

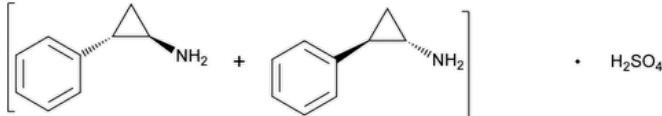
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Tranylcypromine Sulfate

Change to read:

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▲ (ERR 1-Jun-2020)

$(C_9H_{11}N)_2 \cdot H_2SO_4$ 364.46

Cyclopropanamine, 2-phenyl-, *trans*-(±)-, sulfate (2:1);
 (±)-*trans*-2-Phenylcyclopropylamine sulfate (2:1) CAS RN®: 13492-01-8; UNII: 7ZAT6ES870.

DEFINITION

Tranylcypromine Sulfate contains NLT 98.0% and NMT 102.0% of $(C_9H_{11}N)_2 \cdot H_2SO_4$, calculated on the dried basis.

IDENTIFICATION

- A. [SPECTROSCOPIC IDENTIFICATION TESTS \(197\), Infrared Spectroscopy: 197K](#)
- B. The retention time of the major peak of the *Sample* solution corresponds to that of the *Standard* solution, as obtained in the Assay.
- C. [IDENTIFICATION TESTS—GENERAL, Sulfate\(191\)](#): Meets the requirements

ASSAY

• PROCEDURE

Buffer: Dissolve 3.4 g of monobasic ammonium phosphate in about 900 mL of water in a 1-L volumetric flask. Adjust with phosphoric acid to a pH of 2.2 ± 0.1 , and dilute with water to volume.

Mobile phase: Methanol and *Buffer* (3:7)

0.05 N sulfuric acid: Cautiously add 1.3 mL of sulfuric acid to 100 mL of water, cool to room temperature, and dilute to 1000 mL.

Diluent: Methanol, water, and 0.05 N sulfuric acid (1:3:1)

Standard stock solution: Using a sonicator, dissolve [USP Tranylcypromine Sulfate RS](#) in 0.05 N sulfuric acid and methanol (about 30% of the final volume of each solvent). Dilute with *Diluent* to obtain a 400 μ g/mL solution.

Standard solution: 40 μ g/mL of [USP Tranylcypromine Sulfate RS](#) in *Diluent*, prepared from the *Standard stock solution*

Sample stock solution: Using a sonicator, dissolve Tranylcypromine Sulfate in methanol and 0.05 N sulfuric acid (about 30% of the final volume of each solvent). Dilute with *Diluent* to obtain a 400 μ g/mL solution.

Sample solution: 40 μ g/mL tranylcypromine sulfate in *Diluent*, prepared from the *Sample stock solution*

Chromatographic system

(See [Chromatography \(621\), System Suitability](#).)

Mode: LC

Detector: UV 220 nm

Column: 4.6-mm \times 25-cm; 4- μ m packing L11

Column temperature: 30°

Flow rate: 1 mL/min

Injection size: 20 μ L

System suitability

Sample: *Standard* solution

Suitability requirements

Tailing factor: NMT 2.0

Relative standard deviation: NMT 2.0%

Analysis

Samples: Standard solution and Sample solution

Calculate the percentage of $(C_9H_{11}N)_2 \cdot H_2SO_4$ in the portion taken:

$$\text{Result} = (r_u/r_s) \times (C_s/C_u) \times 100$$

r_u = peak response from the Sample solution

r_s = peak response from the Standard solution

C_s = concentration of [USP Tranylcypromine Sulfate RS](#) in the Standard solution (μg/mL)

C_u = concentration of tranylcypromine sulfate in the Sample solution (μg/mL)

Acceptance criteria: 98.0%–102.0% on the dried basis

IMPURITIES**INORGANIC IMPURITIES**

- [RESIDUE ON IGNITION \(281\)](#): NMT 0.1%

ORGANIC IMPURITIES

- **PROCEDURE**

Buffer, 0.05 N sulfuric acid, and Diluent: Proceed as directed in the Assay.

Solution A: Methanol and Buffer (3:17)

Solution B: Methanol and Buffer (3:7)

Mobile phase: See the gradient table below.

Time (min)	Solution A (%)	Solution B (%)
0	100	0
20	100	0
25	0	100
37	0	100
39	100	0
45	100	0

Standard stock solution: 14 μg/mL of [USP Tranylcypromine Sulfate RS](#), and 60 μg/mL each of [USP Tranylcypromine Related Compound A RS](#) and [USP Tranylcypromine Related Compound B RS](#) in Diluent. [NOTE—Sonicate as needed.]

Standard solution: Transfer a portion of the Standard stock solution to a suitable volumetric flask containing methanol and 0.05 N sulfuric acid (30% of the final volume of each solvent). Dilute with Diluent to volume to obtain a solution containing 0.7 μg/mL of [USP Tranylcypromine Sulfate RS](#) and 3.0 μg/mL each of [USP Tranylcypromine Related Compound A RS](#) and [USP Tranylcypromine Related Compound B RS](#).

Sample solution: Using a sonicator, dissolve Tranylcypromine Sulfate in methanol and 0.05 N sulfuric acid (about 30% of the final volume of each solvent). Dilute with Diluent to obtain a solution containing 680 μg/mL of tranylcypromine sulfate.

Chromatographic system

(See [Chromatography \(621\), System Suitability](#).)

Mode: LC

Detector: UV 249 nm for tranylcypromine related compound B; 220 nm for all other specified and unspecified impurities

Column: 4.6-mm × 15-cm; 3-μm packing L1

Column temperature: 35°

Flow rate: 1.2 mL/min

Injection size: 25 μL

System suitability

Sample: Standard solution

Suitability requirements

Resolution: NLT 2.0 between tranylcypromine and tranylcypromine related compound A, determined at 220 nm

Tailing factor: NMT 2.0 for tranylcypromine, determined at 220 nm

Relative standard deviation: NMT 6.0% for tranylcypromine and tranylcypromine related compound A, determined at 220 nm; NMT 6.0% for tranylcypromine related compound B, determined at 249 nm.

Analysis

Samples: Standard solution and Sample solution

[NOTE—Disregard the signal for tranylcypromine related compound B observed at 220 nm.]

Identify tranylcypromine related compound A, *cis*-hydrazide, *trans*-hydrazide, and any unspecified impurity using the data in [Impurity](#).

[Table 1](#). Calculate the percentage of each impurity in the portion of sample taken:

$$\text{Result} = (r_U/r_S) \times (C_S/C_U) \times 100/F$$

r_U = peak response of the impurity from the *Sample solution*

r_S = peak response of the *Standard solution*

C_S = concentration of [USP Tranylcypromine Sulfate RS](#) in the *Standard solution* ($\mu\text{g/mL}$)

C_U = concentration of Tranylcypromine Sulfate in the *Sample solution* ($\mu\text{g/mL}$)

F = response factor relative to tranylcypromine sulfate

Calculate the percentage of tranylcypromine related compound B in the portion of sample taken:

$$\text{Result} = (r_U/r_S) \times (C_S/C_U) \times (M_{r1}/M_{r2}) \times 100$$

r_U = peak response at 249 nm of tranylcypromine related compound B from the *Sample solution*

r_S = peak response at 249 nm of tranylcypromine related compound B from the *Standard solution*

C_S = concentration of [USP Tranylcypromine Related Compound B RS](#) in the *Standard solution* ($\mu\text{g/mL}$)

C_U = concentration of Tranylcypromine Sulfate in the *Sample solution* ($\mu\text{g/mL}$)

M_{r1} = molecular weight of 3-phenylallylamine free base, 133.19

M_{r2} = molecular weight of 3-phenylallylamine hydrochloride ([USP Tranylcypromine Related Compound B RS](#)), 169.66

Acceptance criteria

Individual impurities: See [Impurity Table 1](#). [NOTE—Tranylcypromine related compound B is determined at 249 nm. All other specified and unspecified impurities are determined at 220 nm.]

Total impurities: NMT 1.0% for sum of tranylcypromine related compound A, tranylcypromine related compound B, *cis*-hydrazide, *trans*-hydrazide, and any unspecified impurity.

Impurity Table 1

Name	Relative Retention Time	Relative Response Factor	Acceptance Criteria, NMT (%)
Tranylcypromine related compound A ^a	0.82	0.74	0.50
Tranylcypromine sulfate	1.0	1.0	—
Tranylcypromine related compound B ^b	1.2	—	0.50
<i>cis</i> -Hydrazide ^c	1.4	0.86	0.50

Name	Relative Retention Time	Relative Response Factor	Acceptance Criteria, NMT (%)
trans-Hydrazide ^d	2.5	1.1	0.50
Any unspecified impurity	—	1.0	0.10

^a (±)-*cis*-2-Phenylcyclopropanamine (*cis*-Cypromine).

^b 3-Phenylallylamine (Cinnamylamine), monitored at 249 nm.

^c (±)-*cis*-2-Phenylcyclopropanecarbohydrazide.

^d (±)-*trans*-2-Phenylcyclopropanecarbohydrazide.

SPECIFIC TESTS

- **Loss on Drying (731):** Dry a sample in vacuum at 60° for 2 h: it loses NMT 0.5% of its weight.

ADDITIONAL REQUIREMENTS

- **PACKAGING AND STORAGE:** Preserve in well-closed containers.

- **USP REFERENCE STANDARDS (11):**

[USP Tranylcypromine Sulfate RS](#)

[USP Tranylcypromine Related Compound A RS](#)

(±)-*cis*-2-Phenylcyclopropanamine hydrochloride; *cis*-tranylcypromine hydrochloride.

C₉H₁₁N · HCl 169.65

[USP Tranylcypromine Related Compound B RS](#)

3-Phenylallylamine hydrochloride; cinnamylamine hydrochloride.

C₉H₁₁N · HCl 169.65

Auxiliary Information - Please [check for your question in the FAQs](#) before contacting USP.

Topic/Question	Contact	Expert Committee
TRANYLCYPROMINE SULFATE	Documentary Standards Support	SM42020 Small Molecules 4
REFERENCE STANDARD SUPPORT	RS Technical Services RSTECH@usp.org	SM42020 Small Molecules 4

Chromatographic Database Information: [Chromatographic Database](#)

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