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Application Note

Analysis of Genotoxic Impurities by an ACQUITY QDa Mass Detector with Integrated ACQUITY Diverter Valve

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This is an Application Brief and does not contain a detailed Experimental section.

Abstract

This Application brief demonstrates the use of an ACQUITY QDa Mass Detector integrated with an ACQUITY Diverter Valve for the analysis of aryl esters of p-toluenesulfonic acid in monitoring of potential genotoxic impurities.

Benefits

Minimize potential sources of contamination and mitigate chances of ion suppression by diverting selected analysis segments to waste.

Introduction

Genotoxic impurities (GTIs) have the potential to react with DNA and induce genetic mutation, which may

consequently lead to cancer. It is therefore essential to have a reliable and highly sensitive method for the low level detection of these mutagenic impurities in both drug substance and drug product assays.

For low levels analysis, it is important to show that impurities can be separated and accurately measured in the presence of the sample matrix. The sample matrix can often enhance or suppress the MS signal, leading to inconsistencies in quantitation from sample-to-sample. In addition, the presence of a high concentration of active pharmaceutical ingredient (API) may contaminate the MS source, generating inaccurate results. Incorporating a diverter valve into the flow path will enable the matrix components to be selectively directed to waste, avoiding contamination of the MS source and ensuring the consistency and accuracy of the response.

In this study, we will demonstrate the use of an ACQUITY QDa Mass Detector integrated with an ACQUITY Diverter Valve for the analysis of the potentially genotoxic impurities, methyl and ethyl esters of ptoluenesulfonic acid.

Results and Discussion

The ACQUITY Diverter Valve connected to an ACQUITY QDa Mass Detector enables users to switch the mobile phase flow to waste to direct an unwanted section of the chromatographic injection away from the MS source.

For the analysis of potentially genotoxic impurities, methyl and ethyl esters of p-toluenesulfonic acid, we used the diverter valve to direct the API portion of the injection to waste. A sample solution containing 1 mg/mL of bretylium tosylate API was spiked with the esters at 0.1% and analyzed using an ACQUITY QDa Mass Detector. The MS TIC data without a diverter valve (Figure 1a) shows that API elutes before the analytes. Using a valve, we switched the flow of the mobile phase to waste at 0.15 minute to divert the API portion of the chromatographic injection away from the MS source (Figure 1b). At 2.00 minutes, we switched the flow back to the mass detector to analyze the esters. We programmed the instrument method of the ACQUITY QDa Mass Detector to automatically switch the valve to "ON / Flow to QDa" and "OFF / Flow to Waste" positions (Figure 2). For our method, the diverter valve was switched to an "OFF / Flow to Waste" position between 0.15 and 2.0 minutes to divert the flow into waste.

Instrument Method in Empower

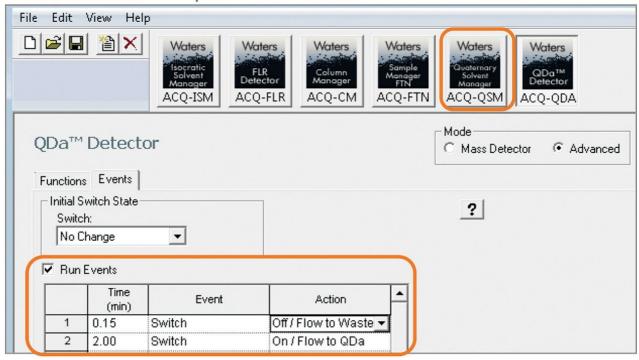


Figure 1. MS TIC (140–280 Da) data of bretylium tosylate API spiked with methyl and ethyl p-toluenesulfonate esters (MTS and ETS). Data acquired with an ACQUITY QDa Detector: A) without and B) integrated with a divert valve.

Methyl p-toluenesulfonate

System Suitability

Sample Set ID: 5702 Result Set ID: 6523 Processed Channel Descr.: 2: QDa Positive(+) SIR Ch1 204.00 Da, CV=6, Smoothed by 9 point Mean

	4	Nam	e: MTS	5		
	Name	Sample Name	RT	Area	USP Tailing	K Prime
1	MTS	SS: 1ug/mL	2.327	11508409	1.2	8.3
2	MTS	SS: 1ug/mL	2.327	11420567	1.2	8.3
3	MTS	SS: 1ug/mL	2.328	11564739	1.2	8.3
4	MTS	SS: 1ug/mL	2.328	11501751	1.2	8.3
5	MTS	SS: 1ug/mL	2.329	11470791	1.2	8.3
Mean			2.328	11493251		
Std. Dev.			0.001	52915.196		
% RSD			0.03	0.46		

Ethyl p-toluenesulfonate

	System Suitability									
NEOU OF	Sample Set ID:	5702	Result Set ID: 6523							
ubawei	Processed Chann	nel Descr.:	2: QDa Positive(+) SIR							
	Ch2 218 00 Da	CV=6 Sm	oothed by 9 point Mean							

	Nan	ne: Els)			
Name	Sample Name	RT	Area	USP Tailing	K Prime	
ETS	SS: 1ug/mL	2.606	20054524	1.2	12.0	
ETS	SS: 1ug/mL	2.609	20368670	1.2	12.0	
ETS	SS: 1ug/mL	2.610	20353450	1.2	12.1	
ETS	SS: 1ug/mL	2.610	20102186	1.2	12.1	
ETS	SS: 1ug/mL	2.611	20179382	1.2	12.1	
		2.609	20211642		l.	
		0.002	143592.202			
		0.07	0.71			
	ETS ETS ETS ETS	Name Sample Name ETS SS: 1ug/mL ETS SS: 1ug/mL ETS SS: 1ug/mL ETS SS: 1ug/mL	Name Sample Name RT ETS SS: 1ug/mL 2.606 ETS SS: 1ug/mL 2.609 ETS SS: 1ug/mL 2.610 ETS SS: 1ug/mL 2.610 ETS SS: 1ug/mL 2.611 ETS SS: 1ug/mL 0.002	ETS SS: 1ug/mL 2.606 20054524 ETS SS: 1ug/mL 2.609 20368670 ETS SS: 1ug/mL 2.610 20353450 ETS SS: 1ug/mL 2.610 20102186 ETS SS: 1ug/mL 2.611 20179382 2.609 20211642 0.002 143592.202	Name Sample Name RT Area USP Tailing ETS SS: 1ug/mL 2.606 20054524 1.2 ETS SS: 1ug/mL 2.609 20368670 1.2 ETS SS: 1ug/mL 2.610 20353450 1.2 ETS SS: 1ug/mL 2.610 20102186 1.2 ETS SS: 1ug/mL 2.611 20179382 1.2 ETS SS: 1ug/mL 2.609 20211642 0.002 143592.202 143592.202	

Figure 2. Instrument method in Empower 3 Software. The events table of an ACQUITY QDa Mass Detector programmed to automatically turn the diverter valve on and off. The diverter valve is in "Off / Flow to waste" position between 0.15 and 2.00 minutes to direct the flow of the mobile phase into waste.

System suitability

Performance of the MS method was verified by evaluating system suitability according to the specifications defined in the USP General Chapter, <621> Chromatography. System suitability results of five replicate injections of sample containing 1 μ g/mL of methyl and ethyl esters are shown in Figure 3. The retention times and peak areas repeatability met the USP specifications of less than 2.0% RSD. The USP peak tailing is less than the USP recommendations of \leq 1.5.

	Methyl p-toluenesulfonate							Ethyl p-toluenesulfonate							
System Suitability Sample Set ID: 5702 Result Set ID: 6523 Processed Channel Descr.: 2; QDa Positive(+) SIR							Sample Set ID. 5702 Result Set ID. 6523 Processed Channel Descr.: 2: QDa Positive(+) SIR								
						E									
	Ch12	04.00 Da. CV	=6. Sm	poothed by	9 poin	t Mean		Ch2 2	18.00 Da. CV	=6. Sn	noothed by	9 point	Mean		
		Nam	e: MTS	,				Name: ETS							
	Name	Sample Name	RT	Area	USP Tailing	K Prime		Name	Sample Name	RT	Area	USP Tailing	K Prime		
1	MTS	SS: 1ug/mL	2.327	11508409	1.2	8.3	1	ETS	SS: 1ug/mL	2.606	20054524	1.2	12.0		
2	MTS	SS: tug'mL	2.327	11420567	1.2	8.3	2	ETS	SS; fug/mL	2.609	20368670	1.2	12.0		
3	MTS	SS: fug/mL	2.328	11564739	1.2	8.3	3	ETS	SS: 1ug/mL	2.610	20353450	1.2	12.1		
4	MTS	SS: fug/mL	2.328	11501751	1.2	8.3	4	ETS	SS: 1ug/mL	2.610	20102186	1.2	12.1		
5	MTS	SS: fug/mL	2.329	11470791	1.2	8.3	5	ETS	SS: fug/mL	2,611	20179382	1.2	12.1		
Mean			2 328	11493251			Mean		-	2.609	20211642				
Std Dev.			0.001	52915, 196			Std Dev			0.002	143592 202				
% RSD			0.03	0.46	1	100	% RSD			0.07	0.71				

Figure 3. System suitability results for five replicate sample injections acquired using an ACQUITY QDa Mass Detector integrated with a diverter valve. Data processed using MS SIR acquisition mode.

Conclusion

An ACQUITY Diverter Valve was used to divert an unwanted API portion of the chromatographic injection away from the source of the ACQUITY QDa Mass Detector to waste. The flow was redirected back to the mass detector for analysis of genotoxic impurities, methyl and ethyl esters of p-toluenesulfonic acid.

Diverting the sample matrix or a high concentration of API minimizes potential ionization enhancement/suppression and contamination of the mass detector, improving the accuracy and consistency of the results for the low-level detection of genotoxic impurities.

Featured Products

ACQUITY QDa Mass Detector https://www.waters.com/134761404

Empower 3 Chromatography Data Software https://www.waters.com/513188

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