

### 1. GENERAL INFORMATION

*IUPAC Name*: 3-(1,3-benzenodioxol-5-yl)-*N*,2-dimethylpropan-1-amine

**CFR:** Not Scheduled (as of 2/2013)

*CAS #:* N/A

*Synonyms:* 3,4-methylenedioxymethamphetamine methylene homolog,

3,4-methylenedioxymethamphetamine methyl homolog,

Heliomethylamine

Source: DEA Reference Material Collection

Appearance: White powder (HCl)

**Kovat's Index:** Pending

 $UV_{max}(nm)$ : Not Determined

### 2. CHEMICAL AND PHYSICAL DATA

# 2.1 CHEMICAL DATA

Form	Chemical Formula	Molecular Weight	Melting Point (°C)
Base	C <sub>12</sub> H <sub>17</sub> NO <sub>2</sub>	207	Not Determined
HC1	C <sub>12</sub> H <sub>17</sub> NO <sub>2</sub> ·HCl	243	144.9

### 3. ADDITIONAL RESOURCES

No resources identified as of 02/15/2013.

## 4. QUALITATIVE DATA

### 4.1 NUCLEAR MAGNETIC RESONANCE

## Method NMR D<sub>2</sub>O

Sample Preparation: Dilute analyte to ~5 mg/mL in D<sub>2</sub>O containing TSP for 0 ppm reference and maleic acid as quantitative internal standard.

Instrument: Varian Mercury 400 MHz NMR spectrometer with proton detection probe

**Parameters:** Spectral width: at least containing -3 ppm through 13 ppm

Pulse angle: 90°

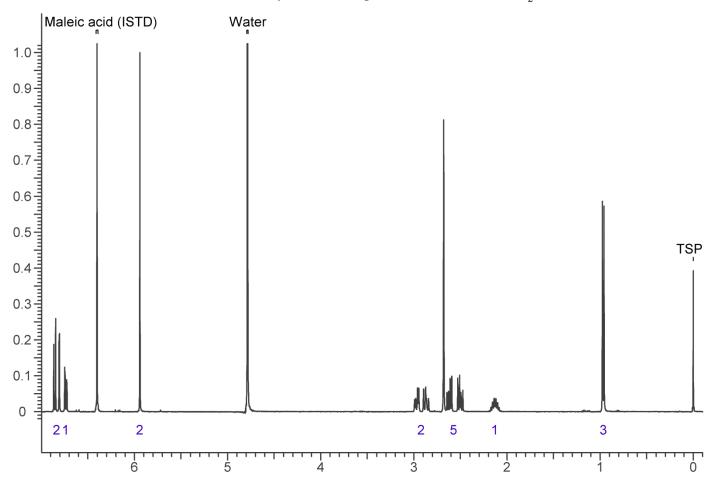
Delay between pulses: 45 seconds

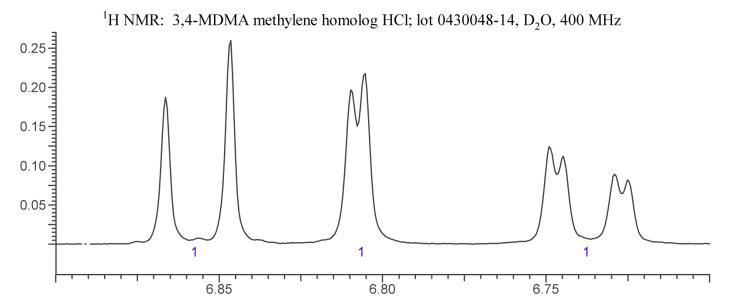
Number of scans (NT): 8 Number of steady state scans: 0 Oversampling: 4 or more

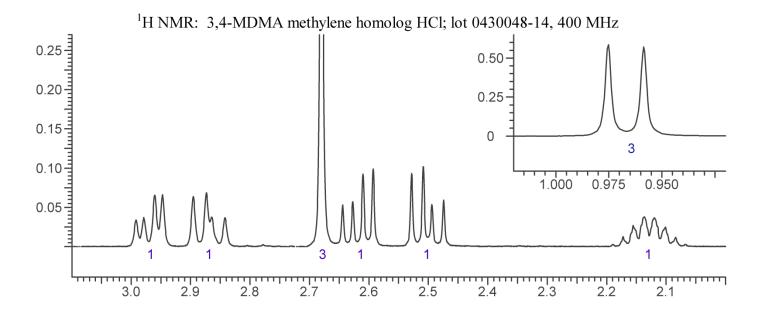
Shimming: automatic gradient shimming of Z1-4 shims

Phasing, Drift Correction: automatic or manual

<sup>1</sup>H NMR: 3,4-MDMA methylene homolog HCl; lot 0430048-14, D<sub>2</sub>O, 400 MHz







### 4.2 GAS CHROMATOGRAPHY/MASS SPECTROMETRY

Sample Preparation: Dilute analyte to ~4 mg/mL in chloroform.

Instrument: Agilent gas chromatograph operated in split mode with MS detector

**Column:** HP-1 MS; 30m x 0.25 mm x 0.25 μm

Carrier Gas: Helium at 1 mL/min Temperatures: Injector: 280°C

MSD transfer line: 280°C

MS Source: 230°C MS Quad: 150°C Oven program:

1) 100°C initial temperature for 1.0 min

2) Ramp to 300°C at 12°C/min

3) Hold final temperature for 9.0 min

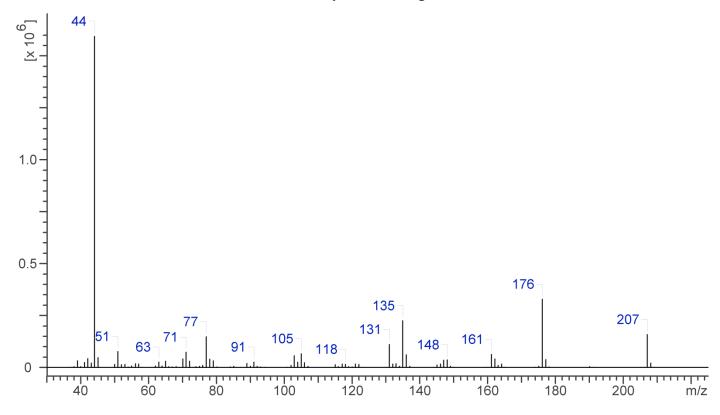
Injection Parameters: Split Ratio = 20:1, 1  $\mu$ L injected

MS Parameters: Mass scan range: 30-550 amu

Threshold: 90 Tune file: stune.u Acquisition mode: scan

**Retention Time:** 9.471 min

GC-MS: 3,4-MDMA methylene homolog HCl; lot 0430048-14



### 4.3 INFRARED SPECTROSCOPY (FTIR)

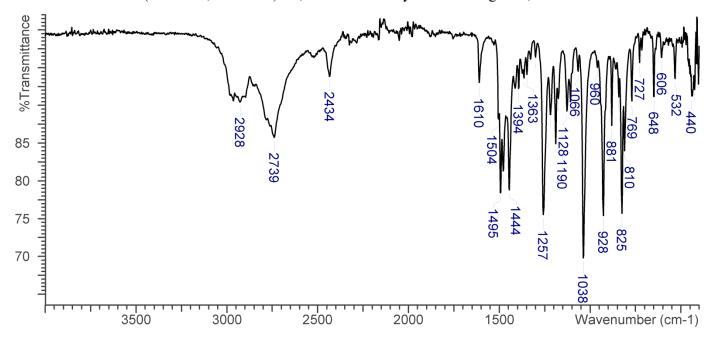
Instrument: FTIR with diamond ATR attachment (3 bounce)

Scan Parameters: Number of scans: 32

Number of background scans: 32

Resolution: 4cm<sup>-1</sup> Sample gain: 8 Aperture: 150

FTIR ATR (Diamond, 3 bounce): 3,4-MDMA methylene homolog HCl; lot 0430048-14



FTIR ATR (Diamond, 3 bounce): 3,4-MDMA methylene homolog HCl; lot 0430048-14

