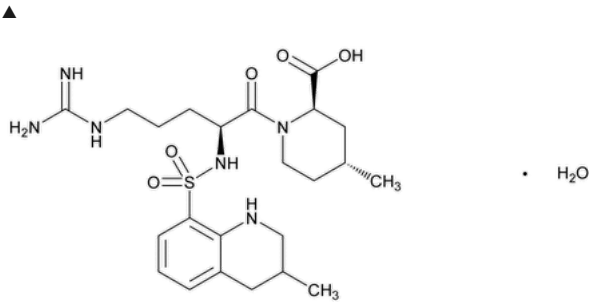


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Change to read:

Argatroban



▲ (ERR 1-Mar-2019)
 $C_{23}H_{36}N_6O_5S \cdot H_2O$ 526.65
2-Piperidinecarboxylic acid, 1-[(S)-5-[(aminoiminomethyl)amino]-1-oxo-2-[(1,2,3,4-tetrahydro-3-methyl-8-quinolyl)sulfonyl]amino]pentyl]-4-methyl-, (2R,4R)-monohydrate;
(2R,4R)-4-Methyl-1-{N²-[(1,2,3,4-tetrahydro-3-methyl-8-quinolyl)sulfonyl]-L-arginyl}pipecolic acid, monohydrate CAS RN®: 141396-28-3; UNII: IY90U61Z3S.

DEFINITION
Argatroban contains NLT 98.0% and NMT 102.0% of argatroban ($C_{23}H_{36}N_6O_5S$), calculated on the anhydrous and solvent-free basis.

IDENTIFICATION

Change to read:

- **A.** ▲ [SPECTROSCOPIC IDENTIFICATION TESTS <197>](#), *Infrared Spectroscopy*: **197K** ▲ (CN 1-MAY-2020)
- **B.** The retention times of the major peaks of the *Sample solution* correspond to those of the *Standard solution*, as obtained in the Assay.

ASSAY

• PROCEDURE

[NOTE—It is recommended to keep all solutions containing argatroban at about 4°.]

Solution A: 10 mM ammonium acetate and 5 mM sodium 1-heptanesulfonate

Solution B: Acetonitrile and methanol (500:300)

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

Table 1

Time (min)	Solution A (%)	Solution B (%)
0	60	40
20	60	40
35	50	50
50	20	80
60	20	80
60.1	60	40
72.1	60	40

Standard solution: 4 mg/mL of [USP Argatroban RS](#) in methanol

Sample solution: 4 mg/mL of Argatroban in methanol

Chromatographic system

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

Mode: LC

Detector: UV 259 nm

Column: 4.6-mm × 25-cm; 3-μm packing L1

Temperatures

Column: 50°

Autosampler: 4°

Flow rate: 0.6 mL/min

Injection volume: 10 μL

System suitability

Sample: *Standard solution*

Suitability requirements

Tailing factor: NMT 1.5 for both peaks

Relative standard deviation: NMT 1.0% for the sum of the peak responses of (R)-argatroban and (S)-argatroban

Analysis

Samples: *Standard solution* and *Sample solution*

Calculate the percentage of argatroban ($C_{23}H_{36}N_6O_5S$) in the portion of Argatroban taken:

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

r_U = sum of the peak responses of (R)-argatroban and (S)-argatroban from the *Sample solution*

r_S = sum of the peak responses of (R)-argatroban and (S)-argatroban from the *Standard solution*

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

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

Acceptance criteria: 98.0%–102.0% on the anhydrous and solvent-free basis

IMPURITIES

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

• **ORGANIC IMPURITIES**

[NOTE—It is recommended to keep all solutions containing argatroban at about 4°.]

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

Sensitivity solution: 2 μg/mL of [USP Argatroban RS](#) in methanol

Standard solution: 4 μg/mL each of [USP Argatroban RS](#), [USP Argatroban Related Compound A RS](#), and [USP Argatroban Related Compound B RS](#) in methanol

System suitability

Samples: *Sensitivity solution* and *Standard solution*

Suitability requirements

Resolution: NLT 1.2 between (R)-argatroban and (S)-argatroban, *Standard solution*

Signal-to-noise ratio: NLT 10 for (R)-argatroban, *Sensitivity solution*

Relative standard deviation: NMT 5% for all peaks. For argatroban, use the sum of the peak responses of (R)-argatroban and (S)-argatroban, *Standard solution*.

Analysis

Samples: *Standard solution* and *Sample solution*

Calculate the percentage of argatroban related compound A and argatroban related compound B in the portion of Argatroban taken:

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

r_U = peak response of argatroban related compound A or argatroban related compound B from the *Sample solution*

r_S = peak response of argatroban related compound A or argatroban related compound B from the *Standard solution*

C_S = concentration of [USP Argatroban Related Compound A RS](#) or [USP Argatroban Related Compound B RS](#) in the *Standard solution* (mg/mL)

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

Calculate the percentage of any unspecified impurity in the portion of Argatroban taken:

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

r_U = peak response of any unspecified impurity from the *Sample solution*

r_S = sum of the peak responses of (R)-argatroban and (S)-argatroban from the *Standard solution*

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

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

Acceptance criteria: See [Table 2](#). Disregard any peak below 0.05%.

Table 2

Name	Relative Retention Time	Acceptance Criteria, NMT (%)
Argatroban related compound A ^a	0.23	0.15
Argatroban related compound B ^b	0.39	0.15
(R)-Argatroban ^c	1.00	—
(S)-Argatroban ^d	1.03	—
Any unspecified impurity	—	0.10
Total impurities ^e	—	0.5

^a (2R,4R)-1-[N⁸-Nitro-N²-(3-methylquinoline-8-sulfonyl)-L-arginyl]-4-methylpiperidine-2-carboxylic acid.

^b Ethyl (4R)-1-[N⁸-nitro-L-arginyl]-4-methylpiperidine-2-carboxylate.

^c (2R,4R)-4-Methyl-1-{N²-[[(R)-1,2,3,4-tetrahydro-3-methyl-8-quinolyl)sulfonyl]-L-arginyl}pipecolic acid.

^d (2R,4R)-4-Methyl-1-{N²-[[(S)-1,2,3,4-tetrahydro-3-methyl-8-quinolyl)sulfonyl]-L-arginyl}pipecolic acid.

^e Total impurities include specified and unspecified impurities and argatroban related compound C from the test for *Content of Argatroban Related Compound C*.

• CONTENT OF ARGATROBAN RELATED COMPOUND C

[NOTE—It is recommended to keep all solutions containing argatroban at about 4°.]

Buffer: 10 mM ammonium acetate and 5 mM sodium 1-heptanesulfonate

Solution A: Acetonitrile, dehydrated alcohol, and *Buffer* (80:240:680)

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

Table 3

Time (min)	Solution A (%)	Acetonitrile (%)
0	100	0
70	100	0
71	30	70
91	30	70
92	100	0
102	100	0

Sensitivity solution: 4 µg/mL of [USP Argatroban Related Compound C RS](#) in methanol

System suitability solution: 10 mg/mL of [USP Argatroban RS](#) and 0.1 mg/mL of [USP Argatroban Related Compound C RS](#) in methanol

Sample solution: 10 mg/mL of Argatroban in methanol

Chromatographic system: Proceed as directed in the Assay.

System suitability

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

Suitability requirements

Resolution: NLT 1.4 between argatroban related compound C and (R)-argatroban, *System suitability solution*

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

Relative standard deviation: NMT 2.5% for argatroban related compound C, *System suitability solution*

Analysis

Sample: *Sample solution*

Calculate the percentage of argatroban related compound C in the portion of Argatroban taken:

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

r_U = peak response of argatroban related compound C from the *Sample solution*

r_T = total of all peak responses from the *Sample solution*

Acceptance criteria: See [Table 4](#).

Table 4

Name	Relative Retention Time	Acceptance Criteria, NMT (%)
Argatroban related compound C ^a	0.94	0.15
(R)-Argatroban	1.00	—
(S)-Argatroban	1.07	—

^a (2R,4R)-1-[N⁸-Amino-N²-(3-methyl-1,2,3,4-tetrahydroquinoline-8-sulfonyl)-L-arginyl]-4-methylpiperidine-2-carboxylic acid.

• **CONTENT OF STEREOISOMERS**

[NOTE—It is recommended to keep all solutions containing argatroban at about 4°.]

Mobile phase: Methanol and water (520:480)

Standard solution: 0.16 mg/mL of [USP Argatroban RS](#) in methanol

Sample solution: 0.16 mg/mL of Argatroban in methanol

Chromatographic system

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

Mode: LC

Detector: UV 259 nm

Column: 4.6-mm × 25-cm; 3-μm packing L1

Column temperature: 50°

Flow rate: 0.6 mL/min

Injection volume: 10 μL

Run time: NLT 1.4 times the retention time of (R)-argatroban

System suitability

Sample: *Standard solution*

Suitability requirements

Resolution: NLT 1.2 between (R)-argatroban and (S)-argatroban

Relative standard deviation: NMT 2.0% for (R)-argatroban and (S)-argatroban

Analysis

Sample: *Sample solution*

Calculate the percentage of (R)-argatroban and (S)-argatroban in the portion of Argatroban taken:

$$\text{Result} = [(r_U \text{ or } r_S)/(r_U + r_S)] \times 100$$

r_U = peak response of (R)-argatroban from the *Sample solution*

r_S = peak response of (S)-argatroban from the *Sample solution*

Acceptance criteria: See [Table 5](#).

Table 5

Name	Relative Retention Time	Acceptance Criteria, (%)
(R)-Argatroban ^a	1.00	63–67
(S)-Argatroban ^b	1.06	33–37

^a (2*R*,4*R*)-4-Methyl-1- $\{N^2-[(R)-1,2,3,4\text{-tetrahydro-3-methyl-8-quinolyl)sulfonyl}]\text{-L-arginyl}\}$ pipecolic acid.

^b (2*R*,4*R*)-4-Methyl-1- $\{N^2-[(S)-1,2,3,4\text{-tetrahydro-3-methyl-8-quinolyl)sulfonyl}]\text{-L-arginyl}\}$ pipecolic acid.

SPECIFIC TESTS

- **WATER DETERMINATION** (921), *Method Ia*: 3.0%–6.0%
- **BACTERIAL ENDOTOXINS TEST** (85): NMT 2.0 USP Endotoxin Units/mg of argatroban
- **MICROBIAL ENUMERATION TESTS** (61) and **TESTS FOR SPECIFIED MICROORGANISMS** (62): The total aerobic microbial count is NMT 10^2 cfu/g, and the total combined molds and yeasts count is NMT 10^2 cfu/g.

ADDITIONAL REQUIREMENTS

- **PACKAGING AND STORAGE:** Preserve in tight, light-resistant containers, and store at controlled room temperature.
- **USP REFERENCE STANDARDS** (11).

USP Argatroban RS

USP Argatroban Related Compound A RS

(2*R*,4*R*)-1- $[N^8\text{-Nitro-}N^2\text{-(3-methylquinoline-8-sulfonyl)-L-arginyl}]\text{-4-methylpiperidine-2-carboxylic acid}$.

$C_{23}H_{31}N_7O_7S$ 549.60

USP Argatroban Related Compound B RS

Ethyl (4*R*)-1- $[N^8\text{-nitro-L-arginyl}]\text{-4-methylpiperidine-2-carboxylate dihydrochloride}$.

$C_{15}H_{28}N_6O_5 \cdot 2HCl$ 445.34

USP Argatroban Related Compound C RS

(2*R*,4*R*)-1- $[N^8\text{-Amino-}N^2\text{-(3-methyl-1,2,3,4-tetrahydroquinoline-8-sulfonyl)-L-arginyl}]\text{-4-methylpiperidine-2-carboxylic acid}$.

$C_{23}H_{37}N_7O_5S$ 523.65

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

Topic/Question	Contact	Expert Committee
ARGATROBAN	Documentary Standards Support	SM22020 Small Molecules 2
REFERENCE STANDARD SUPPORT	RS Technical Services RSTECH@usp.org	SM22020 Small Molecules 2

Chromatographic Database Information: [Chromatographic Database](#)

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