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# Propranolol Hydrochloride Injection

## DEFINITION

Propranolol Hydrochloride Injection is a sterile solution of Propranolol Hydrochloride in Water for Injection. It contains NLT 90.0% and NMT 110.0% of the labeled amount of propranolol hydrochloride ( $C_{16}H_{21}NO_2 \cdot HCl$ ).

## IDENTIFICATION

• **A.** The retention time of the major peak of the *Sample solution* corresponds to that of the *Standard solution*, as obtained in the Assay.

**Add the following:**

▲ • **B.** The UV spectrum of the major peak of the *Diluted sample solution* corresponds to that of the *Diluted standard solution*, as obtained in the Assay. ▲ (USP 1-Dec-2022)

## ASSAY

**Change to read:**

### PROCEDURE

▲ **Mobile phase:** Dissolve 1.6 g of [sodium dodecyl sulfate](#) and 0.3 g of [tetrabutylammonium phosphate](#) in a mixture consisting of 1 mL of [sulfuric acid](#), 450 mL of [water](#), and 550 mL of [acetonitrile](#). Adjust with 2 N [sodium hydroxide](#) solution to a pH of 3.3.

**Standard solution:** 0.1 mg/mL of [USP Propranolol Hydrochloride RS](#) in *Mobile phase*. Sonicate to dissolve, if necessary.

**Diluted standard solution:** 0.02 mg/mL of [USP Propranolol Hydrochloride RS](#) in *Mobile phase* from *Standard solution*

**Sample solution:** Nominally 0.1 mg/mL of propranolol hydrochloride in *Mobile phase* from a volume of the Injection

**Diluted sample solution:** Nominally 0.02 mg/mL of propranolol hydrochloride in *Mobile phase* from *Sample solution*

### Chromatographic system

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

**Mode:** LC

**Detector:** UV 292 nm. For *Identification B*, use a diode array detector in the range of 200–400 nm.

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

**Flow rate:** 1.8 mL/min

**Injection volume:** 50 μL

**Run time:** NLT 9 times the retention time of propranolol

### System suitability

**Sample:** *Standard solution*

#### Suitability requirements

**Tailing factor:** NMT 2.0

**Relative standard deviation:** NMT 1.0%

## Analysis

**Samples:** *Standard solution*, *Diluted standard solution*, *Sample solution*, and *Diluted sample solution*

[NOTE—The *Diluted standard solution* and *Diluted sample solution* are used for *Identification B*.]

Calculate the percentage of the labeled amount of propranolol hydrochloride ( $C_{16}H_{21}NO_2 \cdot HCl$ ) in the portion of Injection taken:

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

$r_U$  = peak response of propranolol from the *Sample solution*

$r_S$  = peak response of propranolol from the *Standard solution*

$C_S$  = concentration of [USP Propranolol Hydrochloride RS](#) in the *Standard solution* (mg/mL)

$C_U$  = nominal concentration of propranolol hydrochloride in the *Sample solution* (mg/mL)▲ (USP 1-Dec-2022)

**Acceptance criteria:** 90.0%–110.0%

**Add the following:**

#### ▲IMPURITIES

##### • ORGANIC IMPURITIES

**Mobile phase and Chromatographic system:** Proceed as directed in the Assay.

**System suitability solution:** 1 µg/mL of [USP Propranolol Related Compound A RS](#) and 500 µg/mL of [USP Propranolol Hydrochloride RS](#) in *Mobile phase*

**Standard solution:** 1 µg/mL of [USP Propranolol Hydrochloride RS](#) in *Mobile phase*

**Sensitivity solution:** 0.5 µg/mL of [USP Propranolol Hydrochloride RS](#) in *Mobile phase* from *Standard solution*

**Sample solution:** Nominally 500 µg/mL of propranolol hydrochloride in *Mobile phase* from a volume of the Injection.

#### System suitability

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

[NOTE—The relative retention times for propranolol related compound A and propranolol are 0.6 and 1.0, respectively.]

#### Suitability requirements

**Resolution:** NLT 3.0 between propranolol and propranolol related compound A, *System suitability 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 any unspecified degradation product in the portion of Injection taken:

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

$r_U$  = peak response of any unspecified degradation product from the *Sample solution*

$r_S$  = peak response of propranolol from the *Standard solution*

$C_S$  = concentration of [USP Propranolol Hydrochloride RS](#) in the *Standard solution* (µg/mL)

$C_U$  = nominal concentration of propranolol hydrochloride in the *Sample solution* (µg/mL)

**Acceptance criteria:** See [Table 1](#). The reporting threshold is 0.1%.

**Table 1**

Name	Relative Retention Time	Acceptance Criteria, NMT (%)
Any unspecified degradation product	—	0.2
Total degradation products	—	0.5▲ (USP 1-Dec-2022)

#### SPECIFIC TESTS

**Change to read:**

• [BACTERIAL ENDOTOXINS TEST \(85\)](#): ▲Meets the requirements▲ (USP 1-Dec-2022)

**Add the following:**

▲• [PARTICULATE MATTER IN INJECTIONS \(788\)](#): Meets the requirements▲ (USP 1-Dec-2022)

• [pH \(791\)](#): 2.8–4.0

**Add the following:**

▲• [STERILITY TESTS \(71\)](#): Meets the requirements▲ (USP 1-Dec-2022)

• **OTHER REQUIREMENTS:** It meets the requirements in [Injections and Implanted Drug Products \(1\)](#).

ADDITIONAL REQUIREMENTS

Change to read:

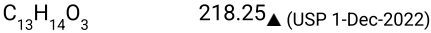
- **PACKAGING AND STORAGE:** Preserve in single-dose, light-resistant containers, preferably of Type I glass. ▲ Store at controlled room temperature. Protect from freezing or excessive heat. ▲ (USP 1-Dec-2022)

Change to read:

- **USP REFERENCE STANDARDS (11).**  
[USP Propranolol Hydrochloride RS](#)

▲ [USP Propranolol Related Compound A RS](#)

3-(Naphthalen-1-yloxy)propane-1,2-diol.



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

Topic/Question	Contact	Expert Committee
PROPRANOLOL HYDROCHLORIDE INJECTION	<a href="#">Documentary Standards Support</a>	SM22020 Small Molecules 2
REFERENCE STANDARD SUPPORT	RS Technical Services <a href="mailto:RSTECH@usp.org">RSTECH@usp.org</a>	SM22020 Small Molecules 2

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

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