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Comparison of UPLC-QTOF and GCMS for Detection of Designer Drugs in Urine Samples

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Disclaimer

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Project Background

- Attendees of electronic dance music festivals (EDM) demonstrate high rates of experimental drug use
- Collection of biological specimens of EDM attendees increases treatment ability for those experiencing adverse reactions and increases ability of toxicology labs to detect compounds



Project Objectives

- Analyze samples to obtain information regarding:
 - New drugs on the market
 - Prevalence of designer drugs
 - Identification of novel designer drugs and metabolites
 - Correlations and comparisons of designer drugs in blood, urine and oral fluid specimens

Sample Collection

- Approached participants on their way to EDM festival
- Location was ~100 yards from entrance gate
- Participants signed consent forms and were asked survey questions
- Samples collected included:
 - Oral Fluid Collection
 - Alere DDS2 Cartridge
 - Quantisal
 - Urine
 - Blood



Disclosure: Participants were not required to donate all 4 samples, and only donated samples based on their comfort level. The gift card incentive was only given if the participant donated a blood sample.

Urine Results

- Total number of urine samples collected: 104
- Samples underwent a battery of screen tests:
 - Immunoassay
 - Volatiles
 - RapidFire-MS/MS
 - **GC/MS**
 - **LC-QTOF**
 - LC-MS/MS

COMPARISON BETWEEN GC/MS AND LC-QTOF AS SCREENING TECHNIQUES

Sample Preparation (GC/MS)

- To 2 mL urine, add internal standard, 100 mM phosphate buffer (pH 6.0)
- To a copolymeric bonded phase extraction column:
 - **Condition:** Methanol, Water, 100 mM phosphate buffer
 - **Apply Sample**
 - **Wash:** Water, 20% Acetonitrile/Water, 100 mM Acetic Acid, then DRY
 - **Wash:** Hexane, Methanol, then DRY
 - **Elute:** Isopropanol, Ammonium Hydroxide, Methylene Chloride
- **Evaporate** (add 10% HCl) and **Reconstitute** with Acetonitrile

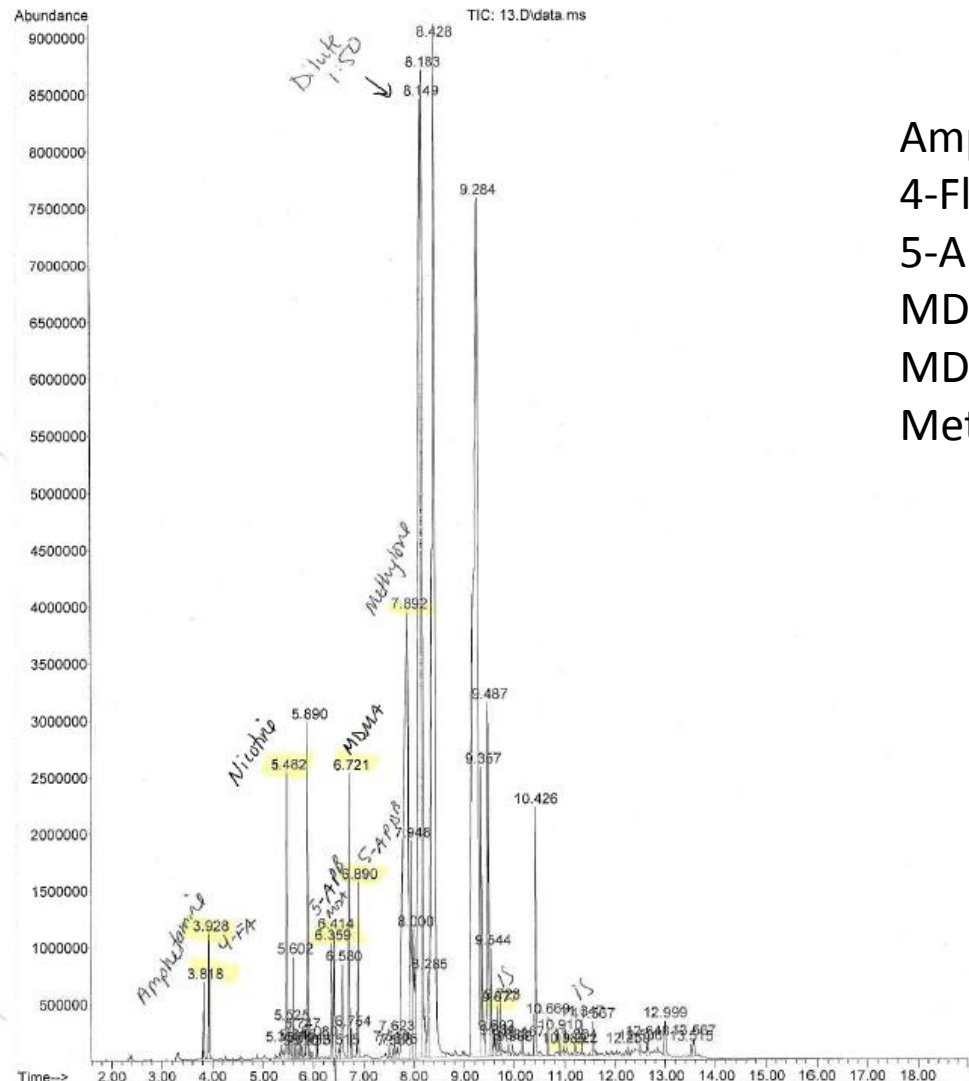
GC/MS Parameters

- Agilent GC (6890)/ MS (5975)
- Column: DB5MS 20m x 0.18mm x 0.18 μ m
- Split Ratio: 10:1
- Injection Temperature: 250°C
- Injection Volume: 2 μ L
- GC Oven Programming:
 - Initial 70°C (1 min)
 - Ramp 20°C/min
 - Final 300°C (5.5 min)
- Total Run Time: 17.5 min
- MS Acquisition: 42-550 m/z

Acceptability Criteria (GC/MS)

- Chromatographic peak must be clearly identifiable, as well as internal standard peak
- Chromatographic peak must be within $\pm 2\%$ of analyte in standard
 - If analyte is not present in a standard, standard is analyzed under same conditions to verify retention time
- Mass spectrum minimum confidence of 70% compared to reference library spectrum

Chromatogram of MS124 (GC/MS)



- Amphetamine
- 4-Fluoroamphetamine
- 5-APB
- MDA
- MDMA
- Methyone

Sample Preparation (LC-QTOF)

- To 0.5 mL urine, add internal standard, water, 100 mM phosphate buffer (pH 6.0)
- To a copolymeric bonded phase extraction column:
 - **Condition:** Methanol, Water, 100 mM phosphate buffer
 - **Apply Sample**
 - **Wash:** Water, 100 mM Acetic Acid, Methanol, then DRY
 - **Elute:** Isopropanol, Ammonium Hydroxide, Methylene Chloride
- **Evaporate** (add 10% HCl) and **Reconstitute** with Mobile Phase

LC-QTOF Parameters

- Waters Acquity I-Class UPLC Conditions:
 - Mobile phase A: 5mM ammonium formate (pH 3.0)
 - Mobile phase B: 0.1% formic acid in acetonitrile
 - Column: Waters Acquity HSS C18 150mm x 2.1mm x 1.8 μ m
 - Flow rate: 0.4 mL/min
 - Column Temperature: 50°C
 - Injection Volume: 2 μ L

Time (min)	%A	%B
Initial	87	13
0.5	87	13
10.0	50	50
10.75	5	95
12.25	5	95
12.5	87	13
15.0	87	13

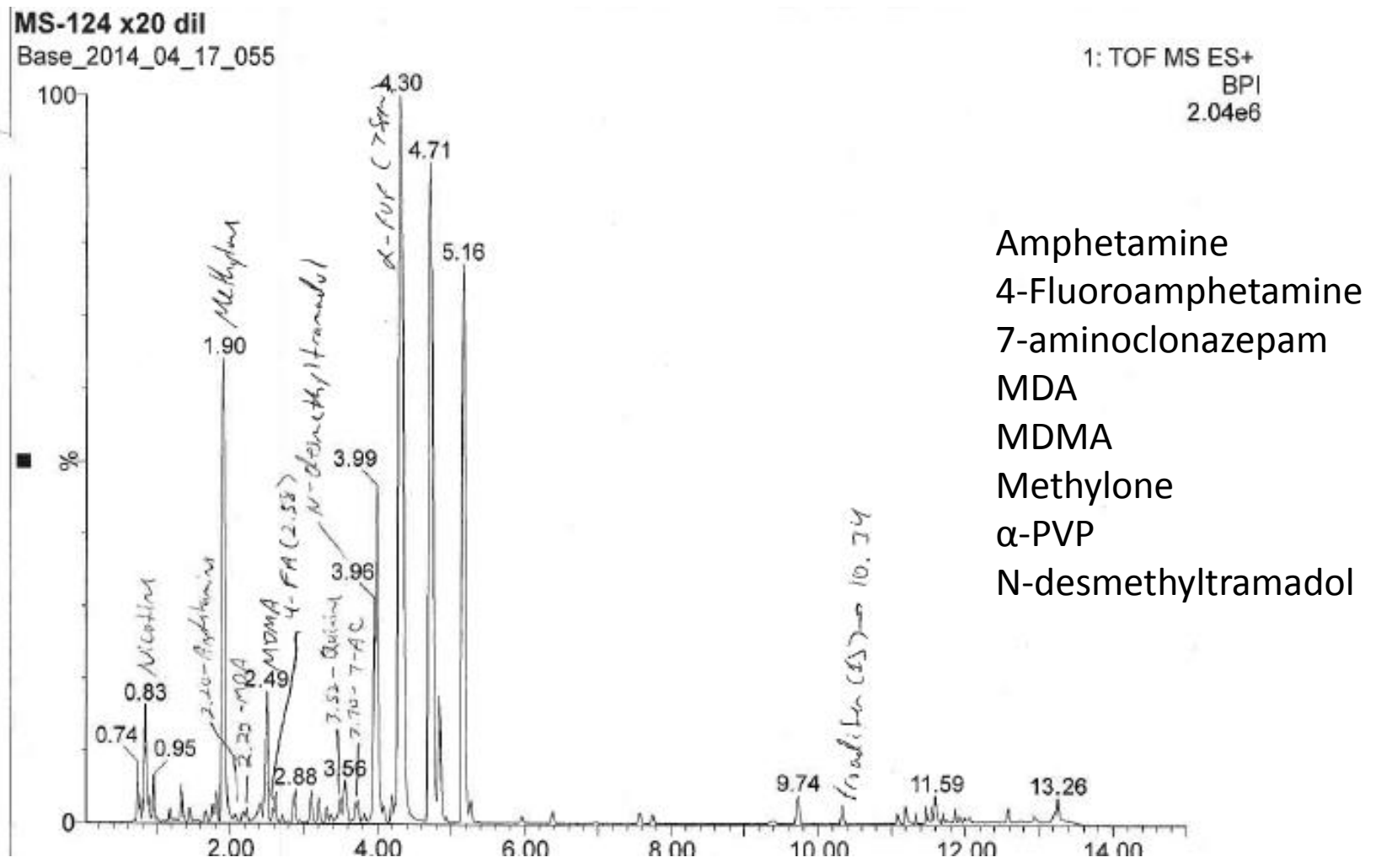
LC-QTOF Parameters

- Xevo G2 QTOF Conditions:
 - Ionization: Positive electrospray
 - Capillary voltage: 0.8 kV
 - Sample Cone Voltage: 20 V
 - Extraction Cone Voltage: 4 V
 - Source Temperature: 140°C
 - Desolvation Temperature/Flow: 500°C/900 L/h
 - Resolution Mode: 50-1000 m/z
 - Collision Energy (Function 1) – 6eV
 - Collision Energy (Function 2) – 10-40eV

Acceptability Criteria (LC-QTOF)

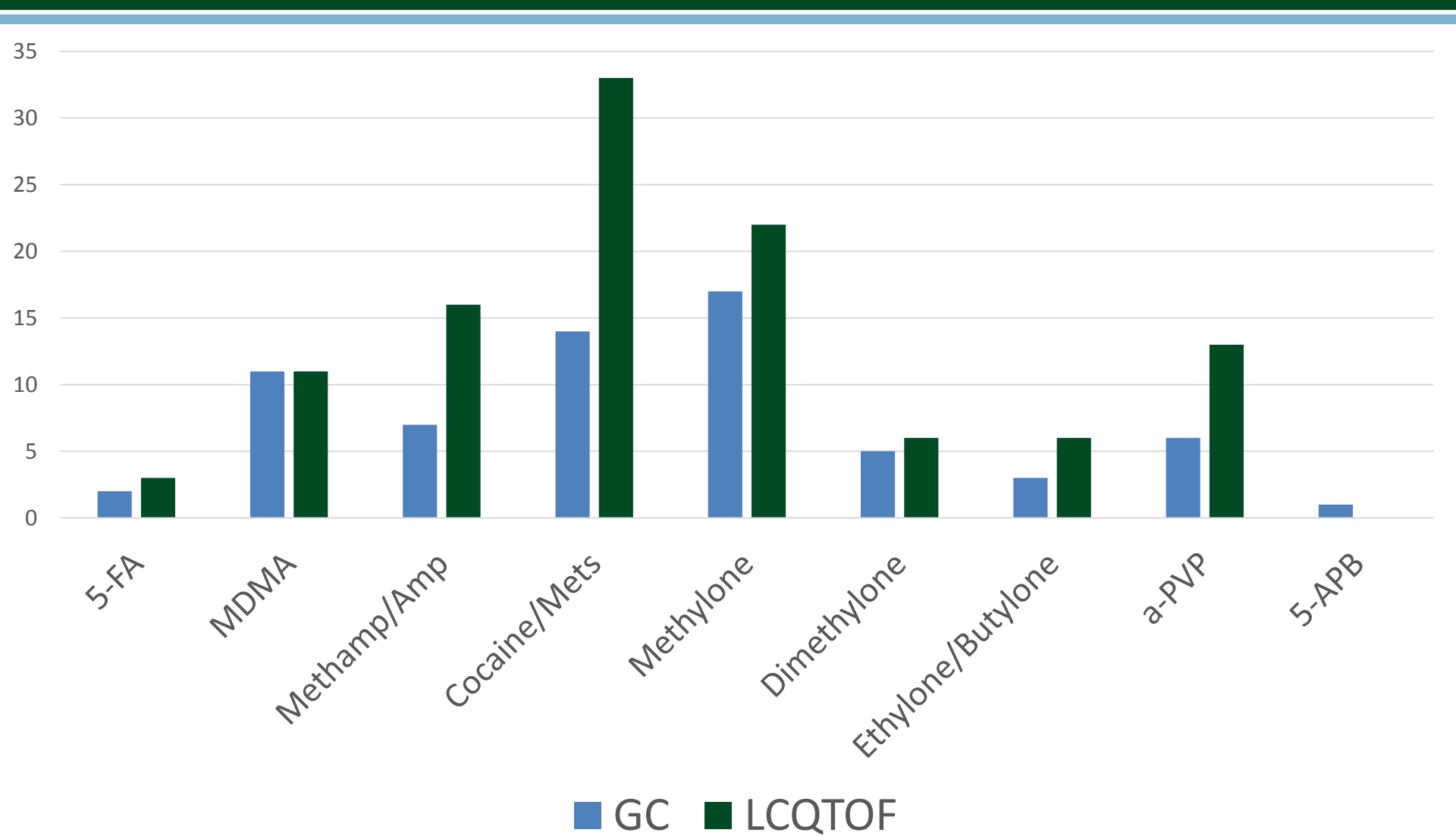
- Chromatographic peak must be clearly identifiable, as well as internal standard peak
- Chromatographic peak must be within $\pm 2\%$ of analyte in standard or within ± 0.3 min of analyte in database
 - If analyte is not present in a standard or database, standard is analyzed under same conditions to verify retention time
- Observed mass of molecular ion must be within ± 5 ppm of mass in database
- Observed mass of fragment ion must be within ± 5 ppm of mass in database

Chromatogram of MS124 (LC-QTOF)

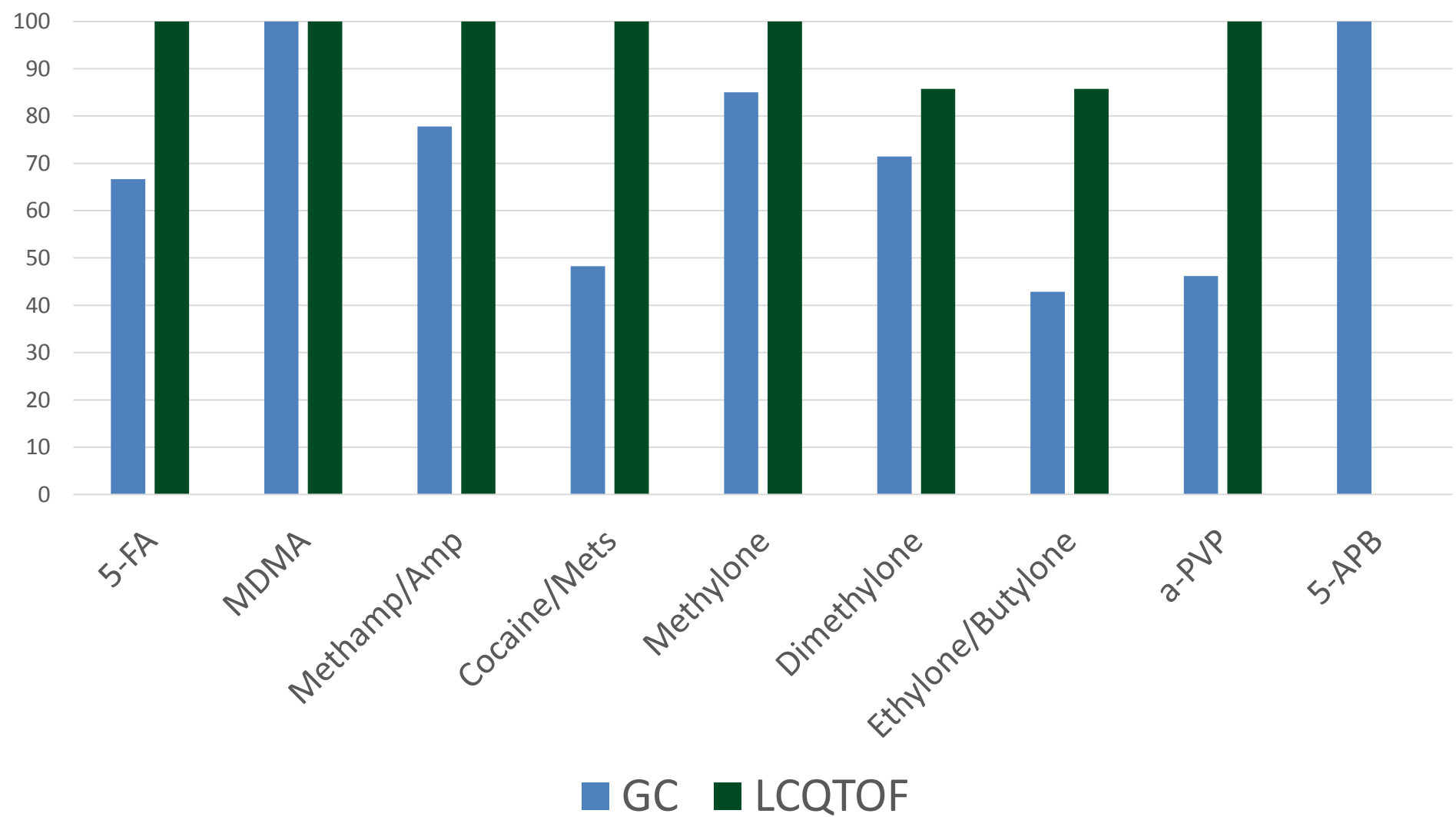


GC/MS AND LC-QTOF RESULTS

GC/MS vs. LC-QTOF Positive Screens



GC/MS vs. LC-QTOF Confirmation Rate



Unconfirmed Positives

Analytes	# Positives	# Confirm Positive
Methamp/Amp	16	9
Cocaine/Mets	33	29
Methylone	22	20

GC/MS vs. LC-QTOF

Rate	GC/MS		LC-QTOF	
	#	% of Total	#	% of Total
Sample Positivity Rate	49	80.3	63	103.3
False Negative Rate	12	19.6	0	0.0
Total Positive Samples	61	--	61	--

Alcohol Only Positives: 8

THC Only Positives: 16

Total Number of Positive Sample: $85 / 104 = 82\%$

GC/MS Results

- Missed analytes:
 - Benzoyllecgonine, THC, Cyclobenzaprine, DMAA, Alprazolam, Oxazepam, 7-aminoclonazepam, Psilocin, Buprenorphine, Azacyclonol, 3,4,5 Trimethoxy cocaine, PMMA, 2-CB
- Missed analytes due to sensitivity, no derivatization reagents used, poor chromatography on GC

LC-QTOF Results

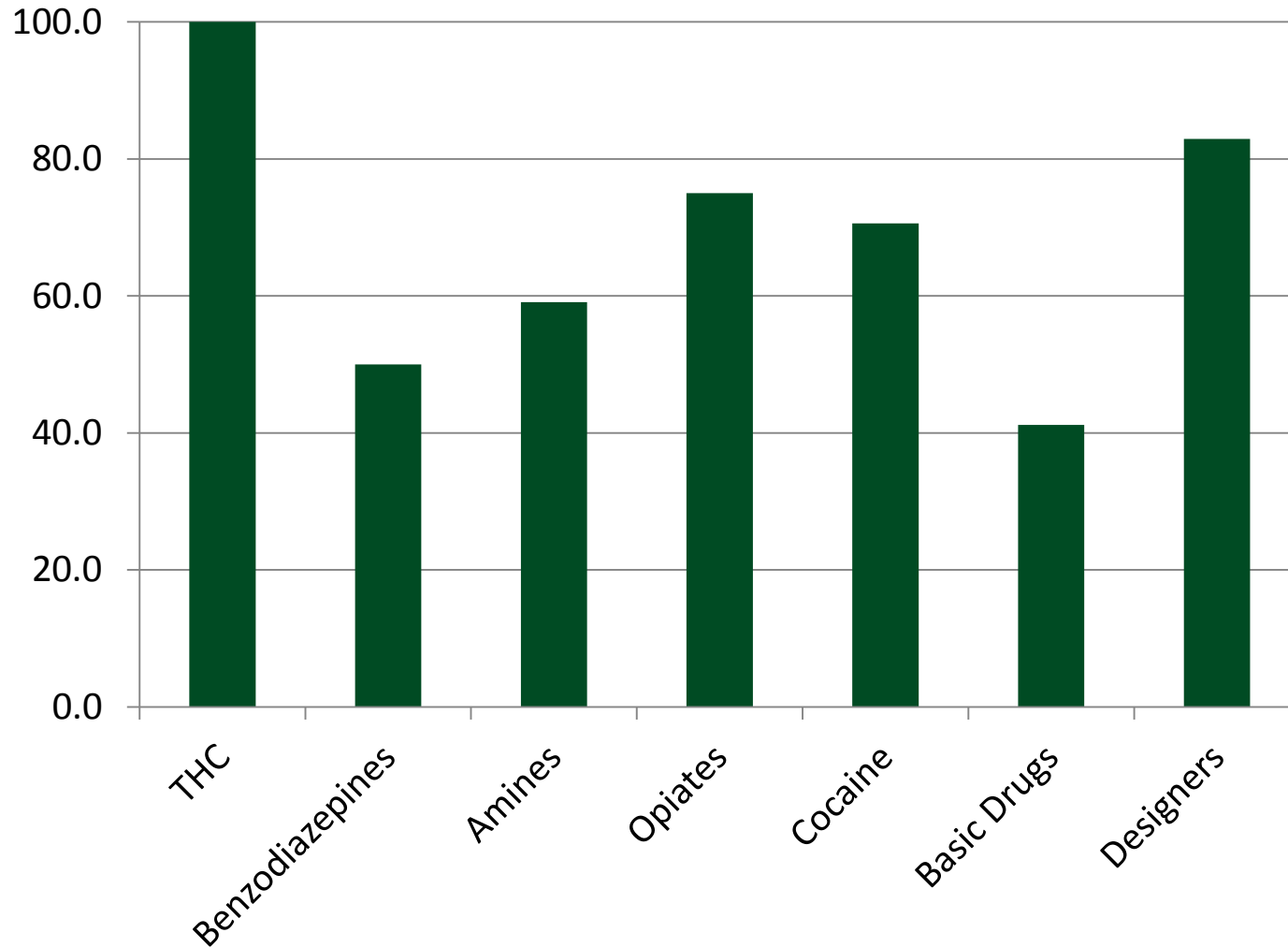
- Missed or poor chromatography analytes:
 - Ecgonine Methyl Ester, THC, 5-APB, Nicotine, Cotinine
- Extra analytes detected due to: increased sensitivity of QTOF vs. confirmation technique, compounds not analyzed for in confirmation technique

Comparison Conclusion

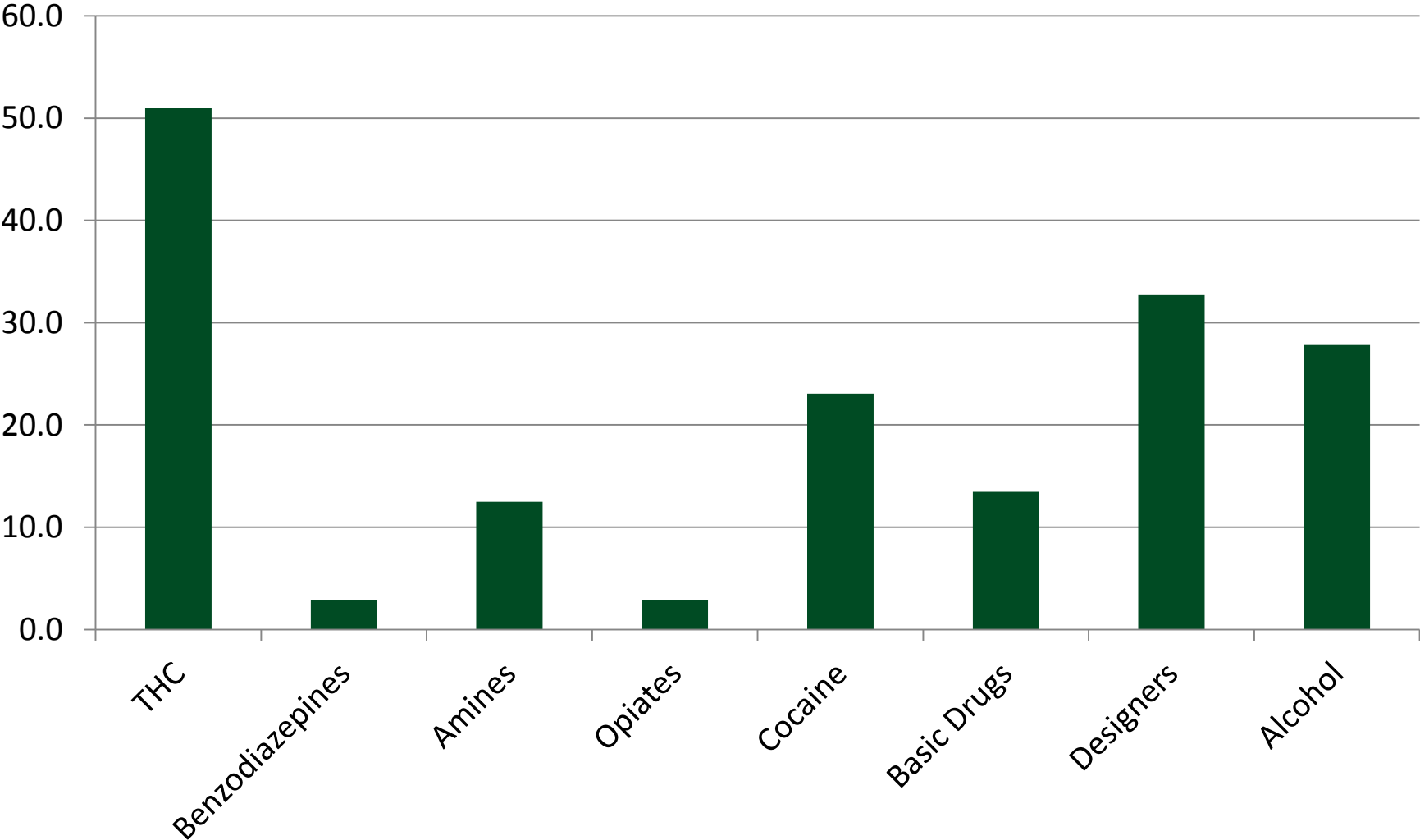
- GC/MS
 - Decreased sensitivity
 - Library search capabilities
 - More false negatives
 - Identified less designer drugs
 - Data interpretation requires less training
- LC-QTOF
 - Increased sensitivity
 - Targeted screen
 - More unconfirmed positives
 - Identified more designer drugs
 - Data interpretation requires increased training

OVERALL RESULTS FOR ANALYTICAL TESTING

Combined % Confirmation Rate



% Positive Rate in Sample Population



“Molly”

- Several participants indicated they had taken “Molly” in the last week
- Samples of subjects (9) who reported taking “Molly” contained:
 - MDMA
 - Methylone
 - Alpha-PVP
- Samples of subjects (15) who reported taking MDMA/Ecstasy contained:
 - MDMA
 - Methylone
 - Dimethylone/Ethylone/Butylone
 - Alpha-PVP

Thank You

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Questions?

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