

## NEW GC IMAGE FEATURE: SIDE-BY-SIDE COMPARISON

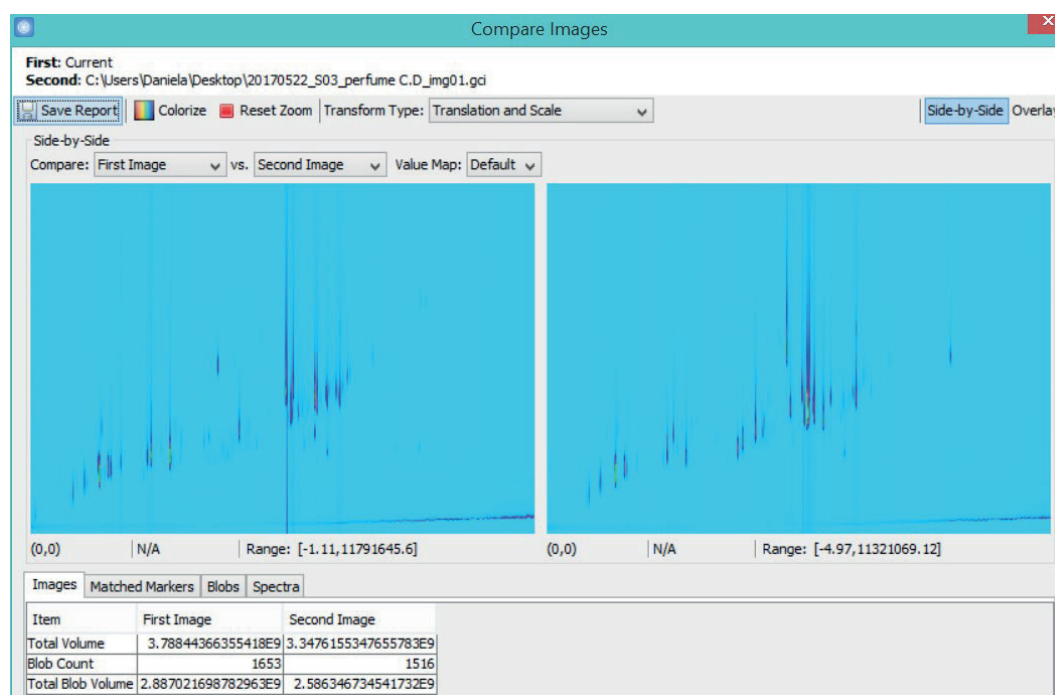
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A new version of the GC Image software package released this year, **GC Image v2.8**, advances software capability and user-friendliness with improvements and new features. An example of newly-added functionality is the **Compare Images - Side-by-Side** tool. This new comparison interface displays two chromatograms next to each other for easy assessment of similarities and differences in the 2D patterns and allows interactive review of blob matching between the two images.

### Brief overview of added possibilities

The improved Compare Images tool now gives a wider selection of comparison modes. Upon loading the reference image, this is now automatically displayed next to the current image. The user can still create differential images (difference, flicker, etc.) like in the previous software versions, and in addition a transparent overlay of the two images, to show the smallest pixel by pixel differences.

**Fig. 1**  
Compare Images - Side-by-side window.



In addition to these graphic options, the Side-by-Side functionality allows peak-based comparison using both chromatographic and spectral information. Initially each chromatogram is processed individually to detect, identify, and quantify blobs. User-identified peaks are used as alignment markers to match all blobs in the two images. The side-by-side window allows reviewing the chromatograms, enables comprehensive peak matching and automates difference detection.

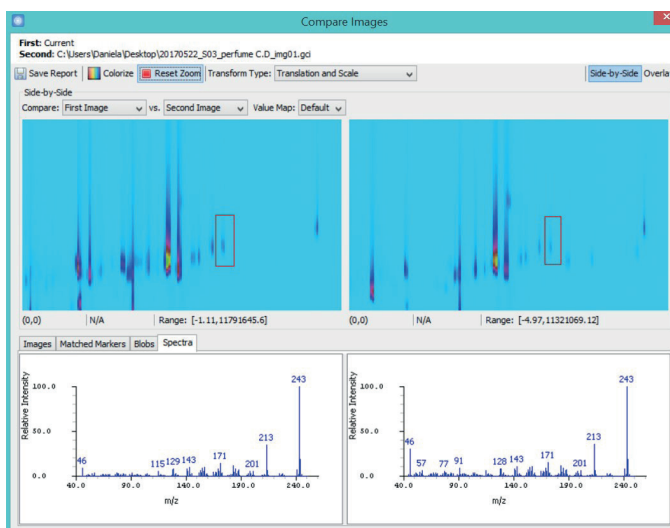
A set of tables provides quantitative comparison statistics:

- The **Matched Markers Tab** reports the marker blobs used to determine the alignment.
- The **Blobs Tab** shows blob tables along with properties and matching scores. A link is shown between the selected matched blobs in the two images to signify correlation. A colorization system gives an overview of the blob status: white blobs are matched bi-directionally, orange indicates a match with conflicting identification and red unmatched blobs unique for one image
- The **Spectra Tab** displays side-by-side the spectra for any selected blobs of the two images.

Fig. 2  
Side-by-side Blobs Tab.

First Image										Second Image										
Compound...	Retention I	Retention II	...	Status	ZD Score	Base Peak Score	Spectral Score	Match		Compound...	Retention I	Retention II	...	Status	ZD Score	Base Peak Score	Spectral Score	Match		
1	60.3	2.79	...	3Two-Way	0.54	...	1	890		7	Isouegenyl...	57.1	3.70	...	None	...	0	927		
2	60.3	2.96	...	3One-Way	0.52	...	1	938		8	Amyl cinn...	61.8	2.88	...	14Two-Way	0.538	...	1	927	
3	25.7	1.45	...	21Two-Way	0.24	...	1	919		9	...	62.9	2.44	...	73Two-Way	0.537	...	1	891	
4	21.2	1.18	...	2Two-Way	0.24	...	1	940		10	beta-Caryo...	46.7	1.72	...	Two-Way	0.531	...	0	790	
5	36.1	1.53	...	23Two-Way	0.24	...	1	935		11	...	64.6	2.25	...	Two-Way	0.526	...	0	642	
6	70.8	2.86	...	4Two-Way	0.54	...	1	938		12	...	69.1	1.64	...	9Two-Way	0.536	...	1	964	
7	32.1	1.55	...	27Two-Way	0.54	...	1	905		13	Ebanol 2	47.8	1.87	...	Two-Way	0.365	...	0	851	
8	66.3	2.73	...	28Two-Way	0.54	...	1	900		14	gamma-Ter...	23.1	1.32	...	13Two-Way	0.238	...	0	964	
9	69.1	1.64	...	12Two-Way	0.54	...	1	964		15	alpha-Isom...	50.5	2.06	...	19Two-Way	0.534	...	1	945	
10	31.2	1.55	...	49Two-Way	0.24	...	1	921		16	...	55.3	3.99	...	37Two-Way	0.530	...	1	946	
11	61.1	2.37	...	1Two-Way	0.75	...	1	922		17	...	33.3	1.99	...	Two-Way	0.532	...	1	771	
12	71.4	2.71	...	19Two-Way	0.75	...	1	958		18	...	57.9	2.50	...	None	...	0	958		
13	23.1	1.32	...	1Two-Way	0.24	...	0	964		19	...	71.4	2.71	...	12Two-Way	0.537	...	1	958	
14	61.7	2.90	...	8Two-Way	0.54	...	1	927		20	beta-Pinene	18.2	1.03	...	17Two-Way	0.238	...	1	960	
15	23.8	1.28	...	26Two-Way	0.24	...	1	937		21	Linalool	25.7	1.45	...	3Two-Way	0.238	...	1	919	
16	66.9	2.60	...	28One-Way	0.04	...	0	591		22	ISO E SUPE...	60.7	2.37	...	90Two-Way	0.530	...	1	891	
17	beta-Pinene	18.2	1.05	...	20Two-Way	0.24	...	1	960		23	Linyl acetate	53.1	1.53	...	5Two-Way	0.723	...	1	935
18	66.4	2.02	...	27Two-Way	0.54	...	0	842		24	...	62.5	2.54	...	73One-Way	0.682	...	1	872	
19	alpha-Isom...	50.5	2.04	...	15Two-Way	0.24	...	1	945		25	MUSK G 2	69.1	2.62	...	22Two-Way	0.754	...	1	926
20	112.0	0.34	...	31One-Way	0.49	...	1	958		26	...	23.8	1.28	...	15Two-Way	0.238	...	1	937	
21	46.0	3.36	...	None	...	...	0	515		27	...	32.1	1.55	...	7Two-Way	0.537	...	1	905	
22	MUSK G 2	69.1	2.62	...	25Two-Way	0.54	...	1	926		28	...	66.3	2.71	...	8Two-Way	0.538	...	1	900
23	21.0	1.36	...	2One-Way	0.03	...	0	515		29	...	47.3	1.76	...	One-Way	0.069	...	0	781	
24	111.7	0.31	...	31Two-Way	0.73	...	1	947		30	...	69.9	2.71	...	87Two-Way	0.538	...	1	847	

Fig. 3  
Side-by-side comparison with spectra.



## CONCLUSIONS

- The new Side-by-Side comparison tool makes samples comparison easier and more effective.
- Capability is extended from comparative visualization to characterization down to individual blobs.
- The user can perform direct MS comparison for peaks of interest.
- The automatic, color-based classification helps to identify recurring peaks and selective markers.

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