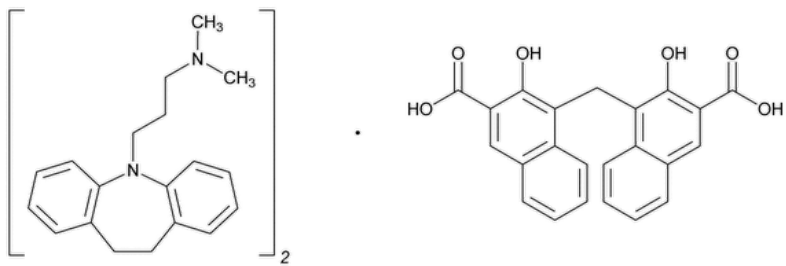


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Imipramine Pamoate



$(C_{19}H_{24}N_2)_2 \cdot C_{23}H_{16}O_6$ 949.18
Naphthalenecarboxylic acid, 4,4'-methylenebis[3-hydroxy-, compd. with 5H-dibenz[b,f]azepine-5-propanamine, 10,11-dihydro-N,N-dimethyl-, 2- (1:2);
5-[3-(Dimethylamino)propyl]-10,11-dihydro-5H-dibenz[b,f]azepine compound (2:1) with 4,4'-methylenebis(3-hydroxy-2-naphthoic acid);
3-[10,11-Dihydro-5H-dibenzo[b,f]azepin-5-yl)-N,N-dimethylpropan-1-amine 4,4'-methylenebis(3-hydroxy-2-naphthoate) salt (2:1) CAS RN®: 10075-24-8; UNII: MC34P30298.

DEFINITION
Imipramine Pamoate contains NLT 98.0% and NMT 102.0% of imipramine pamoate $[(C_{19}H_{24}N_2)_2 \cdot C_{23}H_{16}O_6]$, calculated on anhydrous basis.

IDENTIFICATION
Change to read:
• **A.** **SPECTROSCOPIC IDENTIFICATION TESTS (197), Infrared Spectroscopy: 197K** (CN 1-MAY-2020)
• **B.** The retention times of the major peaks of the *Sample solution* correspond to those of the *Standard solution*, as obtained in the Assay.

ASSAY
• **PROCEDURE**
Buffer: 5.2 g/L of dibasic potassium phosphate in water
Solution A: Acetonitrile and *Buffer* (15:85). Adjust with phosphoric acid to a pH of 8.0.
Solution B: Acetonitrile and *Buffer* (38:62). Adjust with phosphoric acid to a pH of 8.0.
Mobile phase: See [Table 1](#).

Table 1

Time (min)	Solution A (%)	Solution B (%)
0	90	10
10	70	30
20	35	65
30	35	65
31	90	10
35	90	10

Diluent: Acetonitrile and water (75:25)

Standard stock solution: 2 mg/mL of [USP Imipramine Pamoate RS](#) in *Diluent*

Standard solution: 0.2 mg/mL of [USP Imipramine Pamoate RS](#) from *Standard stock solution* in *Solution A*

Sample stock solution: 2 mg/mL of Imipramine Pamoate in *Diluent*

Sample solution: 0.2 mg/mL of Imipramine Pamoate from *Sample stock solution* in *Solution A*

Chromatographic system

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

Mode: LC

Detector: UV 269 nm

Column: 4.6-mm × 15-cm; 5-μm packing L1

Autosampler temperature: 10°

Flow rate: 1.5 mL/min

Injection volume: 20 μL

System suitability

Sample: *Standard solution*

[NOTE—Approximate relative retention times for pamoic acid and imipramine are 0.3 and 1.0, respectively.]

Suitability requirements

Resolution: NLT 2.0 between pamoic acid and imipramine

Tailing factor: NMT 1.8 for imipramine

Relative standard deviation: NMT 2.0% for imipramine

Analysis

Samples: *Standard solution* and *Sample solution*

Calculate the percentage of imipramine pamoate $[(C_{19}H_{24}N_2)_2 \cdot C_{23}H_{16}O_6]$ in the portion of Imipramine Pamoate taken:

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

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

r_S = peak response of imipramine from the *Standard solution*

C_S = concentration of [USP Imipramine Pamoate RS](#) in the *Standard solution* (mg/mL)

C_U = concentration of Imipramine Pamoate in the *Sample solution* (mg/mL)

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

IMPURITIES

• [RESIDUE ON IGNITION \(281\)](#): NMT 0.10%

• ORGANIC IMPURITIES

Buffer: 5.2 g/L of dibasic potassium phosphate in water

Solution A: Acetonitrile and *Buffer* (38:62). Adjust with phosphoric acid to a pH of 7.9.

Solution B: Acetonitrile and *Buffer* (70:30). Adjust with phosphoric acid to a pH of 6.0.

Mobile phase: See [Table 2](#).

Table 2

Time (min)	Solution A (%)	Solution B (%)
0	100	0
15	100	0
50	0	100
60	0	100
61	100	0

Time (min)	Solution A (%)	Solution B (%)
75	100	0

Diluent: Acetonitrile and water (75:25)

System suitability solution: 0.5 mg/mL of [USP Imipramine Pamoate RS](#), 0.02 mg/mL of [USP Depramine RS](#), and 0.02 mg/mL [USP Iminodibenzyl RS](#) in *Diluent*

Standard solution: 0.002 mg/mL of [USP Imipramine Pamoate RS](#) in *Diluent*

Sample solution: 2 mg/mL of Imipramine Pamoate in *Diluent*

Chromatographic system

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

Mode: LC

Detector: UV 220 nm

Column: 4.6-mm × 15-cm; 5-μm packing L1

Temperatures

Column: 40°

Autosampler: 10°

Flow rate: 1.0 mL/min

Injection volume: 10 μL

System suitability

Samples: *System suitability solution* and *Standard solution*

[NOTE—See [Table 3](#) for relative retention times.]

Suitability requirements

Resolution: NLT 5.0 between imipramine and depramine, *System suitability solution*

Relative standard deviation: NMT 5.0%, *Standard solution*

Analysis

Samples: *Standard solution* and *Sample solution*

Calculate the percentage of each impurity in the portion of Imipramine Pamoate taken:

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

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

r_S = peak response of imipramine from the *Standard solution*

C_S = concentration of [USP Imipramine Pamoate RS](#) in the *Standard solution* (mg/mL)

C_U = concentration of Imipramine Pamoate in the *Sample solution* (mg/mL)

Acceptance criteria: See [Table 3](#). Disregard peaks that are less than 0.05% of the imipramine peak.

Table 3

Name	Relative Retention Time	Acceptance Criteria, NMT (%)
Pamoic acid (Pamoate) ^a	0.2	—
Desipramine ^b	0.5	0.10
Depramine	0.7	0.10
Imipramine	1.0	—
Iminodibenzyl	3.7	0.10

Name	Relative Retention Time	Acceptance Criteria, NMT (%)
Any individual, unspecified impurity	—	0.10
Total impurities	—	0.75

^a Pamoic acid is not an impurity. It is listed for identification purposes only.

^b 10,11-Dihydro-5-[3-(methylamino)propyl]-5*H*-dibenz[*b,f*]azepine.

SPECIFIC TESTS

- [WATER DETERMINATION, Method I \(921\)](#): NMT 2.0%

ADDITIONAL REQUIREMENTS

- **PACKAGING AND STORAGE:** Preserve in tight containers, and store at controlled room temperature.
- [USP REFERENCE STANDARDS \(11\)](#).

[USP Depramine RS](#)
 3-(5*H*-Dibenzo[*b,f*]azepin-5-yl)-*N,N*-dimethylpropan-1-amine.
 C₁₉H₂₂N₂ 278.39

[USP Iminodibenzyl RS](#)
 10,11-Dihydro-5*H*-dibenzo[*b,f*]azepine.
 C₁₄H₁₃N 195.28

[USP Imipramine Pamoate RS](#)

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

Topic/Question	Contact	Expert Committee
IMIPRAMINE PAMOATE	Documentary Standards Support	SM42020 Small Molecules 4

Chromatographic Database Information: [Chromatographic Database](#)

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