Waters[™]

Applikationsbericht

CD-R Patent Protection: Brand Characterization to Identify Counterfeit Goods using Simplified Mass Spectrometry

James Morphet, Eleanor Riches

Waters Corporation



Abstract

This application note provides an easy-to-use tool for differentiation between commercially available recordable compact discs (CD-Rs), and to identify counterfeit products.

Introduction

CD-R discs are composed of layers of polycarbonate, a burnable dye, a reflective layer, and a protective coating. The data are written using a laser beam, which follows the grooves in the polycarbonate, burning pits into a special dye that creates a code on the surface. When reading, the laser is set to a lower intensity, and the movement over the edges of the pits causes the reflection of the laser to flicker - which is transformed into a signal by a photo detector. The type of burnable dye used by manufacturers is often patented with premium disc brands using dyes and other additives that give the best performance, while being stable to atmospheric conditions, such as light and humidity. Cheaper products often use less stable dye components, which can change over time, leading to possible data degradation. This burnable dye is sometimes mixed with other dyes to change its color and appearance in order to make the disc more attractive to consumers, and to conceal the type of burnable dye used from competitors.

This competitive environment increases the manufacturers' need for reliable and reproducible testing techniques that can be used to identify differences between brands, and to detect forgeries. Ensuring the consistency of product composition is essential for maintaining product performance and quality. Many manufacturers are seeking ways to profile, or 'fingerprint' their brands by using analytical techniques capable of detecting lower quality counterfeit products that may infringe upon their patents.

A chemometric approach can be adopted for this analysis. This approach can quickly provide information about the similarities and differences within a chromatographic dataset. Automation of this process can greatly reduce the analysis time required, and the probability of error in this assessment.

Advances in ease-of-use mass spectrometer operational software have changed working practices from complex to routine, making day-to-day MS use easier for all. This application note describes the characterization of commercially available recordable compact discs using Time-of-flight Mass Spectrometry, (TOF-MS), especially designed for inexperienced users and simplified for experienced users.

Simplified Workflow

The Waters ACQUITY UPLC with Xevo QTof MS is an LC-MS/MS system that simplifies the process of acquiring and interpreting data to allow new users to feel confident when they have limited experience with TOF-MS. For the first time, Waters' IntelliStart Software has been included with this type of mass spectrometer. With its advanced and basic settings, IntelliStart allows an experienced user to perform the initial instrument setup, which can then be accessed by less practiced analysts.



Xevo QTof MS.

Figure 1 describes this calibration workflow and shows the report generated with a few simple mouse clicks. This workflow indicates if the instrument is performing within preset user-defined parameters. Calibration can be performed infrequently for an instrument situated in a well-regulated, temperature controlled laboratory. LockSpray conditions can also be defined here using a setup wizard that can run in conjunction with the calibration.

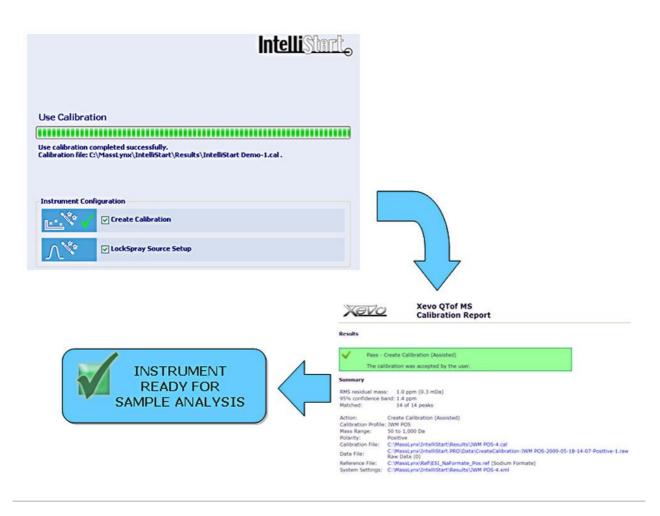


Figure 1. Workflow showing easy calibration of Xevo QTof MS

Once the calibration profile and LockSpray have been set up and saved, further checks can be performed on these files to confirm that the instrument is still within its correct operating range. This will typically be performed before an important batch of samples to ensure precise accurate mass data.

Less-experienced users can verify the calibration and lock mass properties on a more frequent basis to ensure accurate mass precision. IntelliStart Technology uses a simplified computer wizard to run a fully automated procedure. With a few easy mouse clicks it will ready the instrument for use. This check takes the profile, earlier established by an experienced user, and confirms that it still valid and appropriate to use.

Figure 2 shows a previously checked and loaded calibration profile.

	Select Calibration Profile Calibration Profile Calibration Profile IntelliStart Demo (50 - 1.000 Dis, Automatic) I
	Protove Sodium Formate: Negative: (Not available) Available for use Calibration Forlie - IntelliStat Demo Calibration Profile - IntelliStat Demo Cancel Mass Range Mass Range: Mass Range: So to 1,000 Da Calibration Type Calibration Type Calibration Type: Calibration Type: Calibration Type: Calibration Check: Calibration Type: Calibrated and Passed Calibration Check: Status: Calibrated and Passed Calibration Check: Calibrated on: June 3, 2009 10:13:45 AM Calibrated on: June 3, 2009 3:51:09 AM Calibrated on: June 3, 2009 3:51:09 AM Calibrated on: Mot Available Reference Compound O Not Available Reference Compound Defined on:
Use Calibration	
Calibration Profile: IntelliStart Demo Calibration Profile Calibration Profile	INSTRUMENT READY FOR SAMPLE ANALYSIS
Instrument Check Calibration C	

Figure 2. Workflow showing IntelliStart calibration check for inexperienced users on Xevo QTof MS.

A few mouse clicks and IntelliStart runs through its automated checks to give a clear indication that the system is performing correctly. The three green positive confirmation marks (\checkmark) indicate that the system has passed its user defined limits and it is ready to use.

Experimental

Sample Preparation

A simple sample extraction was employed. The CD-R was cut into small segments which were placed in a 100 mL bottle. 50 mL methanol was added, and the bottle was vigorously shaken for 10 min. The supernatant was filtered and placed into a Waters Certified vial, capped, and placed in the ACQUITY Sample Manager for analysis.

LC conditions

LC system:	ACQUITY UPLC
Column:	ACQUITY UPLC HSS C ₁₈ 2.1 x 50 mm, 1.8 μ m
Column temp:	40 °C
Sample temp:	5 °C
Gradient:	0.00 min 95% A
	6.00 min 0% A
	8.00 min 0% A
	8.10 min 95% A
Mobile phase: A	Water + 0.1% formic acid
Mobile phase: B	Methanol + 0.1% formic acid
Weak needle: wash	Water + 0.1% formic acid
Strong needle: wash	Methanol + 0.1% formic acid
Total runtime:	10 min

Injection volume:

5 µL, partial loop injection with needle overfill

MS conditions

MS system:	Xevo QTof MS
Ionization mode:	ESI positive
Capillary voltage:	3 kV
Sample cone:	30 V
Source Temperature:	120 °C
Desolvation gas:	Nitrogen, 800 L/Hr, 400 °C
Cone gas:	Nitrogen, 5 L/Hr
Lock mass compound:	Leucine enkephalin

The design of the Xevo QTof MS allows for three bottles to be utilized by the on-board fluidics system, as shown in Figure 6. Typically, these would be for a lock mass solution, a calibration solution, and a compound of interest that may require infusing. The infusion from these bottles is easily controlled from the instrument page, or automatically from the MS method used for the acquisition of samples, or IntelliStart Software.

A novel feature of Xevo QTof MS is its ability to run methods with a multi point lock mass. This is used to "invisibly" correct the data to give precise accurate masses during the run at the time of ionization. It uses a patented, in-source switching baffle that quickly alternates between analyte and LockSpray flows. This method eliminates the need for any post-column plumbing, or extending sample analysis times, as it uses the on-board fluidics with an automated method of delivery to the source chamber during each injection.

Up to four different points across the mass range can be chosen, giving the analyst a greater choice when looking at compounds across the mass range. Figure 3 shows leucine enkephalin with three user-defined mass values of 556.2771, 425.1825, and 278.1141, all of which will be used to give a precise accurate mass.

ock Mass —			
ock Mass Nam	ne: Lock mass editor	demo	1
Crasta a Lor	ck Mass based on a Re	eference File	1
		ne Enkephalin 🗸 🗸	
]
) Specify your	o <u>w</u> n Lock Mass		
lode ———			
) <u>м</u> s			
) M <u>S</u> MS			
/ M <u>S</u> MS			
ositive Pola	rity		
and a Set N	ng masses for Leucine i 1ass of 556.2771 Da:	Enkephalin were obtained in M	SMS mode, with 21 V Collision Energy,
Include	Mass (Da) 🗸	Relative Intensity (%)	Collision Energy
	556.2771	35	O <u>O</u> ff
	425.1825	45	On Voltage (V): 21
	397.1876	100	
~	278.1141	45	
	221.0926	10	MSMS Set Mass Set Mass (Da): 556.2771
	136.0762	10 🗸	3 <u>e</u> t Mass (Da).
egative Pola			SMS mode, with 21 V Collision Energy,
and a Set M	lass of 554.2615 Da:	Enkephalin were obtained in M.	SMS mode, with 21 V Collision Energy,
	Mass (Da) 🛧	Relative Intensity (%)	Collision Energy
Include	130.0868	15	⊙ O <u>f</u> f
Include		10 =	On Voltage (V): 21
Include	179.0821	10	
Include	179.0821 219.0770	10	
Include /		70	
Include Include	219.0770		MSMS Set Mass Set Mass (Da): 554.2615

Figure 3. The lock mass editor is simple and easy-to-use.

Results and Discussion

The samples were analyzed using ACQUITY UPLC with Xevo QTof MS to show the differences between them. MarkerLynx XS Software was used to process the data into an easily understandable format, with all the calculations and comparisons performed automatically by the software.

Data acquisition and processing methods

Data were acquired using Waters' MassLynx Software, v.4.1. Incorporated into MassLynx, IntelliStart Technology automates calibration and lock mass checks. Its real-time monitoring also gives the user confidence that the system is running at optimum conditions.

The data were processed using MarkerLynx XS Application Manager. This software enables users to overcome the time-consuming problem of identifying patterns in LC-MS data sets, by showing the similarities and differences of the detected MS traces. These characteristics define the markers within the sample set with a "mass-retention time pair". The chromatograms in Figure 4 show the data obtained from two CD-R brands. It would be difficult to characterize all of these differences without MarkerLynx XS, as they look quite similar upon inspection with the naked eye.

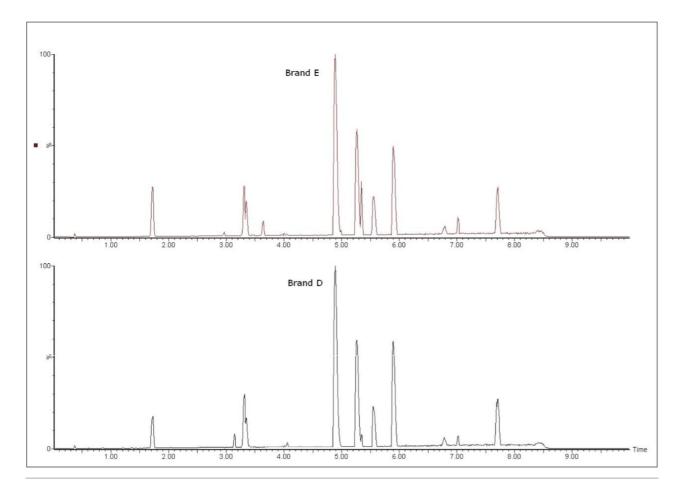


Figure 4. The chromatograms for two different brands are very similar in nature.

These data can then be used for further experiments using the MS/MS capabilities of the instrument to help identify structural information¹.

Principal Component Analysis (PCA) is part of the extended statistics that comes as standard with MarkerLynx XS Application Manager, as shown in Figure 5. This mathematical tool will plot complex data by grouping samples that show similar characteristics (having similar markers). This highlights samples with a similar composition in a matter of seconds.

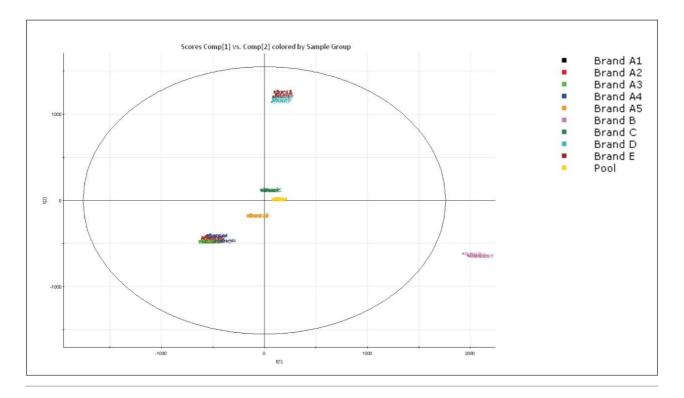


Figure 5. PCA analysis instantly shows differences between sample sets.

The six replicates of ten samples were injected in random order to remove bias. The pool sample (in yellow) was made up of a mixture of equal aliquots of all of the extracted CD-Rs. This is used to show the experiment is valid because this pool should contain properties from all samples and should therefore be found close to the center of the plot. If the pool sample does not appear near the middle, then it is likely that the experiment has not succeeded.

The PCA also illustrates that Brands A1 to A4 (from same manufacturer) all share the same characteristics, except for differences in disc color, showing that separation is not due to this. This shows that the manufacturer is adding extra dyes to their products to disguise the type of dye layer used in production. Three different types of dye layer are generally used and this will be detailed further in Part 2 of this application note¹.

Brand A5 was made by the same manufacturer but was purchased from a European geography. It shares some characteristics of the Japanese CD-Rs A1 to A4 but not all, which implies a variance in composition. The other brands were all characterized and are shown in Figure 5. Brands D and E show many similarities, which means that they share many of the same markers and that their formulations will be similar.

Conclusion

- The ACQUITY UPLC with Xevo QTof MS system was used to show brand variations between commercially available CD-Rs sold in Japan and Europe using accurate mass spectrometry.
- MarkerLynx XS Software offers scientists the unprecedented ability to visualize and interpret the most complex data automatically.
- Counterfeit goods can easily be identified because the location on a MarkerLynx XS PCA plot will clearly show them as a separate group from the genuine product.
- · Multi-point lock mass capability improves mass accuracy across the scanned range.
- This LC-MS/MS System, as shown in Figure 6 reduces laboratory running costs with:
- · IntelliStart ease-of-use Software
 - -Simplified training of new users and daily operation with its background system monitoring.
 - -Automated system setup to accurately prepare the instrument for use without human bias.
- · Robust performance

-Minimal variation over large sample batches (shown by good sample grouping) gives confidence in system reproducibility, and reduces the need for sample retests.



Figure 6. ACQUITY UPLC with Xevo QToF MS.

References

1. Morphet, J, Tatsuya, E. CD-R Patent Protection: Competitor Analysis Made Easy. Waters' Application Note. No. 720003161en, July 2009.

Featured Products

ACQUITY UPLC System <https://www.waters.com/514207> LockSpray Exact Mass Ionization Source <https://www.waters.com/1000396> Progenesis QI Software <https://www.waters.com/134790655> MassLynx MS Software <https://www.waters.com/513662>

720003160, July 2009

©2019 Waters Corporation. All Rights Reserved.