



ANALYTICAL REPORT¹

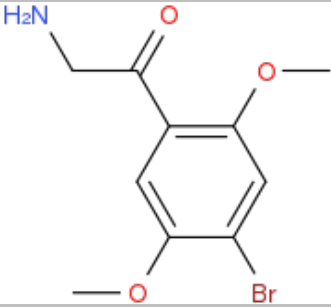
bk-2C-B (

C₁₀H₁₂BrNO₃)

2-amino-1-(4-bromo-2,5-dimethoxyphenyl)ethan-1-one

Remark – other NPS detected: **none**

Sample ID:	1350-15
Sample description:	powder - white
Sample type:	test purchase /RESPONSE -purchasing
Date of sample receipt (M/D/Y):	11/6/2015
Date of entry (M/D/Y) into NFL database:	11/25/2015
Report updates (if any) will be published here:	http://www.policija.si/apps/nfl_response_web/seznam.php

Substance identified - structure ² (base form)	
Systematic name	2-amino-1-(4-bromo-2,5-dimethoxyphenyl)ethan-1-one
Other names	
Formula (per base form)	C ₁₀ H ₁₂ BrNO ₃
M _w (g/mol)	274,11
Salt form	HCl
StdInChIKey	HFYJGAIQIDRPX-UHFFFAOYSA-N
Compound Class	Arylalkylamines
Other NPS detected	none
Add.info (purity..)	GC - MS two small peaks 2%) observed, chloride and bromide anions)

¹ This report has been produced with the financial support of the Prevention of and fight against crime Programme of the European Union (grant agreement number JUST/2013/ISEC/DRUGS/AG/6413). The contents of this report are the sole responsibility of the National Forensic Laboratory and can in no way be taken to reflect the views of the European Commission.

² Created by OPSIN free tool: <http://opsin.ch.cam.ac.uk/> DOI: 10.1021/ci100384d

Report updates

date	comments (explanation)

Instrumental methods (if applied) in NFL

1. GC-MS (Agilent): GC-method is RT locked to tetracosane (RT=9.53 min). Injection volume 1 ml and split mode (1:50) . Injector temperature: 280 °C. Chromatographic separation: on column HP1-MS (100% dimethylpolysiloxane), length 30 m, internal diameter 0.25 mm, film thickness 0.25 mm. Carrier gas He: flow-rate 1.2 ml/min. GC oven program: 170 °C for 1 min, followed by heating up to 293 °C at a rate of 18 °C/min, hold for 6.1 min, than heating at 50 °C/min up to 325 °C and finally 2.8 min isothermal. MSD source EI = 70 eV. GC-MS transfer line T= 235°C, source and quadropole temperatures 280°C and 180°C, respectively. Scan range m/z scan range: from 50 (40) to 550 amu.

2. HPLC-TOF (Agilent): 6230B TOF with Agilent 1260 Infinity HPLC with binary pump, column: Zorbax Eclipse XDB-C18, 50 x 4.6 mm, 1.8 micron. Mobile phases (A) 0.1% formic acid and 1mM ammonium formate in water; (B) 0.1% formic acid in methanol (B). Gradient: starting at 5% B, changing to 40% B over 4 min, then to 70% over 2 min and in 5 min to 100%, hold 1 min and back to 5%, equilibration for 1.7 min. The flow rate: 1.0 ml/min; Injection volume 1 µl. MS parameters: 2GHz, Extended Dynamic range mode to a maximum of 1700 amu, acquisition rate 1.30 spectra/sec. Sample ionisation: by Agilent Jet Stream technology (Dual AJS ESI). Ion source: positive ion scan mode with mass scanning from 82 to 1000 amu. Other TOF parameters: drying gas (N₂) and sheath temperature 325 °C; drying gas flow rate 6 l/min; sheath gas flow rate 8 l/min; nebulizer 25 psig; Vcap. 4000 V; nozzle 2000 V; skimmer 65 V; fragmentor 175 V and Octopole RF 750 V.

3. FTIR-ATR (Perkin Elmer): scan range 4000-400 cm⁻¹; resolution 4cm⁻¹

4. GC- (MS)-IR condensed phase (GC-MS (Agilent) & IR (Spectra analyses-Danny)

GC-method: Injection volume 1 ml and split mode (1:5). Injector temperature 280 °C. Chromatographic separation as above (**1**). Split MS : IR = 1:9.

MSD source EI = 70 eV. GC-MS transfer line T= 235°C, source and quadropole temperatures 280°C and 180°C, respectively. Scan range m/z scan range: from 50 (40) to 550 amu.

IR (condensed phase): IR scan range 4000 to 650, resolution 4 cm⁻¹.

5. IC (anions) (Thermo Scientific, Dionex ICS 2100), Column: IonPac AS19, 2 x 250mm; Eluent: 10mM from 0 to 10 min, 10-58 mM from 10 to 40min; Flow rate: 0.25 ml/min; Temperature: 30°C; Suppressor: AERS 500 2mm, suppressor current 13mA; Inj. Volume: 25 µl

Supporting information

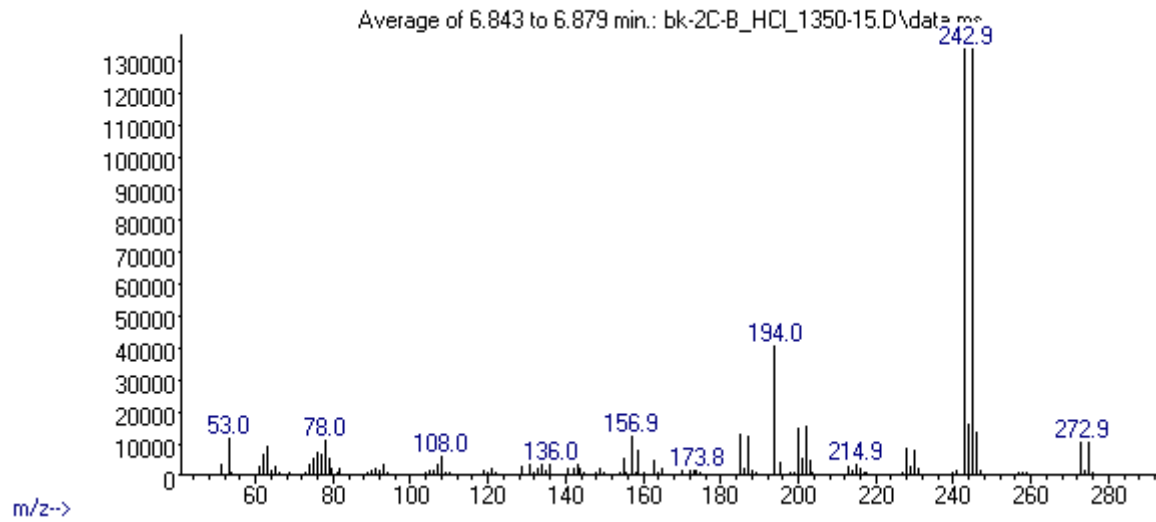
Solubility in	result/remark
CH ₂ Cl ₂	good- few non dissolved particles
MeOH	soluble
H ₂ O	

Analytical technique:	applied	remarks
GC-MS (EI ionization)	+	NFL GC-RT (min): 6,88 BP(1): 243; BP(2): 245, BP(3) :194,
HPLC-TOF	+	Exact mass (theoretical): 273,0001; measured value Δppm: -0,96; formula: <chem>C10H12BrNO3</chem>
FTIR-ATR	+	direct measurement (sample as received)
FTIR (condensed phase) always as base form		
IC (anions)	+	
NMR (in FKKT)	-	no
validation		MS-consistent with SWG DRUG (QM =0.99),
other		

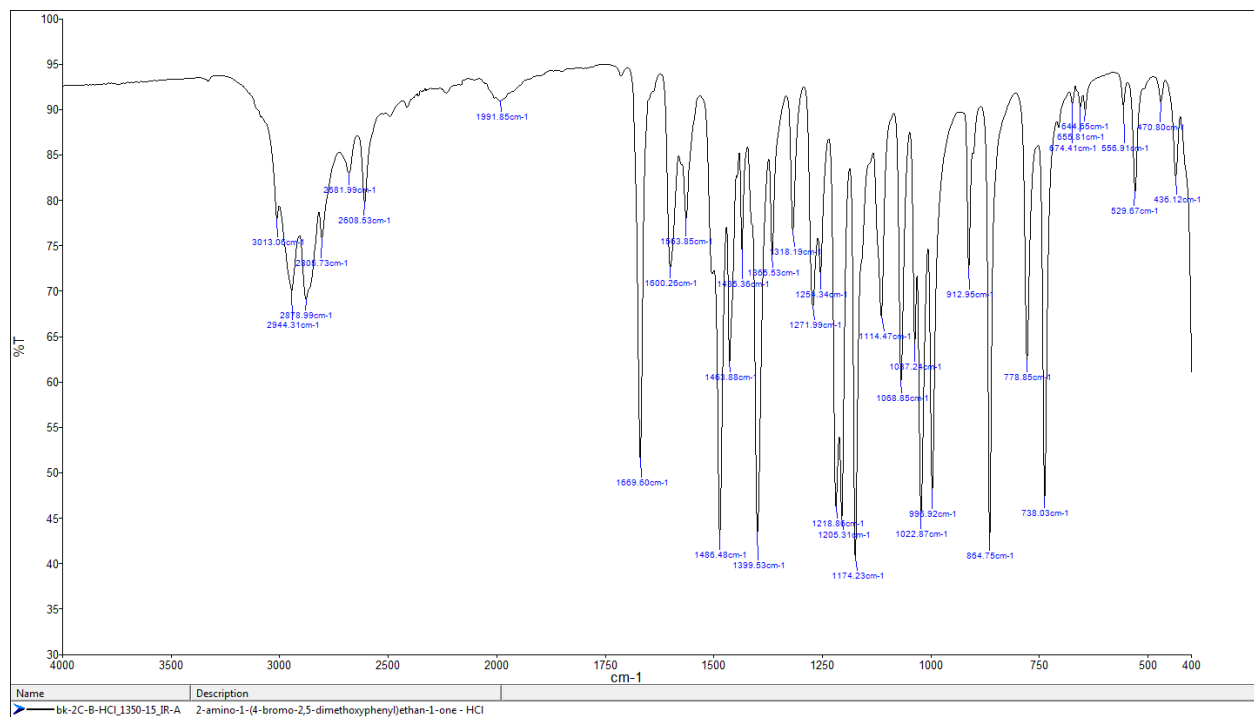
ANALYTICAL RESULTS

MS (EI)

Abundance



FTIR-ATR - direct measurement (sample as received)



IR (condensed phase – after chromatographic separation)

TOF REPORT

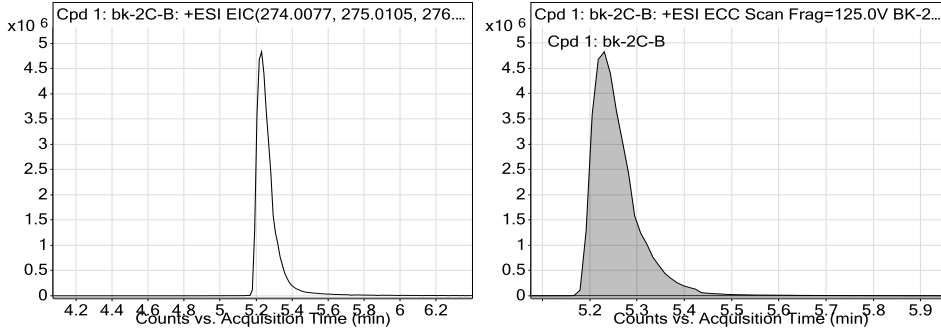
Data File	BK-2C-B_1350-15_TOF.d	Sample Name	ID_1350-15
Sample Type	Sample	Position	P1-A5
Instrument Name	6230B TOF LC-MS	User Name	TG
Acq Method	general-17112015-XDB-C18-ESI-poz.m	Acquired Time	11/17/2015 10:14:15 AM
IRM Calibration Status	Success	DA Method	Drugs_NFL.m
Comment	extract in MeOH		

Compound Table

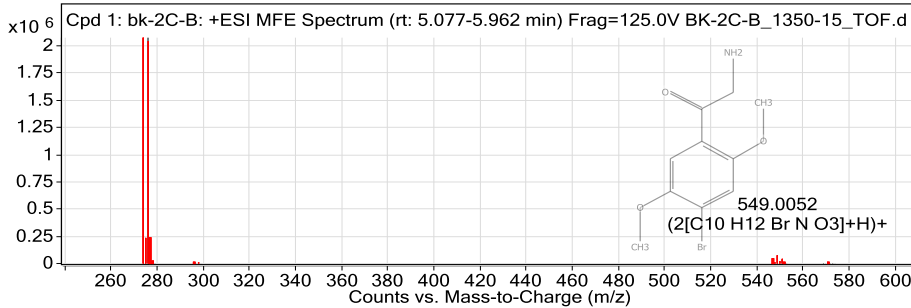
Label	Compound Name	MFG Formula	Obs. RT	Obs. Mass
Cpd 1: bk-2C-B	bk-2C-B	C10 H12 Br N O3	5.235	273.0003

Name	Obs. m/z	Obs. RT	Obs. Mass	DB RT	DB Formula	DB Mass	DB Mass Error (ppm)
bk-2C-B	274.0076	5.235	273.0003	5.24	C10 H12 Br N O3	273.0001	-0.96

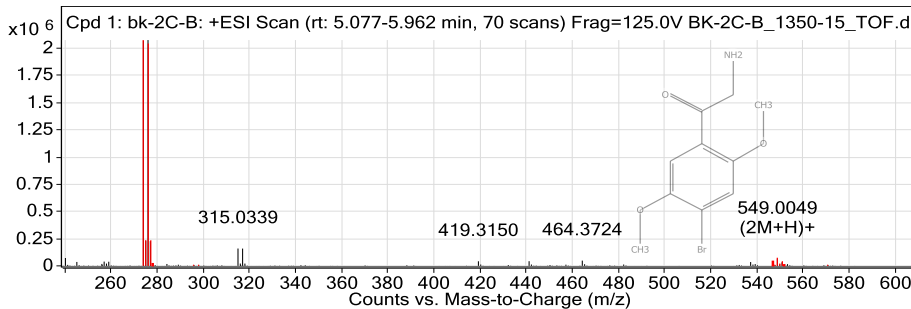
Compound Chromatograms



MFE MS Zoomed Spectrum



MS Zoomed Spectrum



MS Spectrum Peak List

Obs. m/z	Charge	Abund	Formula	Ion/Isotope
274.0076	1	2067959.13	C10 H12 Br N O3	(M+H)+
275.0112	1	231256.97	C10 H12 Br N O3	(M+H)+
276.0055	1	2075617.38	C10 H12 Br N O3	(M+H)+
277.0091	1	227021.2	C10 H12 Br N O3	(M+H)+
278.0108	1	23229.75	C10 H12 Br N O3	(M+H)+
547.007	1	36460.88	C10 H12 Br N O3	(2M+H)+
549.0052	1	77291.05	C10 H12 Br N O3	(2M+H)+
550.0079	1	16559.73	C10 H12 Br N O3	(2M+H)+
551.0036	1	37477.05	C10 H12 Br N O3	(2M+H)+
552.0067	1	8086.31	C10 H12 Br N O3	(2M+H)+

--- End Of Report ---

Peak Integration Report

Sample Name:	bk-2C-B_1350-15_IC	Inj. Vol.:	25,00
Injection Type:	Unknown	Dilution Factor:	1,0000
Program:	ANIONI	Operator:	kemija
Inj. Date / Time:	17-nov-2015 / 19:35	Run Time:	42,00

No.	Time min	Peak Name	Peak Type	Area $\mu\text{S}\cdot\text{min}$	Height μS	Amount mg/L
1,00	9,24	Chloride	BMB	5,94	25,34	n.a.
2,00	14,43	Bromide	BMB	3,27	6,10	n.a.
TOTAL:				9,22	31,43	0,00

