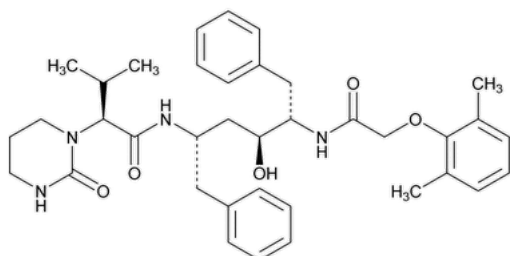


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## Lopinavir



$C_{37}H_{48}N_4O_5$  628.80

[1S-[1*R*\*(*R*\*),3*R*\*,4*R*\*]]-N-[4[[2,6-Dimethylphenoxy] acetyl]amino]-3-hydroxy-5-phenyl-1-(phenylmethyl)pentyl]-tetrahydro- $\alpha$ -(1-methylethyl)-2-oxo-1(2*H*)-pyrimidineacetamide;

( $\alpha$ S)-Tetrahydro-N-[( $\alpha$ S)- $\alpha$ -[(2*S*,3*S*)-2-hydroxy-4-phenyl-3-[2-(2,6-xylyloxy)acetamido]butyl]phenethyl]- $\alpha$ -isopropyl-2-oxo-1(2*H*)-pyrimidineacetamide CAS RN<sup>®</sup>: 192725-17-0; UNII: 2494G1JF75.

### DEFINITION

Lopinavir contains NLT 98.0% and NMT 102.0% of lopinavir ( $C_{37}H_{48}N_4O_5$ ), calculated on the anhydrous basis.

### IDENTIFICATION

**Change to read:**

- **A.** ▲ [SPECTROSCOPIC IDENTIFICATION TESTS \(197\)](#), [Infrared Spectroscopy: 197A](#) ▲ (CN 1-MAY-2020)
- **B.** The retention time of the lopinavir peak of the *Sample solution* corresponds to that of the *Standard solution*, as obtained in the Assay.

### ASSAY

#### PROCEDURE

**Buffer:** 2.7 g/L of monobasic potassium phosphate and 0.9 g/L of dibasic potassium phosphate in water. Adjust with phosphoric acid to a pH of 6.0. Pass the solution through a suitable filter of 0.45- $\mu$ m pore size.

**Diluent:** Acetonitrile and water (1:1)

**Solution A:** Acetonitrile and *Buffer* (9:11)

**Mobile phase:** *Solution A*

**Standard solution:** 0.025 mg/mL of [USP Lopinavir RS](#) in *Diluent*

**Sample solution:** 0.025 mg/mL of Lopinavir in *Diluent*

#### Chromatographic system

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

**Mode:** LC

**Detector:** UV 215 nm

**Column:** 4.6-mm  $\times$  25-cm; 4- $\mu$ m packing L1

**Column temperature:** 50°

**Flow rate:** 1 mL/min

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

**Run time:** 60 min

#### System suitability

**Sample:** *Standard solution*

#### Suitability requirements

**Column efficiency:** NLT 8000 theoretical plates

**Capacity factor:** NLT 15

**Tailing factor:** 0.8–1.5

**Relative standard deviation:** NMT 2.0%

#### Analysis

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

Calculate the percentage of lopinavir ( $C_{37}H_{48}N_4O_5$ ) in the portion of Lopinavir taken:

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 Lopinavir RS](#) in the *Standard solution* (mg/mL)

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

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

**IMPURITIES**

- [RESIDUE ON IGNITION \(281\)](#): NMT 0.2%
- **ORGANIC IMPURITIES: PROCEDURE 1**

[NOTE—For early-eluting impurities.]

**Buffer, Diluent, and Solution A:** Prepare as directed in the Assay.

**Solution B:** Acetonitrile and *Buffer* (3:1)

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

**Table 1**

Time (min)	Solution A (%)	Solution B (%)
0	100	0
60	100	0
61	0	100
81	0	100
82	100	0
100	100	0

**System suitability solution:** 0.5 mg/mL of [USP Lopinavir System Suitability Mixture RS](#) in *Diluent*

**Standard solution:** 0.005 mg/mL of [USP Lopinavir RS](#) in *Diluent*

**Sample solution:** 0.5 mg/mL of Lopinavir in *Diluent*

**Chromatographic system**

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

**Mode:** LC

**Detector:** UV 215 nm

**Column:** 4.6-mm × 25-cm; 4-μm packing L1

**Column temperature:** 50°

**Flow rate:** 1 mL/min

**Injection volume:** 20 μL

**Run time:** 100 min

[NOTE—Data collection is only for the first 60 min. The remaining gradient steps wash out the late-eluting impurities and re-equilibrate the column.]

**System suitability**

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

[NOTE—The relative retention times are listed in [Table 2](#).]

**Suitability requirements**

**Resolution:** NLT 1.2 between lopinavir *N*-formylphenoxyacetamide and lopinavir *N*-acetylphenoxyacetamide, *System suitability solution*

**Capacity factor:** NLT 15, *Standard solution*

**Column efficiency:** NLT 8000, *Standard solution*

**Tailing factor:** 0.8–1.5, *Standard solution*

**Relative standard deviation:** NMT 3.0%, *Standard solution*

**Analysis**

**Samples:** *Diluent*, *System suitability solution*, *Standard solution*, and *Sample solution*

Calculate the percentage of each lopinavir related impurity and unidentified impurity in the portion of Lopinavir taken:

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

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

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

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

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

$F$  = relative response factor (see [Table 2](#))

**Table 2**

Name	Relative Retention	Relative Response Factor	Acceptance Criteria, NMT (%)
Lopinavir free amine <sup>a</sup>	0.03	0.61	0.1
Lopinavir <i>N</i> -formylaminoalcohol <sup>b</sup>	0.07	0.80	0.2
Lopinavir divalinate <sup>c</sup>	0.10	0.65	0.1
Sulfolopinavir <sup>d</sup>	0.13	0.76	0.1
Lopinavir phenoxyacetamide <sup>e</sup>	0.25	0.96	0.1
Lopinavir <i>N</i> -formylphenoxyacetamide <sup>f</sup>	0.59	1.3	0.1
Lopinavir <i>N</i> -acetylphenoxyacetamide <sup>g</sup>	0.62	1.2	0.1
Lopinavir oxazine <sup>h</sup>	0.90	1.1	0.1
Lopinavir	1.00	—	—
Isolopinavir <sup>i</sup>	1.10	0.99	0.2
Lopinavir 2,4-phenoxy isomer <sup>j</sup>	1.13	0.97	0.1
Lopinavir <i>D</i> -valine diastereomer <sup>k</sup>	1.25	1.1	0.1
<i>Z</i> -Diacyclethenediamine <sup>l</sup>	1.28	1.4	0.1
Lopinavir (2 <i>R</i> ,4 <i>R</i> ) diastereomer <sup>m</sup>	1.32	1.0	0.1
Lopinavir (4 <i>R</i> ) epimer <sup>n</sup>	1.38	0.97	0.1
Any other individual impurity	—	1.0	0.1

<sup>a</sup> (S)-*N*-[(2*S*,4*S*,5*S*)-5-Amino-4-hydroxy-1,6-diphenylhexan-2-yl]-3-methyl-2-[2-oxotetrahydropyrimidin-1(2*H*)-yl]butanamide.

<sup>b</sup> (S)-*N*-[(2*S*,4*S*,5*S*)-5-Formamido-4-hydroxy-1,6-diphenylhexan-2-yl]-3-methyl-2-[2-oxotetrahydropyrimidin-1(2*H*)-yl]butanamide.

<sup>c</sup> (2*S*,2'*S*)-*N,N'*-[(2*S*,3*S*,5*S*)-3-Hydroxy-1,6-diphenylhexane-2,5-diyl]bis(3-methyl-2-[2-oxotetrahydropyrimidin-1(2*H*)-yl]butanamide).

<sup>d</sup> (2*S*,3*S*,5*S*)-2-[2-(2,6-Dimethylphenoxy)acetamido]-5-[(S)-3-methyl-2-[2-oxotetrahydropyrimidin-1(2*H*)-yl]butanamido]-1,6-diphenylhexan-3-yl hydrogen sulfate.

<sup>e</sup> *N*-[(2*S*,3*S*,5*S*)-5-Amino-3-hydroxy-1,6-diphenylhexan-2-yl]-2-(2,6-dimethylphenoxy)acetamide.

<sup>f</sup> 2-(2,6-Dimethylphenoxy)-*N*-[(2*S*,3*S*,5*S*)-5-formamido-3-hydroxy-1,6-diphenylhexan-2-yl]acetamide.

<sup>g</sup> *N*-[(2*S*,3*S*,5*S*)-5-Acetamido-3-hydroxy-1,6-diphenylhexan-2-yl]-2-(2,6-dimethylphenoxy)acetamide.

<sup>h</sup> *N*-[(S)-1-[(4*S*,6*S*)-4-Benzyl-2-oxo-1,3-oxazinan-6-yl]-2-phenylethyl]-2-(2,6-dimethylphenoxy)acetamide.

<sup>i</sup> (S)-*N*-[(2*S*,3*S*,5*S*)-5-[2-(2,6-Dimethylphenoxy)acetamido]-3-hydroxy-1,6-diphenylhexan-2-yl]-3-methyl-2-[2-oxotetrahydropyrimidin-1(2*H*)-yl]butanamide.

- j (S)-N-((2S,4S,5S)-5-[2-(2,4-Dimethylphenoxy)acetamido]-4-hydroxy-1,6-diphenylhexan-2-yl)-3-methyl-2-[2-oxotetrahydropyrimidin-1(2H)-yl]butanamide.
- k (R)-N-((2S,4S,5S)-5-[2-(2,6-Dimethylphenoxy)acetamido]-4-hydroxy-1,6-diphenylhexan-2-yl)-3-methyl-2-[2-oxotetrahydropyrimidin-1(2H)-yl]butanamide.
- l (Z)-N,N'-(Ethene-1,2-diyl)bis[2-(2,6-dimethylphenoxy)acetamide].
- m (S)-N-((2R,4R,5S)-5-[2-(2,6-Dimethylphenoxy)acetamido]-4-hydroxy-1,6-diphenylhexan-2-yl)-3-methyl-2-[2-oxotetrahydropyrimidin-1(2H)-yl]butanamide.
- n (S)-N-((2S,4R,5S)-5-[2-(2,6-Dimethylphenoxy)acetamido]-4-hydroxy-1,6-diphenylhexan-2-yl)-3-methyl-2-[2-oxotetrahydropyrimidin-1(2H)-yl]butanamide.

• ORGANIC IMPURITIES: PROCEDURE 2

[NOTE—For late-eluting impurities.]

**Buffer, Diluent, and Solution A:** Prepare as directed in the Assay.

**Solution B:** Acetonitrile and *Buffer* (3:1)

**Mobile phase:** *Solution A* and *Solution B* (3:7)

**System suitability solution:** 0.5 mg/mL of [USP Lopinavir System Suitability Mixture RS](#) in *Diluent*

**Standard solution:** 0.005 mg/mL of [USP Lopinavir RS](#) in *Diluent*

**Sample solution:** 0.5 mg/mL in *Diluent*

**Chromatographic system**

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

**Mode:** LC

**Detector:** UV 215 nm

**Column:** 4.6-mm × 25-cm; 4-μm packing L1

**Column temperature:** 50°

**Flow rate:** 1 mL/min

**Injection volume:** 20 μL

**Run time:** 50 min

**System suitability**

**Sample:** *Standard solution*

[NOTE—The relative retention times are listed in [Table 3](#).]

**Suitability requirements**

**Capacity factor:** NLT 1.5

**Column efficiency:** NLT 3000

**Tailing factor:** 0.8–1.5

**Relative standard deviation:** NMT 3.0%

**Analysis**

**Samples:** *Diluent*, *System suitability solution*, *Standard solution*, and *Sample solution*

Calculate the percentage of each lopinavir related impurity and unidentified impurity in the portion of Lopinavir taken:

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

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

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

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

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

$F$  = relative response factor (see [Table 3](#))

**Table 3**

Name	Relative Retention	Relative Response Factor	Acceptance Criteria, NMT (%)
Lopinavir	1.00	—	—
Lopinavir O-acetyl <sup>a</sup>	1.49	0.77	0.1
Lopinavir (2R) epimer <sup>b</sup>	1.91	1.1	0.1

Name	Relative Retention	Relative Response Factor	Acceptance Criteria, NMT (%)
Lopinavir diamide <sup>c</sup>	4.39	1.4	0.1
Lopinavir <i>N</i> -acyl <sup>d</sup>	6.01	1.3	0.1
Lopinavir <i>O</i> -phenoxyactyl <sup>e</sup>	7.14	1.1	0.1
Lopinavir amino alcohol urea <sup>f</sup>	8.46	1.3	0.1
Any other individual impurity	—	1.0	0.1
Total impurities from Procedure 1 and Procedure 2	—	1.0	0.7 <sup>g</sup>

- <sup>a</sup> (S)-{(2S,3S,5S)-2-[2-(2,6-Dimethylphenoxy)acetamido]-5-[(S)-3-methyl-2-(2-oxotetrahydropyrimidin-1(2*H*)-yl)butanamido]-1,6-diphenylhexan-3-yl} 3-methyl-2-[2-oxotetrahydropyrimidin-1(2*H*)-yl]butanoate.
- <sup>b</sup> (S)-*N*-{(2*R*,4*S*,5*S*)-5-[2-(2,6-Dimethylphenoxy)acetamido]-4-hydroxy-1,6-diphenylhexan-2-yl}-3-methyl-2-[2-oxotetrahydropyrimidin-1(2*H*)-yl]butanamide.
- <sup>c</sup> *N,N'*-[(2*S*,3*S*,5*S*)-3-Hydroxy-1,6-diphenylhexane-2,5-diyl]bis[2-(2,6-dimethylphenoxy)acetamide].
- <sup>d</sup> (S)-*N*-{(2*S*,4*S*,5*S*)-5-[2-(2,6-Dimethylphenoxy)acetamido]-4-hydroxy-1,6-diphenylhexan-2-yl}-2-{3-[2-(2,6-dimethylphenoxy)acetyl]-2-oxotetrahydropyrimidin-1(2*H*)-yl}-3-methylbutanamide.
- <sup>e</sup> (2*S*,3*S*,5*S*)-2-[2-(2,6-Dimethylphenoxy)acetamido]-5-[(S)-3-methyl-2-[2-oxotetrahydropyrimidin-1(2*H*)-yl]butanamido]-1,6-diphenylhexan-3-yl 2-(2,6-dimethylphenoxy)acetate.
- <sup>f</sup> *N,N'*-(2*S*,2'*S*,3*S*,3'*S*,5*S*,5'*S*)-5,5'-Carbonylbis(azanediyl)bis(3-hydroxy-1,6-diphenylhexane-5,2-diyl)bis[2-(2,6-dimethylphenoxy)acetamide].
- <sup>g</sup> Exclude from *Organic Impurities, Procedure 2*, lopinavir (4*R*) epimer and any other peak eluting prior to this peak because these are already monitored in *Procedure 1*.

Acceptance criteria: See [Table 2](#) and [Table 3](#).

SPECIFIC TESTS

- [WATER DETERMINATION, Method I\(921\)](#): NMT 4.4%

ADDITIONAL REQUIREMENTS

- **PACKAGING AND STORAGE:** Preserve in tight containers. Store at room temperature.

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

[USP Lopinavir RS](#)

[USP Lopinavir System Suitability Mixture RS](#)

Lopinavir System Suitability Mixture contains lopinavir *N*-formylphenoxyacetamide, lopinavir *N*-acetylphenoxyacetamide, and several other minor components.

Lopinavir *N*-formylphenoxyacetamide is (2-(2,6-dimethylphenoxy)-*N*-[(2*S*,3*S*,5*S*)-5-formamido-3-hydroxy-1,6-diphenylhexan-2-yl]acetamide.



Lopinavir *N*-acetylphenoxyacetamide is (*N*-[(2*S*,3*S*,5*S*)-5-acetamido-3-hydroxy-1,6-diphenylhexan-2-yl]-2-(2,6-

dimethylphenoxy)acetamide.  $C_{30}H_{36}N_2O_4$  488.62

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

Topic/Question	Contact	Expert Committee
LOPINAVIR	<a href="#">Documentary Standards Support</a>	SM12020 Small Molecules 1

Chromatographic Database Information: [Chromatographic Database](#)

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