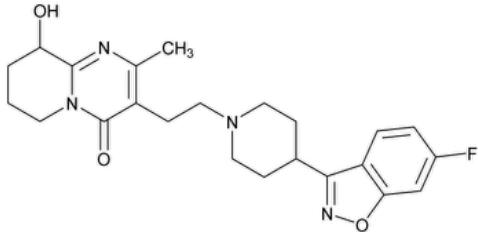


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Paliperidone



$C_{23}H_{27}FN_4O_3$ 426.48

4H-Pyrido[1,2-a]pyrimidin-4-one, 3-[2-[4-(6-fluoro-1,2-benzisoxazol-3-yl)-1-piperidinyl]ethyl]-6,7,8,9-tetrahydro-9-hydroxy-2-methyl-; (9RS)-3-[2-[4-(6-Fluoro-1,2-benzisoxazol-3-yl)piperidin-1-yl]ethyl]-9-hydroxy-2-methyl-6,7,8,9-tetrahydro-4H-pyrido[1,2-a]pyrimidin-4-one CAS RN®: 144598-75-4; UNII: 838F01T721.

DEFINITION

Paliperidone contains NLT 98.0% and NMT 102.0% of paliperidone ($C_{23}H_{27}FN_4O_3$).

IDENTIFICATION

Change to read:

- A. **SPECTROSCOPIC IDENTIFICATION TESTS (197), Infrared Spectroscopy: 197K** ▲ (CN 1-May-2020) Do not dry the sample.
- B. The retention time of the major peak of the *Sample solution* corresponds to that of the *Standard solution*, as obtained in the Assay.

ASSAY

• PROCEDURE

Protect the System suitability solution, Standard solution, and Sample solution from light.

Buffer: 28 mM Tetrabutylammonium hydrogen sulfate

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

Solution A: Methanol and Buffer (10:90)

Solution B: Methanol

Table 1

Time (min)	Solution A (%)	Solution B (%)
0	100	0
5	100	0
35	85	15
37	100	0
45	100	0

Diluent: Dissolve 0.71 g of dibasic sodium phosphate and 0.62 g of monobasic sodium phosphate in 1 L of water.

System suitability solution: 0.5 mg/mL of [USP Paliperidone Resolution Mixture RS](#) prepared as follows. Dissolve first in methanol using 50% of the final flask volume, and dilute with *Diluent*.

Standard solution: 0.5 mg/mL of [USP Paliperidone RS](#) prepared as follows. Dissolve first in methanol using 50% of the final flask volume, and dilute with *Diluent*.

Sample solution: 0.5 mg/mL of Paliperidone prepared as follows. Dissolve first in methanol using 50% of the final flask volume, and dilute with *Diluent*.

Chromatographic system

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

Mode: LC

Detector: UV 275 nm

Column: 4.6-mm × 10-cm; 3-μm packing L1

Column temperature: 40°

Flow rate: 0.9 mL/min

Injection volume: 10 μL

System suitability

Samples: System suitability solution and Standard solution

[NOTE—See [Table 2](#) for relative retention times.]

Suitability requirements

Resolution: NLT 2.0 between paliperidone related compound B and paliperidone; NLT 2.0 between paliperidone and paliperidone hydroxybenzoyl analog, *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 paliperidone ($C_{23}H_{27}FN_4O_3$) in the portion of Paliperidone 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 Paliperidone RS](#) in the *Standard solution* (mg/mL)

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

Acceptance criteria: 98.0%–102.0%

IMPURITIES

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

• **ORGANIC IMPURITIES**

Protect the System suitability solution, Standard solution, and Sample solution from light.

Diluent, Mobile phase, System suitability solution, Standard solution, Sample solution, and Chromatographic system: Proceed as directed in the Assay.

Sensitivity solution: 0.5 μg/mL of [USP Paliperidone RS](#) in *Diluent* from *Standard solution*

System suitability

Samples: System suitability solution and Sensitivity solution

[NOTE—See [Table 2](#) for relative retention times.]

Suitability requirements

Resolution: NLT 2.0 between paliperidone related compound B and paliperidone; NLT 2.0 between paliperidone and paliperidone hydroxybenzoyl analog, *System suitability solution*

Signal-to-noise ratio: NLT 10, *Sensitivity solution*

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

Analysis:

Samples: Standard solution and Sample solution

Calculate the percentage of any individual impurity in the portion of Paliperidone taken:

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

r_u = peak response for each impurity from the *Sample solution* r_s = peak response of paliperidone from the *Standard solution* C_s = concentration of [USP Paliperidone RS](#) in the *Standard solution* (mg/mL) C_u = concentration of Paliperidone in the *Sample solution* (mg/mL) F = relative response factor for the corresponding impurity (see [Table 2](#))**Acceptance criteria:** See [Table 2](#). Disregard any peak with area less than 0.05%.**Table 2**

Name	Relative Retention Time	Relative Response Factor	Acceptance Criteria, NMT (%)
Paliperidone related compound C ^a	0.57	1.0	0.10
Paliperidone related compound B ^b	0.83	1.0	0.10
Paliperidone	1.00	—	—
Paliperidone hydroxybenzoyl analog ^c	1.1	1.0	0.10
Paliperidone ketone ^d	1.27	0.58	0.50
Any other individual unspecified impurity	—	1.0	0.10
Total impurities	—	—	0.70

^a 3-(2-Chloroethyl)-9-hydroxy-2-methyl-6,7,8,9-tetrahydro-4*H*-pyrido[1,2-*a*]pyrimidin-4-one.^b 6-Fluoro-3-(piperidin-4-yl)benzoisoxazole.^c 3-{2-[4-(4-Fluoro-2-hydroxybenzoyl)piperidin-1-yl]ethyl}-9-hydroxy-2-methyl-6,7,8,9-tetrahydro-4*H*-pyrido[1,2-*a*]pyrimidin-4-one.^d 3-{2-[4-(6-Fluorobenzo[*d*]isoxazol-3-yl)piperidin-1-yl]ethyl}-2-methyl-7,8-dihydro-4*H*-pyrido[1,2-*a*]pyrimidine-4,9(6*H*)-dione.**SPECIFIC TESTS**

- [WATER DETERMINATION, Method 1a \(921\)](#): NMT 0.5%

ADDITIONAL REQUIREMENTS

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

- [USP REFERENCE STANDARDS \(11\)](#).

[USP Paliperidone RS](#)[USP Paliperidone Resolution Mixture RS](#)

This contains paliperidone as the major component; it also contains paliperidone related compound B, paliperidone related compound C, paliperidone hydroxybenzoyl analog, and paliperidone ketone as minor components.

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

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

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

Chromatographic Database Information: [Chromatographic Database](#)

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