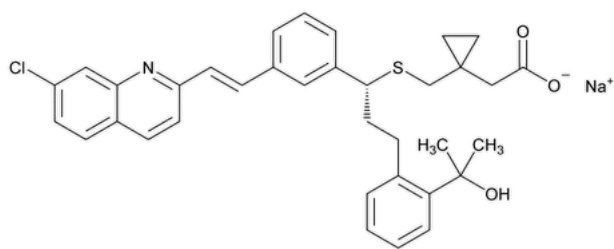


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Montelukast Sodium



$C_{35}H_{35}ClNaO_3S$ 608.17
Cyclopropaneacetic acid, 1-[[[1-[3-[2-(7-chloro-2-quinolinyl)ethenyl]phenyl]-3-[2-(1-hydroxy-1-methylethyl)phenyl]propyl]thio]methyl]-, sodium salt, [R, (E)];
Sodium 1-[[[(R) -m-[(E)-2-(7-chloro-2-quinolyl)vinyl]-α-[o-(1-hydroxy-1-methylethyl)phenethyl]benzyl]thio]-methyl]cyclopropaneacetate CAS RN®: 151767-02-1; UNII: U103J18SFL.
 $C_{35}H_{36}ClNO_3S$ 586.18
Montelukast CAS RN®: 158966-92-8; UNII: MHM278SD3E.

DEFINITION
Montelukast Sodium contains NLT 98.0% and NMT 102.0% of $C_{35}H_{35}ClNaO_3S$, calculated on the anhydrous and solvent-free basis.

IDENTIFICATION

Change to read:

- **A.**▲ [SPECTROSCOPIC IDENTIFICATION TESTS \(197\)](#), *Infrared Spectroscopy*: 197A, 197K, or 197M▲ (CN 1-May-2020)
- **B.** [IDENTIFICATION TESTS—GENERAL, Sodium \(191\)](#).
Sample: 100 mg
Analysis: Ignite the *Sample* in a crucible until an almost white residue is obtained. Take up the residue in 2 mL of water, and filter.
Acceptance criteria: The filtrate meets the requirements of test A.
- **C.** Meets the requirements of the test for *Enantiomeric Purity*.

ASSAY

[NOTE—Avoid exposure of the samples to light. Use low-actinic glassware.]

• **PROCEDURE**

- Solution A:** Add 1.5 mL of trifluoroacetic acid to 1 L of water.
Solution B: Add 1.5 mL of trifluoroacetic acid to 1 L of acetonitrile.
Mobile phase: See [Table 1](#). Return to original conditions and re-equilibrate the column.

Table 1

Time (min)	Solution A (%)	Solution B (%)
0	60	40
3.0	60	40
16.0	49	51

Diluent: Methanol and water (9:1)
Standard solution: 0.13 mg/mL of [USP Montelukast Dicyclohexylamine RS](#) in *Diluent*
Sample solution: 0.1 mg/mL of Montelukast Sodium in *Diluent*
Chromatographic system
(See [Chromatography \(621\), System Suitability.](#))
Mode: LC

Detector: UV 238 nm**Column:** 4.6-mm × 5-cm; 1.8-μm packing L11**Column temperature:** 30°**Flow rate:** 1.2 mL/min**Injection size:** 10 μL**System suitability****Sample:** *Standard solution***Suitability requirements****Relative standard deviation:** NMT 0.73%**Analysis****Samples:** *Standard solution* and *Sample solution*Calculate the percentage of montelukast sodium (C₃₅H₃₅ClNNaO₃S) in the portion of Montelukast Sodium taken:

$$\text{Result} = (r_U/r_S) \times (C_S/C_U) \times (M_{r1}/M_{r2}) \times 100$$

 r_U = peak area from the *Sample solution* r_S = peak area from the *Standard solution* C_S = concentration of the *Standard solution* (mg/mL) C_U = concentration of the *Sample solution* (mg/mL) M_{r1} = molecular weight of montelukast sodium, 608.17 M_{r2} = molecular weight of montelukast dicyclohexylamine, 767.50**Acceptance criteria:** 98.0%–102.0% on the anhydrous and solvent-free basis**IMPURITIES****• ORGANIC IMPURITIES**

[NOTE—Avoid exposure of the samples to light. Use low-actinic glassware.]

Solution A, Solution B, Mobile phase, Diluent, and Chromatographic system: Proceed as directed in the Assay.**Impurity solution:** 1 mg/mL of [USP Montelukast for Peak Identification RS](#) in *Diluent***System suitability solution:** Transfer 1 mL of the *Impurity solution* to a colorless glass vial, and expose to ambient light for approximately 20 min to generate the *cis*-isomer of montelukast.**Sample solution:** 1 mg/mL of Montelukast Sodium in *Diluent***Sensitivity solution:** 0.5 μg/mL of Montelukast Sodium in *Diluent* from the *Sample solution***System suitability****Samples:** *System suitability solution* and *Sensitivity solution***Suitability requirements****Resolution:** NLT 2.5 between the *cis*-isomer and montelukast; NLT 1.5 between montelukast and the methylketone impurity, *System suitability solution***Signal-to-noise ratio:** NLT 10, *Sensitivity solution***Analysis****Sample:** *Sample solution*

Calculate the percentage of each impurity in the portion of Montelukast Sodium taken:

$$\text{Result} = (r_U/r_T) \times 100$$

 r_U = peak response of each impurity from the *Sample solution* r_T = sum of all the peak responses from the *Sample solution***Acceptance criteria:** See [Table 2](#).**Reporting level for impurities:** 0.05%**Table 2**

Name	Relative Retention Time	Acceptance Criteria, NMT (%)
Sulfoxide impurity ^a	0.4	0.2

Name	Relative Retention Time	Acceptance Criteria, NMT (%)
Cis-isomer ^b	0.8	0.15
Michael Adducts 1 ^c and 2 ^d	0.9	0.15 [*]
Montelukast	1.0	—
Methylketone impurity ^e	1.2	0.15
Methylstyrene impurity ^f	1.9	0.3
Any other individual impurity	—	0.10
Total impurities	—	0.6

* These two impurities are not resolved by the method and need to be integrated together to determine conformance.

^a [1-[[[1-[3-[(E)-2-(7-Chloroquinolin-2-yl)ethenyl]phenyl]-3-[2-(1-hydroxy-1-methylethyl)phenyl]propyl]sulfinyl]methyl]cyclopropyl]acetic acid.

^b [1-[[[1(R)-1-[3-[(Z)-2-(7-Chloroquinolin-2-yl)ethenyl]phenyl]-3-[2-(1-hydroxy-1-methylethyl)phenyl]propyl]sulfonyl]methyl]cyclopropyl]acetic acid.

^c 1-[[[1(R)-1-[3-[(1R)-1-[[[1-(Carboxymethyl)cyclopropyl]methyl]sulfonyl]-2-(7-chloroquinolin-2-yl)ethyl]phenyl]-3-[2-(1-hydroxy-1-methylethyl)phenyl]propyl]sulfonyl]methyl]cyclopropyl]acetic acid.

^d 1-[[[1(R)-1-[3-[(1S)-1-[[[1-(Carboxymethyl)cyclopropyl]methyl]sulfonyl]-2-(7-chloroquinolin-2-yl)ethyl]phenyl]-3-[2-(1-hydroxy-1-methylethyl)phenyl]propyl]sulfonyl]methyl]cyclopropyl]acetic acid.

^e [1-[[[1(R)-3-(2-Acetylphenyl)-1-[3-[(E)-2-(7-chloroquinolin-2-yl)ethenyl]phenyl]propyl]sulfonyl]methyl]cyclopropyl]acetic acid.

^f [1-[[[1(R)-1-[3-[(E)-2-(7-Chloroquinolin-2-yl)ethenyl]phenyl]-3-[2-(1-methylethenyl)phenyl]propyl]sulfonyl]methyl]cyclopropyl]acetic acid.

• ENANTIOMERIC PURITY

[NOTE—Avoid exposure of the samples to light. Use low-actinic glassware.]

Solution A: 2.3 g/L of ammonium acetate in water. Adjust with glacial acetic acid to a pH of 5.7.

Solution B: Methanol and acetonitrile (60:40)

Mobile phase: See [Table 3](#).

Table 3

Time (min)	Solution A (%)	Solution B (%)
0	70	30
30	60	40
35	60	40

Diluent: Acetonitrile and water (1:1)

System suitability solution: 0.1 mg/mL of [USP Montelukast Racemate RS](#) in Diluent

Sample solution: 1 mg/mL of Montelukast Sodium in Diluent

Sensitivity solution: 1 µg/mL of Montelukast Sodium in Diluent from the Sample solution

Chromatographic system

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

Mode: LC

Detector: UV 280 nm

Column: 4.0-mm × 15-cm; 5-µm packing L41

Column temperature: 30°

Flow rate: 0.9 mL/min

Injection size: 10 µL

System suitability

Samples: System suitability solution and Sensitivity solution

[NOTE—The relative retention times are 1.0 for montelukast, which is the R-enantiomer, and 0.7 for the S-enantiomer.]

Suitability requirements

Resolution: NLT 2.9 between the S-enantiomer and montelukast, *System suitability solution*

Signal-to-noise ratio: NLT 10 for the montelukast peak, *Sensitivity solution*

Analysis

Sample: *Sample solution*

Calculate the percentage of S-enantiomer in the portion of Montelukast Sodium taken:

$$\text{Result} = (r_U/r_T) \times 100$$

r_U = peak response of the S-enantiomer from the *Sample solution*

r_T = sum of the peak responses of the S-enantiomer and montelukast from the *Sample solution*

Acceptance criteria: NMT 0.2% of the S-enantiomer

SPECIFIC TESTS

- [WATER DETERMINATION, Method Ia\(921\)](#): NMT 4.0%

ADDITIONAL REQUIREMENTS

- **PACKAGING AND STORAGE:** Preserve in tight containers, protected from light. Store at room temperature.

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

[USP Montelukast Sodium RS](#)

[USP Montelukast Dicyclohexylamine RS](#)

$\text{C}_{35}\text{H}_{36}\text{ClNO}_3\text{S} \cdot \text{C}_{12}\text{H}_{23}\text{N}$

767.50

[USP Montelukast Racemate RS](#)

[USP Montelukast for Peak Identification RS](#)

(montelukast containing sulfoxide impurity, michael adducts 1 and 2, methylketone impurity, and methylstyrene impurity)

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

Topic/Question	Contact	Expert Committee
MONTELUKAST SODIUM	Documentary Standards Support	SM52020 Small Molecules 5
REFERENCE STANDARD SUPPORT	RS Technical Services RSTECH@usp.org	SM52020 Small Molecules 5

Chromatographic Database Information: [Chromatographic Database](#)

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