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## Pyridostigmine Bromide



$C_9H_{13}BrN_2O_2$  261.12

Pyridinium, 3-[(dimethylamino)carbonyl]oxy]-1-methyl-, bromide;

3-Hydroxy-1-methylpyridinium bromide dimethylcarbamate CAS RN®: 101-26-8; UNII: KVI301NA53.

### DEFINITION

Pyridostigmine Bromide contains NLT 98.0% and NMT 102.0% of pyridostigmine bromide ( $C_9H_{13}BrN_2O_2$ ), calculated on the dried basis.

### IDENTIFICATION

*Change to read:*

- A. **SPECTROSCOPIC IDENTIFICATION TESTS (197), Infrared Spectroscopy:** 197A or 197K▲ (CN 1-May-2020)
- B. The UV spectrum of the major peak of the *Sample solution* corresponds to that of the *Standard solution*, as obtained in the Assay.
- C. **IDENTIFICATION TESTS—GENERAL (191), Chemical Identification Tests, Bromide**

**Sample solution:** 20 mg/mL of Pyridostigmine Bromide

**Acceptance criteria:** Meets the requirements

### ASSAY

#### • PROCEDURE

**Buffer:** 4.3 g/L of sodium dodecyl sulfate in water. Adjust with phosphoric acid to a pH of 2.0.

**Mobile phase:** Acetonitrile and Buffer (32:68)

**System suitability solution:** 2  $\mu$ g/mL each of USP Pyridostigmine RS, USP Pyridostigmine Related Compound A RS, and USP Pyridostigmine Related Compound B RS in *Mobile phase*

**Standard solution:** 0.3 mg/mL of USP Pyridostigmine RS in *Mobile phase*

**Sample solution:** 0.3 mg/mL of Pyridostigmine Bromide in *Mobile phase*

#### Chromatographic system

(See Chromatography (621), System Suitability.)

**Mode:** LC

**Detector:** UV 220 nm. For *Identification B*, use a diode array detector in the range of 200–400 nm.

**Column:** 4.6-mm  $\times$  15-cm; 5- $\mu$ m packing L1

**Flow rate:** 1 mL/min

**Injection volume:** 20  $\mu$ L

**Run time:** NLT 2.6 times the retention time of pyridostigmine

#### System suitability

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

#### Suitability requirements

**Resolution:** NLT 1.5 between pyridostigmine related compound A and pyridostigmine, *System suitability solution*

**Tailing factor:** NMT 2.0, *Standard solution*

**Relative standard deviation:** NMT 0.73%, *Standard solution*

#### Analysis

**Samples:** *Standard solution* and *Sample solution*

Calculate the percentage of pyridostigmine bromide ( $C_9H_{13}BrN_2O_2$ ) in the portion of Pyridostigmine Bromide 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 Pyridostigmine Bromide RS](#) in the *Standard solution* (mg/mL) $C_u$  = concentration of Pyridostigmine Bromide in the *Sample solution* (mg/mL)**Acceptance criteria:** 98.0%–102.0% on the dried basis

## IMPURITIES

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

- **ORGANIC IMPURITIES**

**Mobile phase and Chromatographic system:** Proceed as directed in the Assay.**Sensitivity solution:** 0.2  $\mu$ g/mL of [USP Pyridostigmine Bromide RS](#) in *Mobile phase***Standard solution:** 0.6  $\mu$ g/mL each of [USP Pyridostigmine Bromide RS](#), [USP Pyridostigmine Related Compound A RS](#), and [USP Pyridostigmine Related Compound B RS](#) in *Mobile phase***Sample solution:** 1000  $\mu$ g/mL of Pyridostigmine Bromide in *Mobile phase***System suitability****Samples:** *Sensitivity solution* and *Standard solution***Suitability requirements****Resolution:** NLT 1.5 between pyridostigmine related compound A and pyridostigmine, *Standard solution***Relative standard deviation:** NMT 5.0% for all the three peaks, *Standard solution***Signal-to-noise ratio:** NLT 10, *Sensitivity solution***Analysis****Samples:** *Standard solution* and *Sample solution*

Calculate the percentage of pyridostigmine related compound A and pyridostigmine related compound B in the portion of Pyridostigmine Bromide taken:

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

 $r_u$  = peak response of pyridostigmine related compound A or pyridostigmine related compound B from the *Sample solution* $r_s$  = peak response of pyridostigmine related compound A or pyridostigmine related compound B from the *Standard solution* $C_s$  = concentration of [USP Pyridostigmine Related Compound A RS](#) or [USP Pyridostigmine Related Compound B RS](#) in the *Standard solution* ( $\mu$ g/mL) $C_u$  = concentration of Pyridostigmine Bromide in the *Sample solution* ( $\mu$ g/mL)

Calculate the percentage of each unspecified impurity in the portion of Pyridostigmine Bromide 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 pyridostigmine from the *Standard solution* $C_s$  = concentration of [USP Pyridostigmine Bromide RS](#) in the *Standard solution* ( $\mu$ g/mL) $C_u$  = concentration of Pyridostigmine Bromide in the *Sample solution* ( $\mu$ g/mL)**Acceptance criteria:** See [Table 1](#). The reporting threshold is 0.02%.**Table 1**

Name	Relative Retention Time	Acceptance Criteria, NMT (%)
Pyridostigmine related compound B	0.78	0.06
Pyridostigmine related compound A	0.92	0.06
Pyridostigmine	1.0	—
Any individual unspecified impurity	—	0.06
Total impurities	—	0.5

**SPECIFIC TESTS**

- [Loss on Drying \(731\)](#).

**Analysis:** Dry at 100° for 4 h in a suitable vacuum drying tube, using phosphorus pentoxide as the desiccant.

**Acceptance criteria:** NMT 2.0%

**ADDITIONAL REQUIREMENTS**

- **PACKAGING AND STORAGE:** Preserve in tight containers.

- [USP Reference Standards \(11\)](#).

[USP Pyridostigmine RS](#)

[USP Pyridostigmine Related Compound A RS](#)

Pyridin-3-yl dimethylcarbamate.

$C_8H_{10}N_2O_2$  166.18

[USP Pyridostigmine Related Compound B RS](#)

3-Hydroxy-1-methylpyridin-1-ium bromide.

$C_6H_8BrNO$  190.04

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

Topic/Question	Contact	Expert Committee
PYRIDOSTIGMINE BROMIDE	<a href="#">Documentary Standards Support</a>	SM42020 Small Molecules 4
REFERENCE STANDARD SUPPORT	RS Technical Services <a href="mailto:RSTECH@usp.org">RSTECH@usp.org</a>	SM42020 Small Molecules 4

**Chromatographic Database Information:** [Chromatographic Database](#)

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