

Identity and Purity

Proton NMR Spectrum

Identity: the signals are consistent with the proposed structure and in accord with literature where available.

Isotopic Labelling: signals at the sites of deuteration are absent, compared with the spectrum of unlabelled material, indicating clean deuteration.

Residual Solvents: a trace (under 0.1 % w/w) of dichloromethane is observed.

Impurities: no significant impurities are evident in the spectrum.

Carbon-13 NMR Spectrum

Identity: the signals are consistent with the proposed structure and in accord with literature where available.

Isotopic Labelling: signals at the sites of deuteration have collapsed to small multiplets compared with the spectrum of unlabelled material, indicating clean deuteration.

High-resolution Mass Spectrum (ESI+)

Found m/z 438.3219. $C_{29}H_{24}D_{10}NO_2 [M]^+$ requires m/z 438.3217. The deviation of 0.5 ppm is within normally accepted limits for the establishment of identity by HRMS. No signal for d_0 material was seen (detection limit about 0.5 %).

HPLC

A somewhat broadened, tailing peak is observed (99.3 %). Note: in the absence of reference materials for preparing calibration curves, it is assumed that all peaks have the same detector response. Where possible, the conditions of analysis follow a pharmacopeial or literature method, or have been adapted from same.

Elemental Analysis

	Found:	C 67.14, H 4.74, D 3.95, N 2.60 %
$C_{29}H_{24}D_{10}NO_2 \cdot Br$	Requires:	C 67.17, H 4.67, D 3.88, N 2.70 %

The elemental analyses fall within generally accepted limits for establishing the molecular formula given. The results may also be taken to imply the absence of significant quantities of water or inorganic salts (which have not been elsewhere tested for because of sample size limitations).

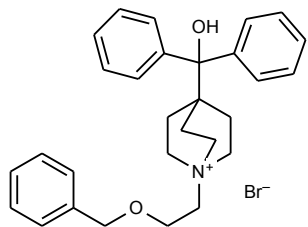
The available quantity of custom-synthesised material is always small, and this limits the extent and type of analytical data which can be obtained. This Certificate is presented in descriptive format for use by analytical chemists who are trained in the use of custom-synthesised materials. Custom materials often contain higher levels of residual solvents and/or water, and we urge you to use the corrected purity where needed rather than the raw HPLC purity. This compound is intended for use as an analytical reference material and it is not for human administration. Structures are shown with relative stereochemistry unless otherwise specified.

The re-test date is assigned from experience gained with the material in the laboratory and/or on storage. It is not possible to perform formal storage studies because of the small amount of material available.



Proton NMR Spectrum of Umeclidinium Bromide (top) and Umeclidinium-d₁₀ Bromide (bottom) in DMSO-d₆

BDG SYNTHESIS



15xArH
15.1H

OH
1.0H

CH₂
2.0H

OCH₂
2.0H

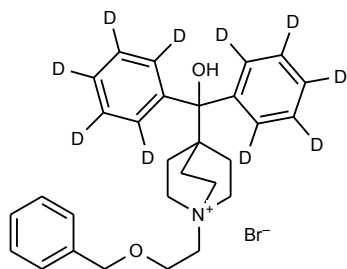
3xNCH₂
6.0H

NCH₂
2.0H

H₂O

NMR
Solvent

3xCH₂
6.0H



5xArH
5.0H

OH
1.0H

CH₂
2.0H

OCH₂
2.0H

3xNCH₂
6.0H

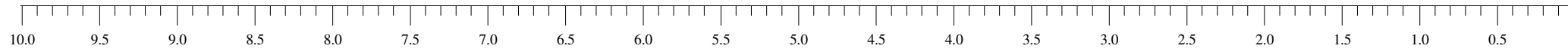
NCH₂
2.0H

H₂O

NMR
Solvent

3xCH₂
6.0H

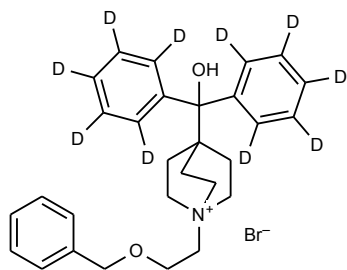
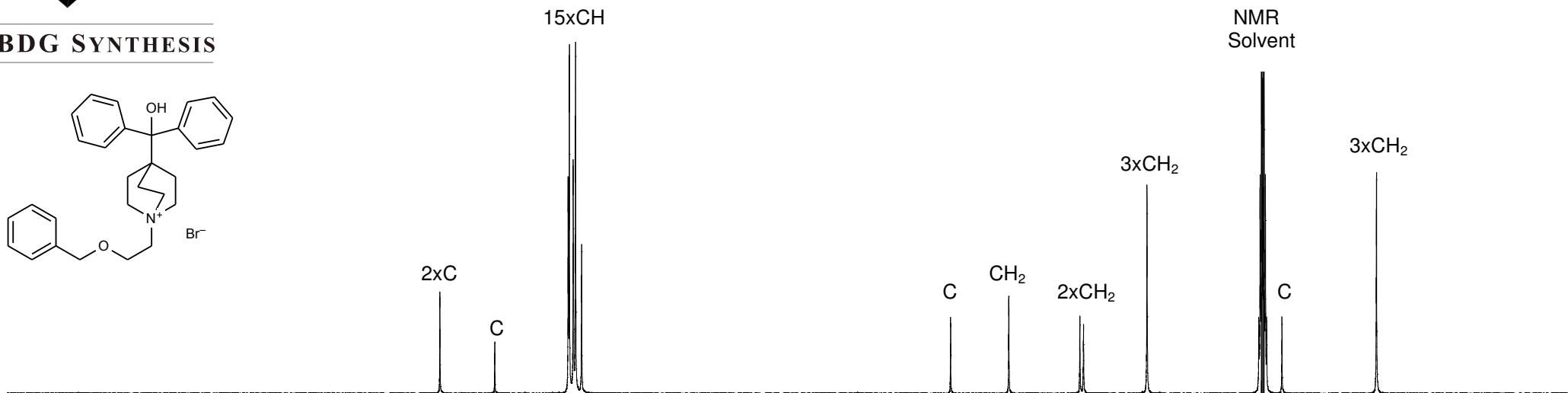
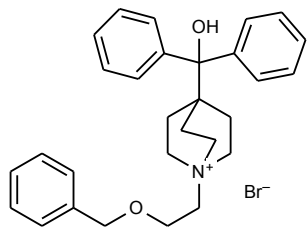
Lot Number: BDG 15055.2



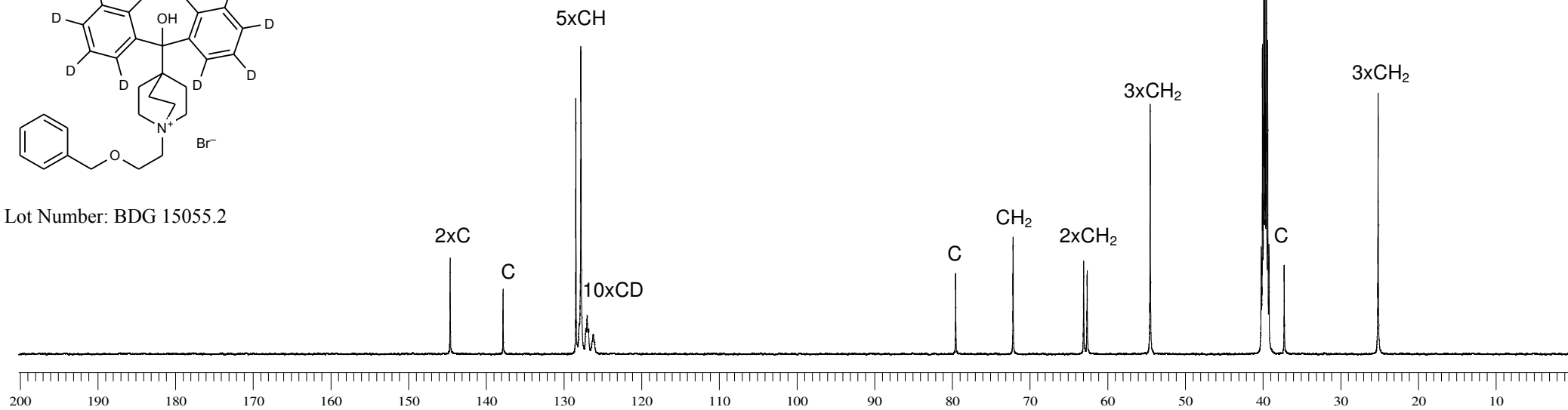


Carbon-13 NMR Spectrum of Umeclidinium Bromide (top) and Umeclidinium-d₁₀ Bromide (bottom) in DMSO-d₆

BDG SYNTHESIS



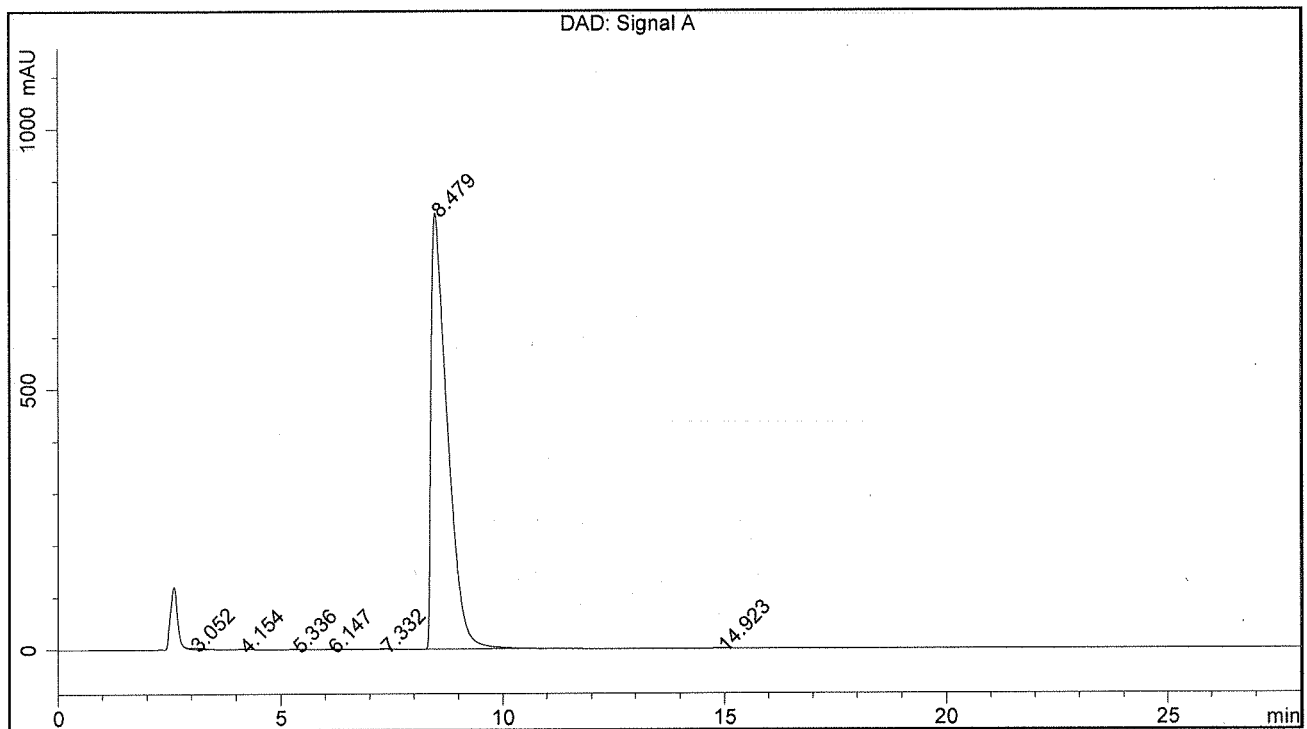
Lot Number: BDG 15055.2



BDG - Analysis of Umeclidinium-d10 Bromide

Column : Phenomenex Luna C18(2) 5um 250 x 4.6 mm
 Guard : Phenomenex Security Guard C18 RP 4 x 3 mm
 Mobile Phase : 55:45:0.05 Water : Acetonitrile : Trifluoroacetic Acid
 Flow Rate : 1.0 mL/min
 Sample Solvent : 70:30 Water : Acetonitrile
 Injection Volume : 10 uL
 Detection: UV 214 nm

Sample Name	BDG 15055.2	Instrument	AnalyticalLC01
Acquisition	23/07/2014, 16:41:27	Method (rev.)	LC10622a (10)
Sequence	BDG_23Jul2014b - Reprocessed	Vial Position	83
Operator	solvation010\cerityadmin	Injection	1 of 1



Area Percent Report

Peak#	RT	Peak Height	Peak Area	Width	Area %
1	3.05 min	3.0935	76.9800	0.3343 min	0.359 %
2	4.15 min	1.1453	15.9967	0.2189 min	0.075 %
3	5.34 min	1.1557	17.0763	0.2044 min	0.080 %
4	6.15 min	0.5637	8.9071	0.2044 min	0.041 %
5	7.33 min	1.3694	22.5132	0.2478 min	0.105 %
6	8.48 min	836.9988	21308.4227	0.3690 min	99.279 %
7	14.92 min	0.5175	13.2202	0.3100 min	0.062 %