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## Xylometazoline Hydrochloride Nasal Solution

### DEFINITION

Xylometazoline Hydrochloride Nasal Solution is an isotonic solution of Xylometazoline Hydrochloride in water. It contains NLT 90.0% and NMT 110.0% of the labeled amount of xylometazoline hydrochloride ( $C_{16}H_{24}N_2 \cdot HCl$ ).

### IDENTIFICATION

#### Change to read:

• **A.** ▲The UV spectrum of the xylometazoline peak of the *Sample solution* corresponds to that of the *Standard solution*, as obtained in the *Assay*.▲ (USP 1-Aug-2019)

#### Add the following:

▲• **B.** The retention time of the major peak of the *Sample solution* corresponds to that of the *Standard solution*, as obtained in the *Assay*.▲ (USP 1-Aug-2019)

### ASSAY

#### Change to read:

#### • PROCEDURE

▲**Solution A:** 1.4 g/L of [monobasic potassium phosphate](#). Adjust with [phosphoric acid](#) to a pH of 3.0.

**Solution B:** Acetonitrile

**Mobile phase:** See [Table 1](#).

Table 1

Time (min)	Solution A (%)	Solution B (%)
0	70	30
5	70	30
20	15	85
35	15	85
37	70	30
45	70	30

**System suitability solution:** 0.01 mg/mL each of [USP Xylometazoline Hydrochloride RS](#) and [USP Xylometazoline Related Compound A RS](#) in [water](#)

**Standard solution:** 0.05 mg/mL of [USP Xylometazoline Hydrochloride RS](#) in [water](#)

**Sample solution:** Nominally 0.05 mg/mL of xylometazoline hydrochloride from a suitable volume of Nasal Solution in [water](#)

#### Chromatographic system

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

**Mode:** LC

**Detector:** UV 220 nm. For *Identification A*, use a diode array detector in the range of 210–300 nm.

**Column:** 4.6-mm × 25-cm; 5-μm packing [L1](#)

**Flow rate:** 1 mL/min**Injection volume:** 20 µL**System suitability****Samples:** *System suitability solution* and *Standard solution*[NOTE—The relative retention times are given in [Table 2](#).]**Suitability requirements****Resolution:** NLT 2.0 between xylometazoline related compound A and xylometazoline, *System suitability solution***Tailing factor:** NMT 2.0, *Standard solution***Relative standard deviation:** NMT 1.0%, *Standard solution***Analysis****Samples:** *Standard solution* and *Sample solution*Calculate the percentage of the labeled amount of xylometazoline hydrochloride ( $C_{16}H_{24}N_2 \cdot HCl$ ) in the portion of Nasal Solution taken:

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

 $r_U$  = peak response of xylometazoline from the *Sample solution* $r_S$  = peak response of xylometazoline from the *Standard solution* $C_S$  = concentration of [USP Xylometazoline Hydrochloride RS](#) in the *Standard solution* (mg/mL) $C_U$  = nominal concentration of xylometazoline hydrochloride in the *Sample solution* (mg/mL)

▲ (USP 1-Aug-2019)

**Acceptance criteria:** 90.0%–110.0%**IMPURITIES****Add the following:**

## ▲ • ORGANIC IMPURITIES

**Solution A, Solution B, Mobile phase, System suitability solution, and Chromatographic system:** Proceed as directed in the Assay.**Standard solution:** 0.001 mg/mL of [USP Xylometazoline Hydrochloride RS](#) in [water](#)**Sensitivity solution:** 0.5 µg/mL of [USP Xylometazoline Hydrochloride RS](#) from the *Standard solution* in [water](#)**Sample solution:** Nominally 0.5 mg/mL of xylometazoline hydrochloride from a suitable volume of Nasal Solution in [water](#)**System suitability****Samples:** *System suitability solution*, *Standard solution*, and *Sensitivity solution*[NOTE—The relative retention times are given in [Table 2](#).]**Suitability requirements****Resolution:** NLT 2.0 between xylometazoline related compound A and xylometazoline, *System suitability solution***Tailing factor:** NMT 2.0, *Standard solution***Relative standard deviation:** NMT 5.0%, *Standard solution***Signal-to-noise ratio:** NLT 10, *Sensitivity solution***Analysis****Samples:** *Standard solution* and *Sample solution*

Calculate the percentage of each degradation product in the portion of Nasal Solution taken:

$$\text{Result} = (r_U/r_S) \times (C_S/C_U) \times (1/F) \times 100$$

 $r_U$  = peak response of each degradation product from the *Sample solution* $r_S$  = peak response of xylometazoline from the *Standard solution* $C_S$  = concentration of [USP Xylometazoline Hydrochloride RS](#) in the *Standard solution* (mg/mL) $C_U$  = nominal concentration of xylometazoline hydrochloride in the *Sample solution* (mg/mL) $F$  = relative response factor for the corresponding degradation product (see [Table 2](#))**Acceptance criteria:** See [Table 2](#). The reporting threshold is 0.10%.**Table 2**

Name	Relative Retention Time	Relative Response Factor	Acceptance Criteria, NMT (%)
Xylometazoline related compound A	0.79	0.67	0.2
Xylometazoline	1.0	—	—
Any individual unspecified degradation product	—	1.0	0.2
Total degradation products	—	—	0.5▲ (USP 1-Aug-2019)

**SPECIFIC TESTS**

- **pH (791):** 5.0–7.5

**Add the following:**

▲ **MICROBIAL ENUMERATION TESTS (61)** and **TESTS FOR SPECIFIED MICROORGANISMS (62)**: The total aerobic microbial count does not exceed  $10^2$  cfu/g, and the total combined molds and yeasts count does not exceed  $10^1$  cfu/g. It meets the requirements of the tests for the absence of *Staphylococcus aureus* and *Pseudomonas aeruginosa*.▲ (USP 1-Aug-2019)

**ADDITIONAL REQUIREMENTS****Change to read:**

- **PACKAGING AND STORAGE:** Preserve in tight, light-resistant containers. ▲Store at controlled room temperature.▲ (USP 1-Aug-2019)

**Change to read:**

- **USP REFERENCE STANDARDS (11)**

[USP Xylometazoline Hydrochloride RS](#)

- ▲ [USP Xylometazoline Related Compound A RS](#)

N-(2-Aminoethyl)-2-[4-(tert-butyl)-2,6-dimethylphenyl]acetamide.

$C_{16}H_{26}N_2O$  262.39▲ (USP 1-Aug-2019)

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

Topic/Question	Contact	Expert Committee
XYLOMETAZOLINE HYDROCHLORIDE NASAL SOLUTION	<a href="#">Documentary Standards Support</a>	SM52020 Small Molecules 5
REFERENCE STANDARD SUPPORT	RS Technical Services <a href="mailto:RSTECH@usp.org">RSTECH@usp.org</a>	SM52020 Small Molecules 5

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

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