

# HR-MS/MS as a way to avoid false positive quantification of 27 psychoactive compounds in venous and VAMS-collected blood

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Modern technique of precise analysis of 27 psychoactive compounds from DRUID list consist of: optimized sample preparation; reversed-phase chromatography on ExionAC LC coupled with ZenoTOF 7600 (SCIEX) and a processing method in SciexOS software for multilevel confirmation of the obtained result.

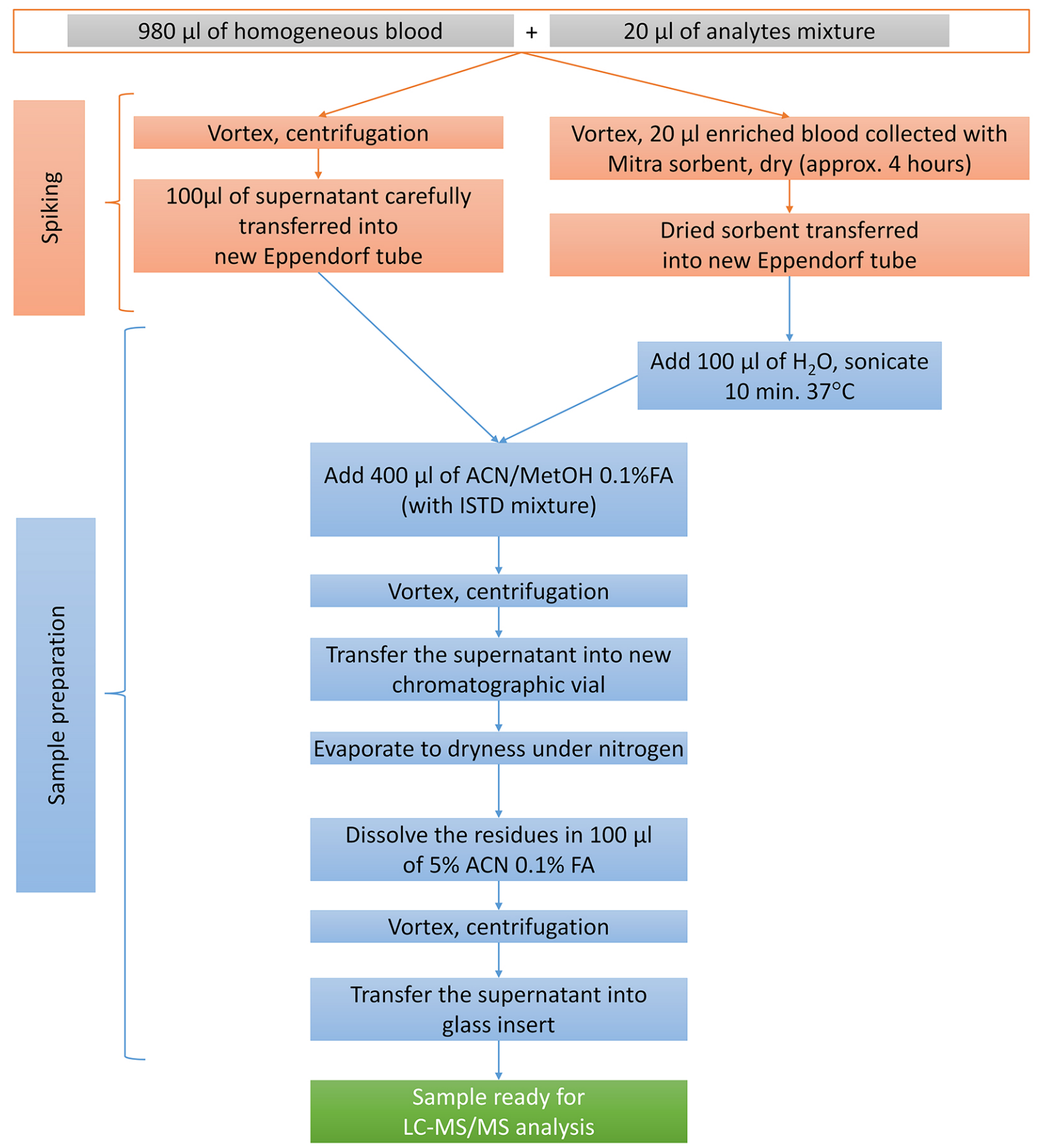


Figure 1. Sample preparation scheme.

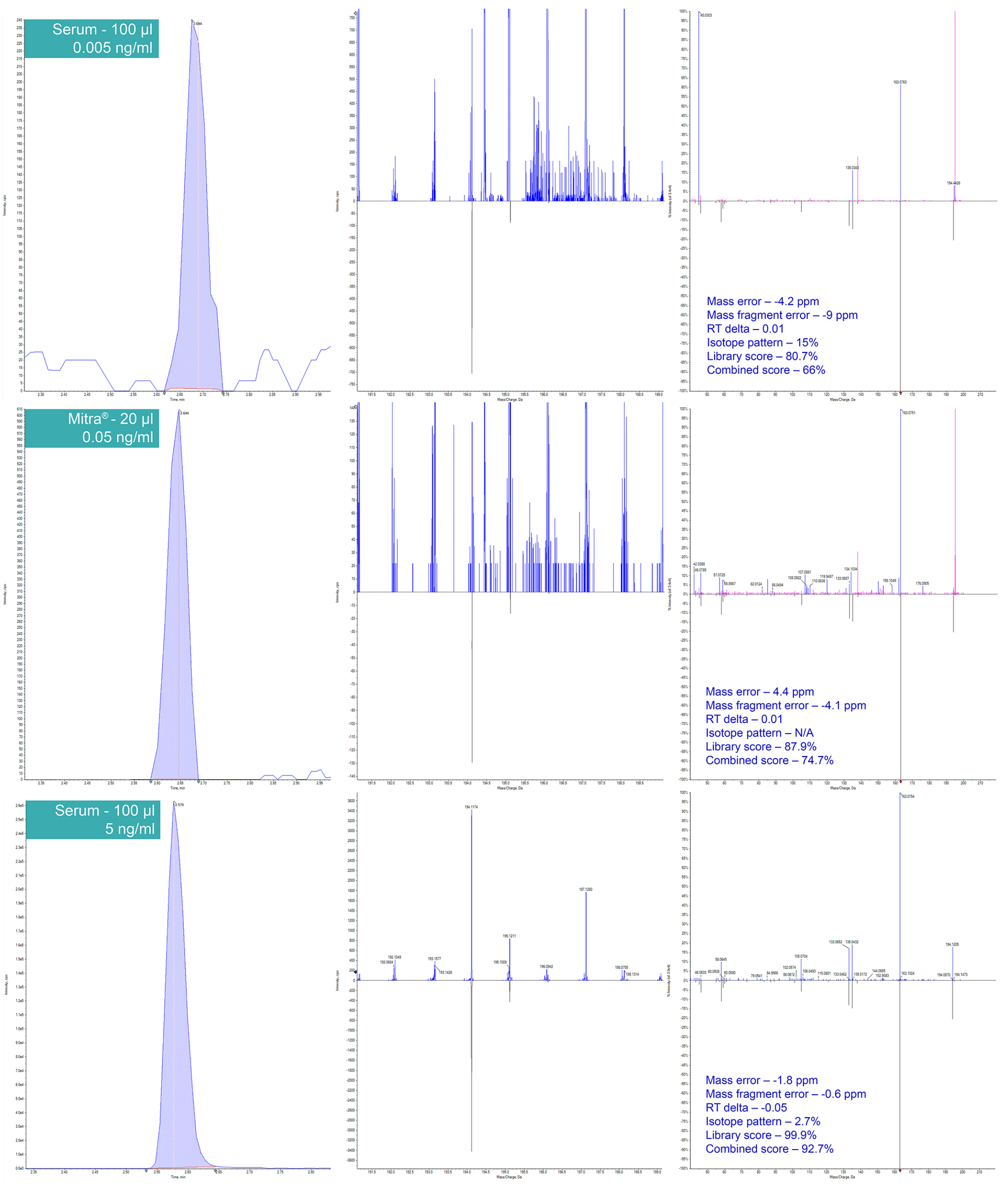


Figure 2. Examples of peak review and scoring data for MDMA in spiked samples.

Table 1. Limits of detection for analytes determined by the Druid method ZenoTOF LC-MS/MS system.

Analyte	LOD (blood) [ng/mL]	LOD (VAMS) [ng/mL]
MDA	0.050	0.100
MDEA	0.005	0.050
Zopiclone	0.005	0.050
Amphetamine	0.500	0.500
THC	0.500	0.500
Clonazepam	0.010	0.050
Cocaine	0.005	0.050
Tramadol	0.001	0.050
Methadone	0.001	0.010
Flunitrazepam	0.010	0.050
Methamphetamine	0.050	0.500
MDMA	0.005	0.050
Hydroxyzine	0.005	0.050
6-acetylmorphine	0.100	0.500
Benzylecgonine	0.005	0.010
Oxazepam	0.050	0.100
Alprazolam	0.005	0.050
Lorazepam	0.010	0.050
Diazepam	0.010	0.050
Nordiazepam	0.005	0.050
7-aminoclonazepam	0.050	0.100
7-aminoflunitrazepam	0.005	0.500
Fentanyl	0.001	0.050
Zolpidem	0.001	0.500
Codeine	0.050	0.100
Morphine	0.050	0.500
THC-COOH	0.500	0.500

The method was validated separately for venous blood and blood collected on VAMS and fulfilled the following criteria for each analyte: linearity (R ≥ 0.995), reproducibility (%CV ≤ 15%, accuracy 80 - 120%) and recovery (%CV ≤ 15%).

Developed processing method use multi-level confirmation and combined scoring based on HR MS and MS/MS data.

Apply	Qualitative Rule	Acceptable Difference	Marginal Difference	Unacceptable Difference	Combined Score Weight (%)
<input checked="" type="checkbox"/>	Mass Error (ppm)	< 5	< 10	>=	20
<input checked="" type="checkbox"/>	Fragment Mass Error (ppm)	< 5	< 10	>=	20
<input checked="" type="checkbox"/>	Error in Retention Time	< 2.5	< 40	>=	5
<input checked="" type="checkbox"/>	% Difference Isotope Ratio	< 5	< 20	>=	5
<input checked="" type="checkbox"/>	Library Hit Score	> 70	> 50	<=	50

After series of analysis the cut-off point of Combined Score estimated at 60% allows flawless reporting of analytes from the DRUID list. The library has been updated with MS/MS spectra obtained from spiked at different levels matrix samples with optimized collision energies for each compound.

Table 2. Example of multi-level confirmation and scoring by the DRUID method in VAMS blood sample on ZenoTOF 7600 LC-MS/MS system.

Formula	Reference Mass	Found At Mass	Mass Error (ppm)	Fragment Mass	Found Fragment Mass	Fragment Mass Error (ppm)	Mass Error Confidence	Fragment Mass Error Confidence	RT Confidence	Isotope Confidence	Library Confidence	Library Hit	Library Score	Combined Score	Isotope Ratio Difference
C12H19NO2	182.15	182.15	2.3	125.05	125.07	1.6	✓	✓	✓	✓	✓	MDA	99.5	92.80	0.8
C12H17NO2	208.13	208.13	-1.3	143.07	143.07	3.4	✓	✓	✓	✓	✓	MDA	100.0	92.80	0.7
C11H15NO2	184.14	184.17	-1.2	143.07	143.07	3.8	✓	✓	✓	✓	✓	MDA	99.9	92.75	1.1
C12H17NO2	184.14	184.14	-0.8	93.06	93.07	4.3	✓	✓	✓	✓	✓	Amphetamine [Smart Confirmation]	84.1	92.79	16.2
C12H17NO2	184.14	184.14	-1.2	205.04	205.07	2.8	✓	✓	✓	✓	✓	Hydroxyzine [Smart Confirmation]	100.0	94.20	0.8
C17H17NO	243.12	243.12	0.2	243.02	243.03	4.7	✓	✓	✓	✓	✓	Zopiclone [Smart Confirmation]	100.0	92.67	6.5
C12H17NO2	184.14	184.14	-1.0	243.02	243.03	4.1	✓	✓	✓	✓	✓	Methadone [Smart Confirmation]	100.0	93.19	3.8
C16H19NO4	285.19	285.19	0.9	168.10	168.10	4.1	✓	✓	✓	✓	✓	Benzylecgonine [Smart Confirmation]	99.3	93.13	0.8
C16H19NO2	264.18	264.18	-0.5	98.05	98.05	3.8	✓	✓	✓	✓	✓	Tramadol	100.0	94.88	0.9
C16H19NO2	264.18	264.18	-0.5	91.04	91.04	3.5	✓	✓	✓	✓	✓	Amphetamine [Smart Confirmation]	100.0	93.51	1.2
C12H17NO2	184.14	184.14	-1.5	198.12	198.12	1.2	✓	✓	✓	✓	✓	THC Positive	98.4	94.14	3.8
C15H17NO2	227.09	227.09	-0.7	241.05	241.05	2.9	✓	✓	✓	✓	✓	Chazepem [Smart Confirmation]	96.7	92.60	3.2
C16H19NO4	285.19	285.19	0.4	163.06	163.07	4.1	✓	✓	✓	✓	✓	6-Monoacetylmorphine [Smart Confirmation]	94.4	93.88	1.2
C16H19NO4	285.19	285.19	0.4	91.04	91.04	5.1	✓	✓	✓	✓	✓	Methamphetamine [Smart Confirmation]	98.8	88.95	0.4
C17H19NO2	267.17	267.17	-0.8	183.18	183.18	1.7	✓	✓	✓	✓	✓	Cocaine O3 [Smart Confirmation]	100.0	93.91	10.4
C17H19NO2	267.17	267.17	-0.7	183.18	183.18	4.2	✓	✓	✓	✓	✓	Cocaine [Smart Confirmation]	100.0	94.00	0.7
C17H19NO4	309.09	309.09	-1.4	281.07	281.07	6.2	✓	✓	✓	✓	✓	Alprazolam	98.7	88.74	6.3
C17H19NO2	267.17	267.17	-0.8	327.19	327.19	-0.9	✓	✓	✓	✓	✓	THC-COOH Positive	98.9	92.73	1.6
C16H19NO2	264.18	264.18	-1.8	133.07	133.07	2.1	✓	✓	✓	✓	✓	7-Aminoclonazepam	87.9	92.89	5.8
C12H17NO2	184.14	184.14	-0.8	188.14	188.14	4.7	✓	✓	✓	✓	✓	Fentanyl [Smart Confirmation]	100.0	93.00	1.4
C15H17NO2	227.09	227.09	-1.0	140.08	140.08	2.3	✓	✓	✓	✓	✓	Nordiazepam	99.3	93.75	5.8
C16H19NO2	264.18	264.18	-0.7	230.09	230.09	0.9	✓	✓	✓	✓	✓	Flunitrazepam	94.7	94.37	2.8
C12H17NO2	184.14	184.14	-1.5	198.14	198.14	-8.2	✓	✓	✓	✓	✓	THC-8 [Smart Confirmation]	85.7	76.18	5.8
C16H19NO2	264.18	264.18	-0.6	193.08	193.08	2.0	✓	✓	✓	✓	✓	Diazepam	98.5	94.50	1.4
C15H17NO2	227.09	227.09	0.9	270.09	270.09	1.7	✓	✓	✓	✓	✓	Cocaine [Smart Confirmation]	97.7	94.26	3.8
C15H17NO2	227.09	227.09	-0.4	121.07	121.07	-0.4	✓	✓	✓	✓	✓	7-Aminoclonazepam [Smart Confirmation]	97.9	96.71	4.5
C15H17NO2	227.09	227.09	-1.0	270.09	270.09	3.5	✓	✓	✓	✓	✓	Lorazepam	99.8	93.37	3.8
C17H19NO2	267.17	267.17	-0.3	133.07	133.07	-1.3	✓	✓	✓	✓	✓	Morphine [Smart Confirmation]	99.1	96.80	4.4
C16H19NO2	264.18	264.18	-0.9	233.12	233.12	6.1	✓	✓	✓	✓	✓	Zolpidem	100.0	93.87	0.8
C17H19NO2	267.17	267.17	-1.1	210.10	210.10	0.1	✓	✓	✓	✓	✓	DS-Alprazolam [Smart Confirmation]	89.2	95.43	3.3
C16H19NO2	264.18	264.18	0.1	133.07	133.07	0.8	✓	✓	✓	✓	✓	Cocaine [Smart Confirmation]	98.4	98.10	2.4
C17H19NO2	267.17	267.17	-0.6	133.07	133.07	1.9	✓	✓	✓	✓	✓	Morphine O3 [Smart Confirmation]	98.7	93.83	1.2

Compound presence confirmation is one of the key aspects of targeted and quantitative analysis in forensics applications. High resolution tandem mass spectrometry (HR-MS/MS) may become a major tool for this kind of analysis because of its mass accuracy resulting in unparalleled specificity, all combined with broad linearity and scanning speed achieved in the newest hardware on the market

**“Sensitive, specific and false positive resistant HR-MS/MS quantitation of 27 psychoactive compounds in blood and blood collected by VAMS.”**