

LDTD-MS/MS in 1.8 seconds with RSD of 2.4 % : Paracetamol in Human Plasma Crash

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Overview

- High-throughput determination of paracetamol in human plasma crash is performed by LDTD-MS/MS;
- Calibration range from 0.6 to 5000 ng/mL with $r^2 = 0.9944$;
- 1008 samples in 1.26 hours;
- Sample-to-sample run time of 4.5 seconds;
- Desorption peak width of 1.8 seconds;
- RSD of 2.4 % for 1008 replicates.

Instrumentation

- Phytronix Technologies LDTD ion source (model T-960);
- Thermo Fisher Scientific TSQ[®] Quantum[™] Ultra AM mass spectrometer.

LDTD ionization process

The LDTD ion source uses an infrared laser diode to desorb sample that have been dried onto a well of a LazWell[™] (96-well plate). The desorbed gas phase molecules are carried into a corona discharge region to undergo APCI, then they are transferred directly into the mass spectrometer for detection.

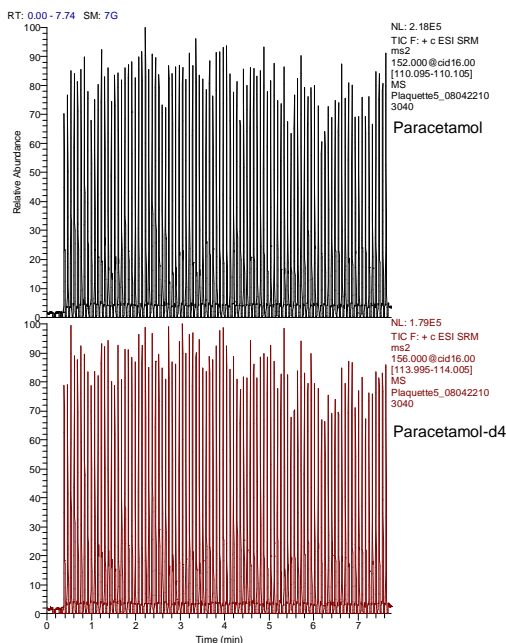
Samples Preparation

Stripped human plasma was spiked with paracetamol and paracetamol-d4 (ISTD). Plasma proteins were precipitated with the addition of acetonitrile (1:4 v/v). A volume of 2.0 μ L of the supernatant was manually transferred into a well of a LazWell[™] and was allowed to dry at room temperature.

Results and Discussion

LDTD-MS/MS allows high-throughput analysis of paracetamol in human plasma as shown in **Figure 1** where 96 replicates are analyzed in 7.2 minutes.

Figure 1 Area count of 96 replicates of paracetamol (40 ng/mL) in human plasma crash determined in LDTD-MS/MS.



Analysis Repeatability

Using only the raw area count of 1008 replicates, the RSD for paracetamol (40 ng/mL) and paracetamol-d4 (40 ng/mL) signal are 9.9 %. With the ISTD correction, the **RSD is lowered to 2.4 %** showing the stability of the LDTD-APCI process (**Figure 2**).

RSD of 2.4% over 1008 replicates

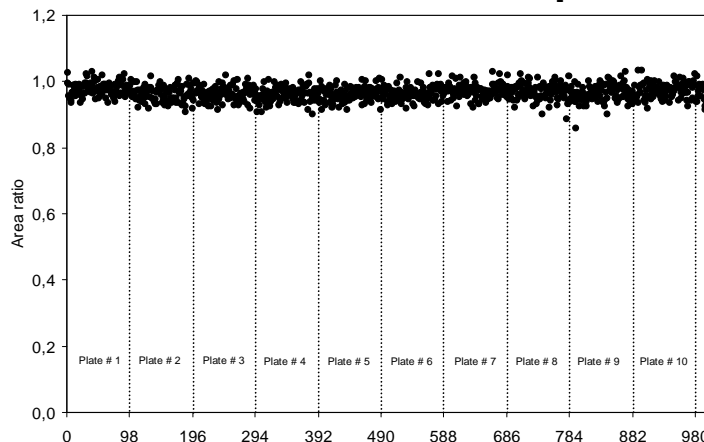


Figure 2 LDTD-MS/MS repeatability on paracetamol/ISTD ratio over 1008 human plasma crash replicates.

Paracetamol Calibration Curve

Quantitative determination of paracetamol in human plasma crash can be achieved over a nominal concentration range of 0.6 to 5000 ng/mL (Figure 3). An excellent linearity is obtained over the concentration range ($R^2 = 0.9944$).

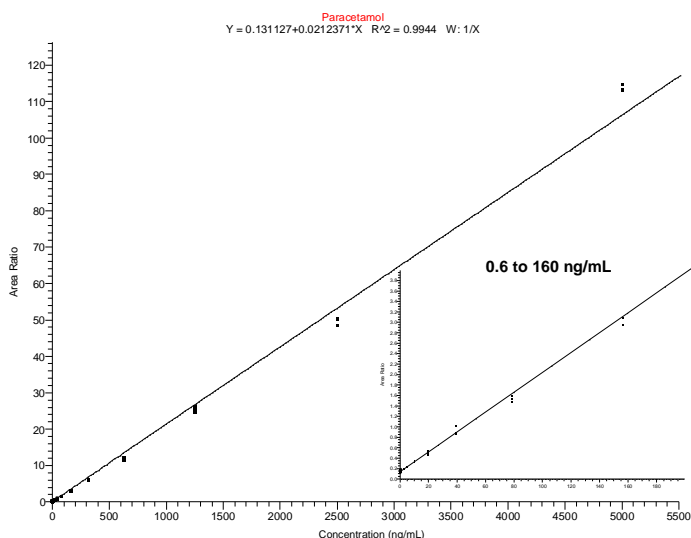


Figure 3 Paracetamol calibration curve in human plasma crash (3 replicates at each concentration).

Paracetamol and Paracetamol-d4 LDTD Desorption Profile

The Laser Diode Thermal Desorption process performed in **1.5 seconds** allows an excellent paracetamol signal width of 1.8 seconds. No carryover, no overlapping peak and no matrix effect are observed (Figure 4).

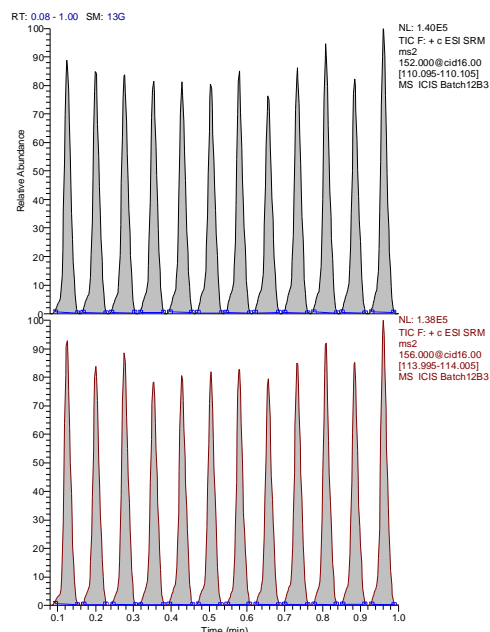


Figure 4 Area count of 12 samples of paracetamol in human plasma crash (40 ng/mL) analyzed in LDTD-MS/MS.

MS Parameters

APCI (+)	
Collision gas pressure	1.5 mTorr (Argon)
Collision energy	16 V
Tube lens	51 V
Scan time	0.050 s
Scan width	0.010 amu
Needle voltage	4500 V
Q1 width	0.30 amu
Q3 width	0.70 amu
Paracetamol SRM transition	152.0 → 110.1 amu
Paracetamol-d4 SRM transition	156.0 → 114.1 amu

LDTD Parameters

Laser power pattern	0 to 25 % in 1.0 s Hold at 25 % for 0.5 s
Carrier gas flow	3.0 L/min (Air)

Conclusions

LDTD-MS/MS allows ultra-fast paracetamol thermal desorption in **1.8 seconds** with a sample-to-sample run time of 4.5 seconds. The LDTD-MS/MS signal is stable over 1008 replicates as show by a RDS of 2.4 % with no matrix effect and no observed carryover. Paracetamol quantification can be performed over a 4-fold concentration range with an excellent linearity.

High-throughput analysis with excellent repeatability and linearity can be achieved using LDTD as ion source in mass spectrometry.

For more information about your specific application, visit www.phytronix.com

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