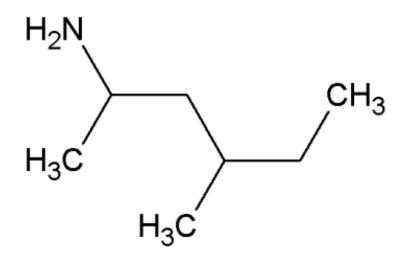
DMAA Latest Revision: April 19, 2013



## 1. GENERAL INFORMATION

*IUPAC Name*: 4-methylhexan-2-amine

CFR: Not Scheduled (4/2013)

*CAS #:* 105-41-9

*Synonyms:* 1,3-dimethylamylamine, methylhexanamine, 2-amino-4-methylhexane,

1,3-dimethylpentylamine, 4-methyl-2-hexylamine

Source: DEA Reference Material Collection

Appearance: White powder (HCl)

Kovat's Index: Pending

*UV<sub>max</sub>*: Not Determined

#### 2. CHEMICAL AND PHYSICAL DATA

## 2.1 CHEMICAL DATA

Form	Chemical Formula	Molecular Weight	Melting Point (°C)
Base	C <sub>7</sub> H <sub>17</sub> N	115	Not Determined
HCl	C <sub>7</sub> H <sub>17</sub> N HCl	151	122.9

#### 3. ADDITIONAL RESOURCES

## Wikipedia

# 4. QUALITATIVE DATA

## 4.1 NUCLEAR MAGNETIC RESONANCE

## Method NMR Dimethylfumarate/DMSO

Sample Preparation: Dilute analyte to ~10 mg/mL in DMSO containing TMS for 0 ppm reference and dimethylfumarate as quantitative internal standard.

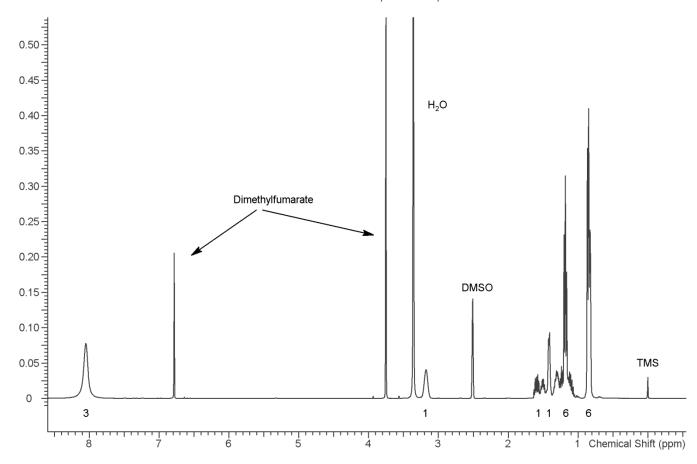
*Instrument:* 400 MHz NMR spectrometer

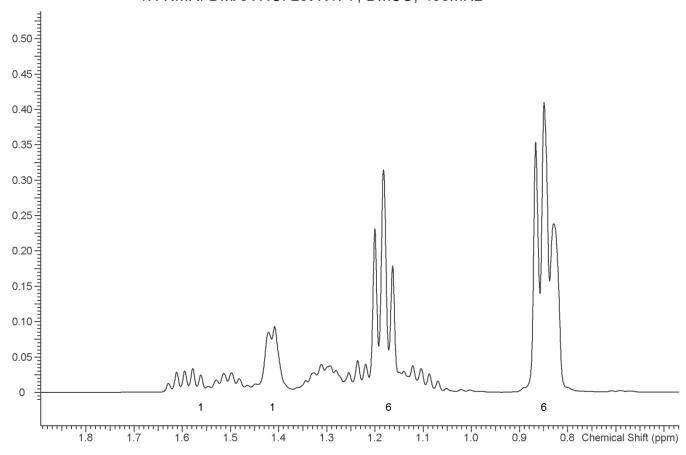
Parameters: Spectral width: at least containing -3 ppm through 13 ppm

Pulse angle: 90°

Delay between pulses: 45 seconds

# 1H NMR: DMAA HCI Lot N1P7; DMSO; 400MHz





## NMR Analytical Observation

DMAA has two chiral carbons; therefore, two diastereomers are possible. Diasteriomers produce slightly different chemical shifts for proton and carbon. Both diastereomers are present in the above spectra.

$$H_2N$$
 $H_3C$ 
 $H_3C$ 
 $H_3C$ 
 $H_3C$ 
 $H_3C$ 

#### 4.2 GAS CHROMATOGRAPHY/MASS SPECTROMETRY

Sample Preparation: Dilute analyte to ~1 mg/mL base extracted into chloroform

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

Column: DB-1 MS or equivalent; 30m x .25mm x .25μm

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

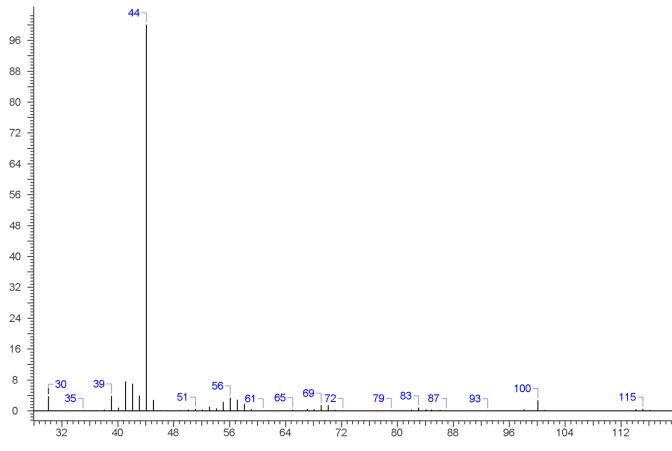
2) Ramp to 300°C at 12°C/min 3) Hold final temperature for 9.0 min

Injection Parameters:Split Ratio = 20:1, 1 μL injectedMS Parameters:Mass scan range: 30-550 amu

Threshold: 100 Tune file: stune.u Acquisition mode: scan

**Retention Time:** 2.100 minutes

El Mass Spectrum: DMAA HCI Lot N1P7



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# 4.3 INFRARED SPECTROSCOPY (FTIR)

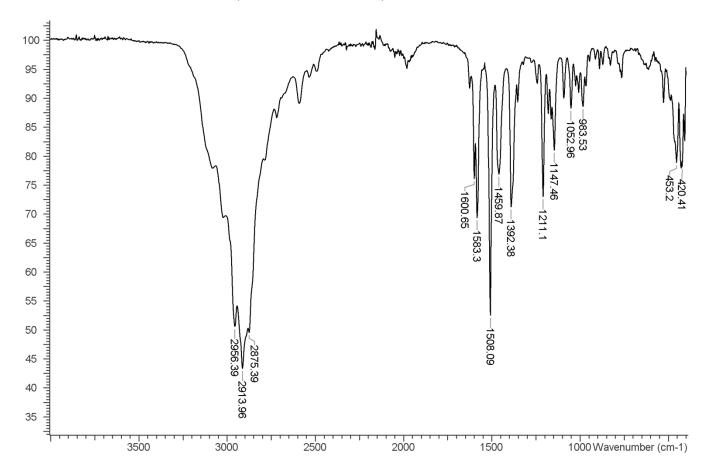
*Instrument:* FTIR with diamond ATR attachment (3 bounce)

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): DMAA HCI Lot N1P7



# FTIR ATR (Diamond, 3 Bounce): DMAA HCI Lot N1P7

