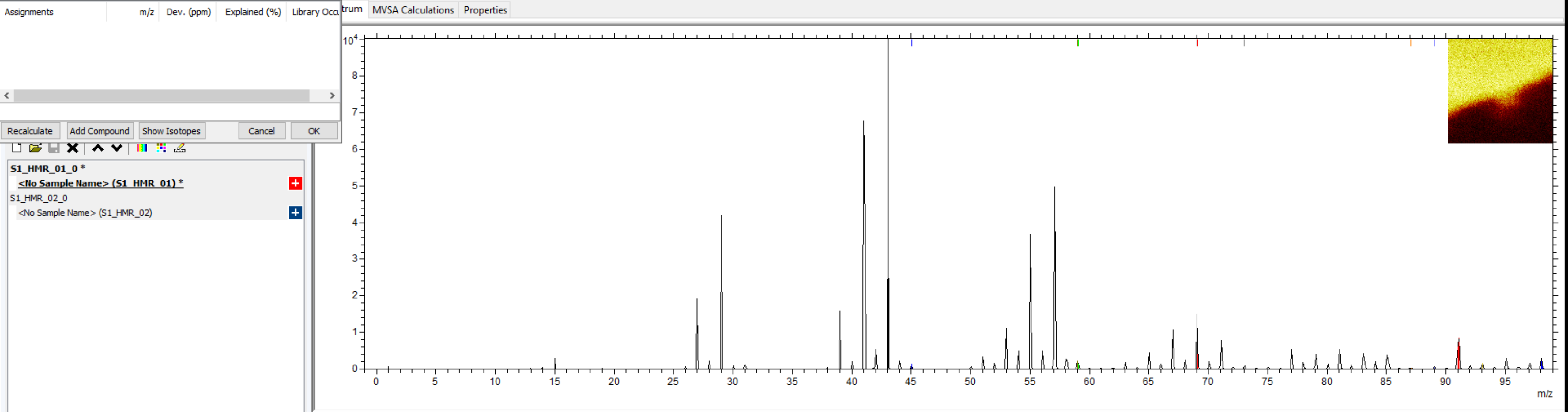
V	No.	m/z	Area / cts	Color	Peak Label	Description
<b>Mass Intervals</b>						
<input checked="" type="checkbox"/>	2	43.0565	854832	Red	43.06 u	C3H7
<input checked="" type="checkbox"/>	3	45.0352	32713	Blue	45.03 u	PEG
<input checked="" type="checkbox"/>	4	59.0136	51836	Olive	59.01 u	PMMA
<input checked="" type="checkbox"/>	5	59.0511	37293	Green	59.05 u	PEG
<input checked="" type="checkbox"/>	6	69.0376	118182	Grey	69.04 u	PMMA
<input checked="" type="checkbox"/>	7	69.0762	360824	Red	69.08 u	
<input checked="" type="checkbox"/>	8	73.0342	16483	Grey	73.03 u	PEG
<input checked="" type="checkbox"/>	9	87.0478	6134	Orange	87.05 u	PEG
<input checked="" type="checkbox"/>	10	89.0617	3870	Blue	89.06 u	PEG
<input checked="" type="checkbox"/>	11	91.0566	238681	Red	91.05 u	PS
<input checked="" type="checkbox"/>	12	93.0730	55170	Yellow	93.07 u	PMMA
<input checked="" type="checkbox"/>	13	98.0763	70210	Blue	98.07 u	GTP
<input checked="" type="checkbox"/>	14	101.0605	26893	Green	101.06 u	PMMA
<input checked="" type="checkbox"/>	15	103.0490	34262	Light Green	103.05 u	PEG



File Edit Spectrum Mass Interval List Peak List View Help

Total Range Total Area All Intervals Find: C3H3

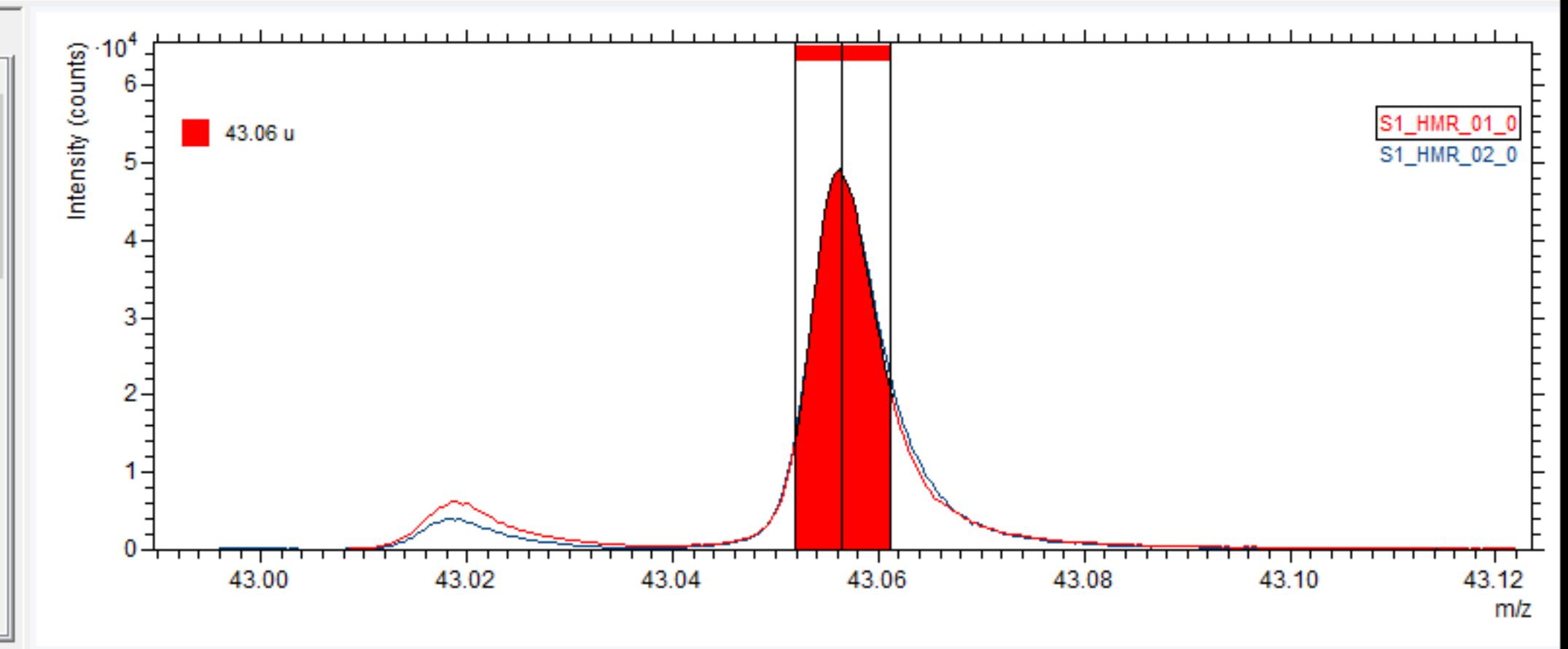
m/z Dev. (ppm) Area (cts) Counts / Shot Explained (%) Resolution Width (ns)



Recalculate Add Compound Show Isotopes Cancel OK

S1\_HMR\_01\_0 \*  
 <No Sample Name> (S1\_HMR\_01) \*  
 S1\_HMR\_02\_0  
 <No Sample Name> (S1\_HMR\_02)

V	No.	m/z	Area / cts	Color	Peak Label	Description
<b>Mass Intervals</b>						
<input checked="" type="checkbox"/>	2	43.0565	854832	Red	43.06 u	C3H7
<input checked="" type="checkbox"/>	3	45.0352	32713	Blue	45.03 u	PEG
<input checked="" type="checkbox"/>	4	59.0136	51836	Olive	59.01 u	PMMA
<input checked="" type="checkbox"/>	5	59.0511	37293	Green	59.05 u	PEG
<input checked="" type="checkbox"/>	6	69.0376	118182	Grey	69.04 u	PMMA
<input checked="" type="checkbox"/>	7	69.0762	360824	Red	69.08 u	
<input checked="" type="checkbox"/>	8	73.0342	16483	Grey	73.03 u	PEG
<input checked="" type="checkbox"/>	9	87.0478				
<input checked="" type="checkbox"/>	10	89.061				



Mass Interval Lists

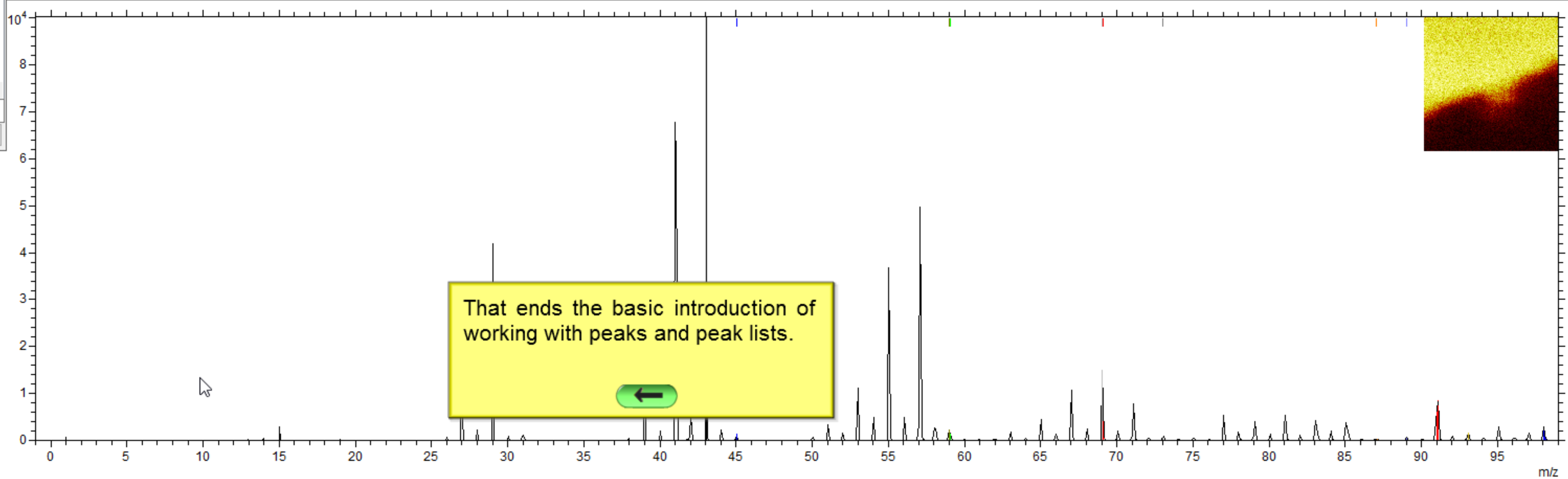
MassListPos (+)

- Add To
- Replace All Existing Peak Lists Of
- Save As...
- Color...
- Remove!

To make sure all the data is ordered properly in all of the programs, right click on the list name here and select 'Replace All Existing Peak Lists Of' -> 'Selected Spectra'.

m/z  Area (cts)  Explained (%)  Resolution   
 Dev. (ppm)  Counts / Shot  Peak Difference...  Width (ns)

- S1\_HMR\_01\_0 \*  
 <No Sample Name> (S1\_HMR\_01) \*  
 S1\_HMR\_02\_0  
 <No Sample Name> (S1\_HMR\_02)



V	No.	m / z	Area / cts	Color	Peak Label	Description
<b>Mass Intervals</b>						
<input checked="" type="checkbox"/>	37	43.0565	854832	Red	43.06 u	C3H7
<input checked="" type="checkbox"/>	2	45.0352	32713	Blue	45.03 u	PEG
<input checked="" type="checkbox"/>	3	59.0136	51836	Olive	59.01 u	PMMA
<input checked="" type="checkbox"/>	4	59.0511	37293	Green	59.05 u	PEG
<input checked="" type="checkbox"/>	5	69.0376	118182	Grey	69.04 u	PMMA
<input checked="" type="checkbox"/>	6	69.0762	360824	Red	69.08 u	
<input checked="" type="checkbox"/>	7	73.0342	16483	Grey	73.03 u	PEG
<input checked="" type="checkbox"/>	8	87.0478	6134	Orange	87.05 u	PEG
<input checked="" type="checkbox"/>	9	89.0617	3870	Blue	89.06 u	PEG
<input checked="" type="checkbox"/>	10	91.0566	238681	Red	91.05 u	PS
<input checked="" type="checkbox"/>	11	93.0730	55170	Yellow	93.07 u	PMMA
<input checked="" type="checkbox"/>	12	98.0763	70210	Blue	98.07 u	GTP
<input checked="" type="checkbox"/>	13	101.0605	26893	Green	101.06 u	PMMA
<input checked="" type="checkbox"/>	14	103.0490	34262	Light Green	103.05 u	PEG

