# Evaluation of Accurate Mass and Dynamic Range Capabilities of Low and High-Resolution Instrumentation in Compound Identification in Drug Discovery.

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ZQ. Single Quadrupole

Rank

2

1

6

MassWorks<sup>™</sup> software calibrates spectral accuracy (SA) using

a known standard. It includes mass accuracy and peak line

The calibrated spectrum is compared with the theoretical one

for a molecular formula candidate. Formula rank is based on

Scan speed plays important role in MassWorks<sup>™</sup> performance

Mass accuracy of all measurements was <10 mDa and <5 mDa

SA improves with slower scan speeds and is not influenced by

whether internal or external calibration is used (figure 2).

Table I. MassWorks molecular formula rank at 350 amu/s

Concentration<sup>b</sup> Compound 1 Compound 2 Compound 3

Rank

1

2

using external<sup>a</sup> spectral accuracy calibration.

Rank

1

1

3

shape calibration of the isotopic signature of the molecule.

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#### Overview

 Reliable accurate mass (AM) measurement plays an important role in compound identification.

 Accurate mass measurement typically requires use of expensive equipment and significant user expertise

• A simpler, more robust, and user friendly system is needed for routine use in the drug discovery environment

 Capabilities and limitations of time-of-flight (TOF) and guadrupole accurate mass measurements and compound molecular formula determination are compared.

## Introduction

 In drug discovery applications. AM is typically measured using orbitrap or TOF instruments. with time-to-digital (TDC) or analog-to-digital (ADC) converters

• TDC-based TOF instruments suffer from limited dynamic range due to dead time detector limitations.

 Replacement of TDC-TOF instruments with ADC instruments or an orbitrap is not always feasible due to budgetary constraints.

 Feasibility of routine <5 ppm AM measurement and molecular formula (MF) determination</li> was investigated using TDC-based TOF and single guadrupole MS instruments.

 For the TOF instrument, useful dynamic range was determined using Dynamic Range Enhancement (DRE)

· For the quadrupole instrument, AM and MF determination was done after spectral calibration with MassWorks<sup>™</sup> software with isotopic pattern calibration.

· Effect of MS scan parameters on AM, formula rank, and dynamic range was investigated.

Three model compounds with typical elemental composition in drug discovery were used.

	10					
(eq	5					
r (mDa		i	1	-		
lass Error	0	I	:		1	<ul> <li>1 mg/mL</li> </ul>
Aass	-5					<ul> <li>100 μg/ml</li> </ul>
~				-		* 10 μg/mL
	-10			*	1000 5000 60	1

Results

Figure 1. Influence of scan speed on mass accuracy.

uracy %	100 98 96 94	:		 	1				: esternal calibration
Spectral Acc	92 90 88		4		*	24			<ul> <li>internal calibration</li> <li>1 mg/mi,</li> <li>100 µg/mi,</li> <li>20 µg/mi,</li> </ul>
	86 84	150	350 J 5			2500 imu/		5000	
	ure ctra					of	sca	an :	speed on

LCTp. Time-of-Flight

• DRE tool alternates between high and low intensity beam to extend the dynamic range of the instrument

 DRE is calibrated using a lock mass in dead time saturation (attenuated lock mass) and normal lock mass (not saturated). Z-focus lens and a magnification factor are calibrated.

Without DRE, dynamic range with AM <5 ppm is 1-2 orders of magnitude (table II).</li>

With DRE, dynamic range extends to 3-4 orders of magnitude.

 Isotopic fit (i-FIT<sup>™</sup>. measure of isotopic pattern accuracy) improves with decreasing concentration. Formula rank is based on i-fit and mass accuracy.

	Conc.ª µg/mL	Compound		11	Compound 2			Compound 3		
		AM (ppm)	i-FIT™	Rank	AM (ppm)	i-FIT™	Rank	AM (ppm)	i-FIT™	Rank
	100	111.3			100.5			125.8		
No	10	15.3			3.9	13.4	3	14.5		
DRE	1	2.9	0.3	1	0.0	0.9	3	-2.6	0.2	1
	0.1	-1.1	2.5	3	7.5	1.1	3	6.8	0.2	3
	100	-0.3	120	1	0.7	4.9	1	-2.3	51.9	2
DRE	10	-4.0	94.8	2	5.5	14.3	2	2.6	11.4	2
	1	2.4	1.3	1	2.6	0.3	2	3.7	3.7	5
	0.1	-2.4	2.2	2	0.9	1.6	2	-13.2		

Compounds not detected <0.1ug

### Conclusions

 TDC based TOF instrument has only 1-2 orders of magnitude dynamic range for accurate mass measurement (<5 ppm).

It limits the usefulness in applications with wide concentration range samples.

DRE extends the dynamic range to 3-4 orders of magnitude.

DRE greatly simplifies instrument operation due to less stringent requirements to control ion beam intensity.

Significant user expertise is still required to achieve the desired results.

Accurate mass measurement and molecular formula ID is possible using a more economical single guadrupole instrument with MassWorks<sup>™</sup> spectral accuracy calibration

It has three orders of magnitude dynamic range, with better performance at higher concentrations.

Scan speed is a key parameter affecting the spectral accuracy.

Internal calibration did not improve spectral accuracy due to high stability of guadrupole mass analyzer as compared to TOF.

Compared to TOF, a single guadrupole instrument with MassWorks<sup>™</sup> is easier to use, more economical and well suited in support of drug discovery chemistry operations.

Compared to a single guadrupole instrument, a TOF instrument provides better sensitivity and is well suited to applications where sample concentration might be limited, such as in metabolite ID.

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# **Materials and Methods**

SA.

SA (%) = 100\*(1-RMSE)

at <2500 amu/s (figure 1).

	Single Quadrupole, ZQ	TOF, LCT Premier	Sample [M+H], Da MF Search criteria AM Criteria*				
	Single Quadrupole, 2Q	TOP, LOT Prennier					
Instrument	Waters ZQ + Agilent 1100	Waters LCT premier + Waters Acquity UPLC®	C <sub>21</sub> H <sub>19</sub> N <sub>3</sub> O <sub>4</sub> 378.1454 C <sub>1+100</sub> H <sub>1-100</sub> N <sub>0-20</sub> O <sub>0-20</sub> 5 ppm/5 mDa				
Column	Varian Polaris™ C18-A, 3 µm, 20 mm x 2.0 mm	Waters Acquity UPLC <sup>®</sup> HSS T3, 1.8 μm, 50 mm x 2.1 mm	C <sub>25</sub> H <sub>26</sub> N <sub>3</sub> O <sub>6</sub> Br 544.1083 C <sub>1-100</sub> H <sub>1-100</sub> N <sub>0-20</sub> O <sub>0-20</sub> Br <sub>0-5</sub> 5 ppm/5 mDa C <sub>21</sub> H <sub>25</sub> N <sub>7</sub> O <sub>6</sub> SCI 574.1042 C <sub>1-100</sub> H <sub>1-100</sub> N <sub>0-20</sub> O <sub>0-20</sub> S <sub>0-8</sub> Br <sub>0-5</sub> 5 ppm/5 mDa				
Flow Rate	1.8 mL/min	1.0 mL/min	* 5ppm TOF, 5mDa quadrupole				
Injection Volume	1 μL	2 μL					
Mobile Phase	A: 0.1% formic acid in H <sub>2</sub> O	A: 0.1% formic acid in H <sub>2</sub> O	Concentrations: 10 ng/mL - 1 mg/mL in methanol				
	B: 0.1% formic acid in CH <sub>3</sub> CN	B: 0.1% formic acid in CH <sub>3</sub> CN	Quadrupole Instrument:				
Gradient	5 to 95% B in 1.75 min	5 to 95% B in 2.0 min	<ul> <li>Compounds mixed with internal spectral accuracy standards.</li> <li>MassWorks<sup>™</sup> (ver. 2,0,2,0) used for spectral accuracy calibration (internal and external calibration)</li> </ul>				
MS detection, ESI+	Scan speed 250, 1000, 2500, and 5000 amu/s covering the appropriate mass range	200 – 800 amu in 0.5 s					
Resolution	Unit, ~0.5 amu peak width half height	~6000 FWHM	TOF Instrument:				
Mass Calibration	External mass calibration Internal and external Spectral Accuracy calibration using MassWorks	Fresh external mass calibration Lock mass using Lock Spray With DRE – two lock masses used	<ul> <li>Same day external mass calibration performed (100-1500 Da) at RMS</li> <li>5 ppm.</li> <li>DRE calibrated with <sup>12</sup>C peaks of reference compounds (in dead time).</li> <li>Z-focus lens and magnification factor determined.</li> <li>Lock masses used were <sup>13</sup>C peaks of reference compounds.</li> <li>Compounds run with and without using DRE.</li> </ul>				
Lock Mass or calibrant	imipramine ([M+H] 281.2018), 100 μg/mL buspirone ([M+H] 386.2556), 100 μg/mL tyr-tyr-tyr ([M+H] 508.2084), 1 mg/mL reserpine ([M+H] 609.2812), 100 μg/mL	buspirone- <sup>13</sup> C* ([M+H] 387.2771), 0.1 μg/mL leucine enkephalin- <sup>13</sup> C* ([M+H] 557.2805), 1 μg/mL *naturally occurring <sup>13</sup> C solopic mass peak					
DRE calibration	N/A	buspirone ([M+H] 386.2556), 0.1 μg/mL leucine enkephalin ([M+H] 556.2771), 1 μg/mL					

µg/mL

1000

100

10

a rank was the same with internal calibration.

<sup>b</sup> compounds not detected <10 µg/ml