

ThermoFisher SCIENTIFIC

Emerging applications in routine mass spectrometry

Michal Godula Special Solutions Center Thermo Fisher Scientific

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Where do we go?





Chromatography is Important....Glyphosate Story





IC-MS/MS Chromatogram of polar pesticides in lettuce at 10 ug/kg





Thermo Scientific™ Integrion™ IC with TSQ Quantis™ MS/MS





Glyphosate analysis in water at ng/l levels

Name	LOD [ppt]	LOQ [ppt]	RSD % (10 ppt) level
Fosetyl-Al	1	2.5	5
Clopyralid	10	50	9
AMPA	2	5	9
Glyphosate	5	10	15
Glufosinate	2	5	4





Total Fipronil and Fipronil Sulfone Analysis Workflow Options



Easy and fast sample preparation





Thermo Scientific[™] TRACE[™] 1310 GC with Thermo Scientific[™] TSQ[™] 8000 Evo triple quadrupole GC-MS/MS



Thermo Scientific[™] TraceFinder[™] 4.1 data processing software



Results: Quantitative and Qualifier Ions (UHPLC-MS/MS)

Fipronil: 0.5 µg/kg



Fipronil sulfone: 0.5 µg/kg





Multi-Residue method for veterinary medicine residues with TSQ Altis

- 150+ compounds included in the method for the following classes of veterinary medicines:
 - Cefalosporins, macrolides, penicillins, quinolones, sulfas, tetracyclines, anthelmintics, nitroimidazoles, NSAIDs, sedatives, avermectins and coccidiostats, dyes (applied to fish), steroids (milk)





- QuEChERS based approach
 - EDTA/NH₄ oxalate solution and acetonitrile added to sample
 - Sample homogenised until sample fully dispersed
 - Sodium sulphate added before centrifugation
 - Dispersive SPE (CEC- C₁₈) clean-up
 - Add 1 mL H2O to 3mL extract, filter, inject



• Non-targeted acquisition:

- Same <u>sensitivity</u> as triple quadrupole
- Increase the <u>scope</u> of the analysis through full-scan
- Simple full-scan acquisition
- Perform targeted live or retrospective data processing

SANTE/11945/2015 Screening, identification, quantitation and confirmation requirements QqQ: 2 SRMs (Quan and Qual), Ion ratios, Peak characteristics HRAM: defined and can be applied (<5ppm, >2 fragment ions)

Solution

1. Screening , quantitation and identification by Q Exactive, ideally FullScan/(vDIA) to provide chromatographic profile of at least 2 HRAM *ions*

2. Confirmation with Q Exactive FullScan/ddMS2 to provide MS2 spectra



Screening Example – Acetamiprid in honey sample

Possible identification parameters



Acceptance criteria: accurate mass at correct Rt + presence 1 additional identification parameter



Results of vDIA – Sensitivity in Tea, 1 ul injection on column!

• Screening Detection Limits – Tea matrix



SDL values in Tea







Pesticide residues screening analysis in tea and honey using a Q Exactive Focus High-Resolution Mass Spectrometer

Authors

Keywords

vDIA

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Q Exactive Focus, Vanguish Flex,

QuEChERS, pesticide residues,

Goal

To develop and test a multi-residue instrumental method that can be applied for high-throughput screening and semi-quantitation of pesticide residues in food matrices at or below the current legislative requirements. A high-resolution, accurate-mass mass spectrometer operated in Full Scan – Variable Data-Independent Analysis (FS-vDIA)* mode providing an option for full spectrum filtering, retrospective analysis, and multi-parameter-based compound identification was applied. The method was validated for 328 target pesticides, with an option for the future extension to a larger number of analytes.

Introduction

European Commission directive EC 396/2005 sets maximum residue levels for pesticides in different products of plant and animal origin. The regulation presents significant analytical challenges with respect to the low limits of quantification and high number of target analytes. Nowadays, many GC- and LC-based methods are in routine use employing mostly low-resolution triple quadrupole mass spectrometric detection.

*vDIA method is not available in the United States of America.





vDIA method not available in the United States of America

Forensic Toxicology/Doping/Drugs of Abuse with ddMS2 workflow





Unknown Workflow – What is different? Can I identify the components?





Targeted Bottom-Up Proteomics Can Be Used For Meat Speciation





Peak Detection and Candidate Matching with GC Orbitrap EI Spectral Acquisition





- Right chromatography
- Total Workflows for LC-MS and GC-MS based applications
- Utilization of high resolution and accurate mass MS for screening, identification and quantification AND confirmation
- Merging of omics approaches with targeted methods allergens, proteins
- Authenticity and adulteration
- Novel data processing software tools Compound Discoverer
- You!

Blagodarja!

