





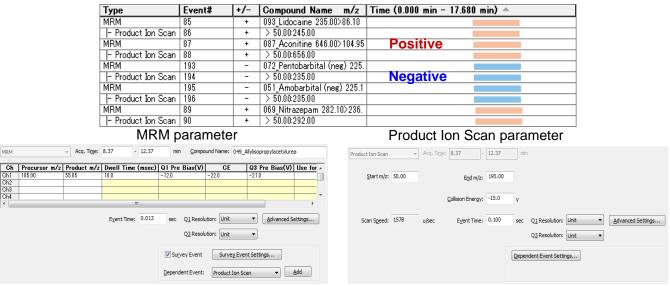
LC-MS Liquid Chromatograph Mass Spectrometer

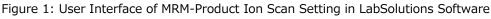
Identification of Drugs in Urine Using Synchronized Survey Scan®

LAAN-J-LM-E013

This report describes an analytical system using ultra-high-speed triple quadrupole mass spectrometry (LCMS-8030) for forensic analysis of drugs in urine. The LCMS-8030 has a Synchronized Survey Scan function that automatically performs MS/MS when a precursor threshold is exceeded, thereby producing a combined MRM and MRM-dependent product ion scan in a single analysis. Utilizing MRMs, it is possible to sensitively detect compounds while simultaneously performing a product ion scan, yielding MS/MS spectra. Furthermore, as the Collision Energy (CE) can be configured for each product scan, it is possible to optimize the CE for each compound, further increasing product ion scan spectra quality.

MRM mass chromatograms and library search results based on MS/MS spectra data from four compounds (allylisopropylacetylurea, diclofenac, amobarbital and thiamylal) spiked in urine are shown in this report.





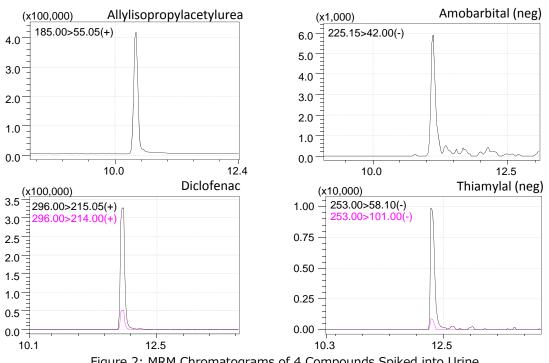


Figure 2: MRM Chromatograms of 4 Compounds Spiked into Urine

Figure 3 shows the LabSolutions Library Search Parameter Settings. Accurate results can be obtained using details, including precursor m/z and polarity, in addition to MS/MS spectral information (product m/z values and intensities).

K Method View - Library Search Parame	ters							63 View 📝 Edit	
Integration Identification Quantitative	Compound Gr	roup Performance	Spectrum	Library	(MSn)Library	/ Custom	QC Check		
☑ Search with (MSn)Library function				Adduct Ion of Target:					
Library File Name:		Min Similarity:	m/z Tolera	nce:	Positive:	H+	+1.00728	Da	
C:¥LabSolutions¥Data¥Method_Package	¥Rapid T	0	0.5		Negative:	-(H+)	-1.00728	Da	
		0	0.5						
		0	0.5						
		0	0.5						
		0	0.5	Ma	ax Hits: 5				
					Exclude comp	oounds of s	same CAS#		
Post Search: 📃 Case Sensitivity On				1	Search with S	Same Precu	ursor m/z		
Index	Parameter			▼ Use The Threshold of Intensity					
2					Rel/Abs.	Intensity:	Relative Intensity	¥ -	
3				E	Т	hreshold:	20		
5					Ignore Isotope	es of Precu	ursor Ion in a MS/	MS Spectrum	
6				E	xolude Range:	-2	- 5		
7				-	Reverse Sear	ch			
				~	Scan with sar	me polarity	,		
					ſ	Advanc	ed Settings		

Figure 3: LabSolutions Library Search Parameter Settings interface

Alternatively, it is possible to retrieve inaccurate information from a library search when precursor m/z and polarity information is excluded (Fig.4 bottom), indicating precursor m/z and polarity are important values when searching the library.

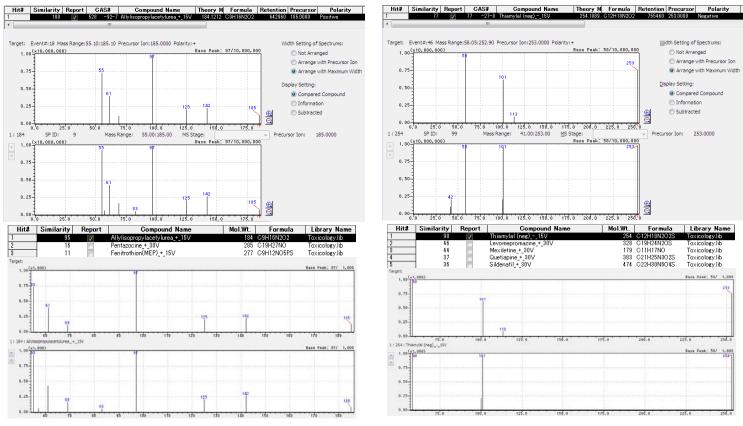


Figure 4: Library Search Results (top : parameter ON, bottom : parameter OFF)

First Edition: June, 2014



For Research Use Only. Not for use in diagnostic procedures.

The content of this publication shall not be reproduced, altered or sold for any commercial purpose without the written approval of Shimadzu. The information contained herein is provided to you "as is" without warranty of any kind including without limitation warranties as to its accuracy or completeness. Shimadzu does not assume any responsibility or liability for any damage, whether direct or indirect, relating to the use of this publication. This publication is based upon the information available to Shimadzu on or before the date of publication, and subject to change without notice.

Shimadzu Corporation www.shimadzu.com/an/