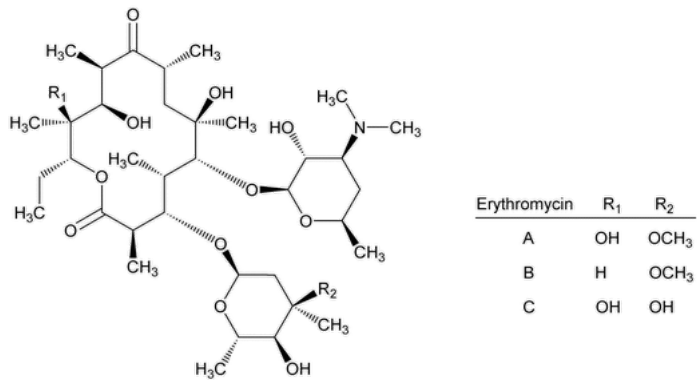


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# Erythromycin

Change to read:



C<sub>37</sub>H<sub>67</sub>NO<sub>13</sub> 733.94

▲Erythromycin A

(3*R*,4*S*,5*S*,6*R*,7*R*,9*R*,11*R*,12*R*,13*S*,14*R*)-4-[(2,6-Dideoxy-3-C-methyl-3-O-methyl-α-*L*-ribo-hexopyranosyl)oxy]-14-ethyl-7,12,13-trihydroxy-3,5,7,9,11,13-hexamethyl-6-[(3,4,6-trideoxy-3-dimethylamino-β-*D*-xylo-hexopyranosyl)oxy]oxacyclotetradecane-2,10-dione;

C<sub>37</sub>H<sub>67</sub>NO<sub>12</sub> 717.94

Erythromycin B

(3*R*,4*S*,5*S*,6*R*,7*R*,9*R*,11*R*,12*R*,13*S*,14*R*)-4-[(2,6-Dideoxy-3-C-methyl-3-O-methyl-α-*L*-ribo-hexopyranosyl)oxy]-14-ethyl-7,12-dihydroxy-3,5,7,9,11,13-hexamethyl-6-[(3,4,6-trideoxy-3-dimethylamino-β-*D*-xylo-hexopyranosyl)oxy]oxacyclotetradecane-2,10-dione;

12-Deoxyerythromycin CAS RN®: 527-75-3.

C<sub>36</sub>H<sub>65</sub>NO<sub>13</sub> 719.91

Erythromycin C

(3*R*,4*S*,5*S*,6*R*,7*R*,9*R*,11*R*,12*R*,13*S*,14*R*)-4-[(2,6-Dideoxy-3-C-methyl-α-*L*-ribo-hexopyranosyl)oxy]-14-ethyl-7,12,13-trihydroxy-3,5,7,9,11,13-hexamethyl-6-[(3,4,6-trideoxy-3-dimethylamino-β-*D*-xylo-hexopyranosyl)oxy]oxacyclotetradecane-2,10-dione;

3'-O-Demethylerythromycin CAS RN®: 1675-02-1.▲ (USP 1-Dec-2024)

Erythromycin

▲▲ (USP 1-Dec-2024) CAS RN®: 114-07-8; UNII: 63937KV33D.

Change to read:

## DEFINITION

Erythromycin consists primarily of erythromycin A (C<sub>37</sub>H<sub>67</sub>NO<sub>13</sub>). The sum of the percentages of erythromycin A, erythromycin B, and erythromycin C is NLT ▲93.0%▲ (USP 1-Dec-2024) and NMT ▲102.0%▲ (USP 1-Dec-2024) calculated on the anhydrous basis.

## IDENTIFICATION

• A. [SPECTROSCOPIC IDENTIFICATION TESTS \(197\)](#), [Infrared Spectroscopy](#): 197S

**Standard solution:** 50 mg/mL of [USP Erythromycin RS](#), previously dried at a pressure not exceeding 5 mm of mercury at 60° for 3 h, in chloroform

**Sample solution:** 50 mg/mL of Erythromycin, previously dried at a pressure not exceeding 5 mm of mercury at 60° for 3 h, in chloroform

**Spectral range:** 4000–2050 cm<sup>-1</sup> and 1980–400 cm<sup>-1</sup>

**Acceptance criteria:** Meets the requirements

• B. The retention times of erythromycin A, erythromycin B, and erythromycin C in the *Sample solution* correspond to those of *Standard solution 1* and *Standard solution 2*, as obtained in the Assay.

## ASSAY

Change to read:

• PROCEDURE

▲Prepare the erythromycin solutions immediately before use.

**Diluted phosphoric acid:** Dilute 7 mL of [phosphoric acid](#) with [water](#) to 100 mL.

**Phosphate buffer solution pH 8.0:** Dissolve 11.5 g of [dibasic potassium phosphate](#) in 900 mL of [water](#). Adjust to a pH of 8.0 with *Diluted phosphoric acid* and dilute with [water](#) to 1000 mL.

**Diluent:** *Phosphate buffer solution pH 8.0* and [methanol](#) (60:40)

**Phosphate buffer solution pH 7.0:** Dissolve 35 g of [dibasic potassium phosphate](#) in 900 mL of [water](#). Adjust to a pH of 7.0 with *Diluted phosphoric acid* and dilute with [water](#) to 1000 mL.

**Solution A:** *Phosphate buffer solution pH 7.0*, [water](#), and [acetonitrile](#) (5:60:35)

**Solution B:** *Phosphate buffer solution pH 7.0*, [water](#), and [acetonitrile](#) (5:45:50)

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

**Table 1**

Time (min)	Solution A (%)	Solution B (%)
0	100	0
$T_R^a$	100	0
$T_R + 2$	0	100
$T_R + 15$	0	100

<sup>a</sup>  $T_R$  = retention time of erythromycin B, determined by injecting 10  $\mu$ L of *Standard solution 2* and eluting with *Solution A*.

**Standard solution 1:** 4 mg/mL of [USP Erythromycin RS](#) in *Diluent*

**Standard solution 2:** 0.2 mg/mL of [USP Erythromycin B RS](#) and [USP Erythromycin C RS](#) in *Diluent*

**Sample solution:** 4 mg/mL of Erythromycin in *Diluent*

**Chromatographic system**

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

**Mode:** LC

**Detector:** UV 210 nm

**Column:** 4.6-mm  $\times$  25-cm; 3.5- $\mu$ m packing [L1](#)

**Temperatures**

**Column:** 65°, preheating the *Mobile phase* may be required, for instance by extending the inlet tubing in the oven to 30 cm

**Sampler:** 4°

**Flow rate:** 1.0 mL/min

**Injection volume:** 100  $\mu$ L

**System suitability**

**Sample:** *Standard solution 1*

**Suitability requirements**

**Tailing factor:** NMT 2.0 for erythromycin A

**Relative standard deviation:** NMT 1.0% for erythromycin A, 6 replicate injections

**Analysis**

**Samples:** *Standard solution 1*, *Standard solution 2*, and *Sample solution*

Calculate the percentage of erythromycin A in the portion of Erythromycin taken:

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

$r_U$  = peak response of erythromycin A from the *Sample solution*

$r_S$  = peak response of erythromycin A from *Standard solution 1*

$C_S$  = concentration of [USP Erythromycin RS](#) in *Standard solution 1* (mg/mL)

$C_U$  = concentration of Erythromycin, calculated on the anhydrous basis, in the *Sample solution* (mg/mL)

$P$  = percentage of erythromycin A in [USP Erythromycin RS](#)

Calculate the percentages of erythromycin B and erythromycin C in the portion of Erythromycin taken:

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

$r_U$  = peak response of the relevant analyte from the *Sample solution*

$r_s$  = peak response of the relevant analyte from *Standard solution 2*

$C_s$  = concentration of the corresponding Reference Standard in *Standard solution 2* (mg/mL)

$C_u$  = concentration of Erythromycin, calculated on the anhydrous basis, in the *Sample solution* (mg/mL)

$P$  = potency of erythromycin B or erythromycin C in the corresponding Reference Standard (mg/mg)

#### Acceptance criteria

**Sum of Erythromycin A, Erythromycin B, and Erythromycin C:** 93.0%–102.0% on the anhydrous basis

**Erythromycin B:** NMT 5.0% on the anhydrous basis

**Erythromycin C:** NMT 5.0% on the anhydrous basis▲ (USP 1-Dec-2024)

#### IMPURITIES

- [RESIDUE ON IGNITION \(281\)](#): NMT 0.2%

#### Change to read:

- **LIMIT OF THIOCYANATE**

Use the *Standard solutions*, *Sample solution*, and *Blank solution* within 30 min.

**Standard stock solutions 1 and 2:** 0.2 mg/mL of potassium thiocyanate prepared in duplicate as follows. Transfer 100 mg of [potassium thiocyanate](#), previously dried at 105° for 1 h and cooled, to a 50-mL volumetric flask. Add about 20 mL of [methanol](#) to each flask, swirl to dissolve, and dilute with [methanol](#) to volume. Transfer 5.0 mL of this solution to a 50-mL volumetric flask and dilute with [methanol](#) to volume.

**Standard solutions 1 and 2:** 0.02 mg/mL of potassium thiocyanate prepared in duplicate as follows. Transfer 5.0 mL of each of the *Standard stock solutions* to separate 50-mL low-actinic volumetric flasks, add 1.0 mL of [ferric chloride TS](#), dilute with [methanol](#) to volume.

**Sample solution:** 2 mg/mL of Erythromycin prepared as follows. Transfer 100 mg of Erythromycin to a 50-mL low-actinic volumetric flask, add 20 mL of [methanol](#), and swirl to dissolve. Add 1.0 mL of [ferric chloride TS](#) and dilute with [methanol](#) to volume.

**Blank solution:** Add 1.0 mL of [ferric chloride TS](#) to a 50-mL low-actinic volumetric flask. Dilute with [methanol](#) to volume.

#### Instrumental conditions

(See [Ultraviolet-Visible Spectroscopy \(857\)](#).)

**Mode:** UV-Vis

**Analytical wavelength:** 492 nm

#### System suitability

**Samples:** *Standard solution 1*, *Standard solution 2*, and *Blank solution*

Use the *Blank solution* to zero the instrument. Measure the absorbance of the two *Standard solutions*.

**Suitability:** 0.985–1.015

Calculate the suitability,  $S$ :

$$\text{Result} = (A_1/W_1) \times (W_2/A_2)$$

$A_1$  = absorbance of *Standard solution 1*

$W_1$  = weight of the potassium thiocyanate taken to prepare *Standard solution 1* (mg)

$W_2$  = weight of the potassium thiocyanate taken to prepare *Standard solution 2* (mg)

$A_2$  = absorbance of *Standard solution 2*

#### Analysis

**Samples:** *Standard solution 1*, *Standard solution 2*, and *Sample solution*

Calculate the percentage of thiocyanate in the portion of Erythromycin taken:

$$\text{▲Result} = (M_{r1}/M_{r2}) \times (A_u/W_u) \times 0.5 \times [(W_1/A_1) + (W_2/A_2)] \text{▲ (USP 1-Dec-2024)}$$

$M_{r1}$  = molecular weight of thiocyanate, 58.08

$M_{r2}$  = molecular weight of potassium thiocyanate, 97.18

$A_u$  = absorbance of the *Sample solution*

$W_u$  = weight of Erythromycin taken to prepare the *Sample solution* (mg)

$W_1$  = weight of the potassium thiocyanate taken to prepare *Standard ▲stock▲ (USP 1-Dec-2024) solution 1* (mg)

$A_1$  = absorbance of *Standard solution 1*

$W_2$  = weight of the potassium thiocyanate taken to prepare *Standard ▲stock▲ (USP 1-Dec-2024) solution 2* (mg)

$A_2$  = absorbance of *Standard solution 2*

**Acceptance criteria:** NMT 0.3%

**Change to read:**

• **ORGANIC IMPURITIES**

▲ **Diluent, Solution A, Solution B, Mobile phase, Standard solution 1, Standard solution 2, Sample solution, and Chromatographic system:** Proceed as directed in the Assay. Prepare the erythromycin solutions immediately before use.

**Diluted standard solution:** 0.04 mg/mL of [USP Erythromycin RS](#) prepared as follows. Dilute 1.0 mL of *Standard solution 1* to a 100-mL volumetric flask and dilute with *Diluent* to volume.

**System suitability solution:** Dissolve 4 mg of [USP Erythromycin System Suitability Mixture RS](#) in 1 mL of *Diluent*.

**System suitability**

**Sample:** *System suitability solution*

[NOTE—See [Table 2](#) for the relative retention times. Use the reference chromatogram provided with [USP Erythromycin System Suitability Mixture RS](#) and the chromatogram obtained with *System suitability solution* to identify the specified impurity peaks. Use the chromatogram obtained with *Standard solution 2* to identify erythromycin B and erythromycin C.]

**Suitability requirements**

**Peak-to-valley ratio:** NLT 1.5 for the ratio of the height of the pseudoerythromycin A enol ether peak to the height of the valley between the pseudoerythromycin A enol ether peak and the erythromycin B peak; NLT 2.0 for the ratio of the height of the erythromycin E peak to the height of the valley between erythromycin E peak and erythromycin A peak

**Resolution:** NLT 1.2 between 3"-N-demethylerythromycin A and erythromycin C

**Analysis**

**Samples:** *Standard solution 2, Sample solution, and Diluted standard solution*

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

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

$r_U$  = peak response of any individual impurity (the peak other than erythromycin A, erythromycin B, and erythromycin C) from the *Sample solution*

$r_S$  = peak response of erythromycin A from the *Diluted standard solution*

$C_S$  = concentration of [USP Erythromycin RS](#) in the *Diluted standard solution* (mg/mL)

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

$P$  = percentage of erythromycin A in [USP Erythromycin RS](#)

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

**Acceptance criteria:** See [Table 2](#). The reporting threshold is 0.2%.

**Table 2**

Name	Relative Retention Time	Relative Response Factor	Acceptance Criteria, NMT (%)
Erythromycin A N-oxide <sup>a</sup>	0.3	1	1.0
Erythromycin F <sup>b</sup>	0.4	1	2.0
3"-N-Demethylerythromycin A <sup>c</sup>	0.5	1	2.0
Erythromycin C	0.55	—	—
3"-N-Demethyl-3"-N-formyl erythromycin A <sup>d</sup>	0.63	9.1	0.4
Erythromycin E <sup>e</sup>	0.9	1	3.0
Erythromycin A	1.0	—	—
Anhydroerythromycin A <sup>f</sup>	1.61	0.5	1.0

Name	Relative Retention Time	Relative Response Factor	Acceptance Criteria, NMT (%)
Erythromycin B	1.75	—	—
Pseudoerythromycin A enol ether <sup>g</sup>	1.81	12.5	1.0
Erythromycin A enol ether <sup>h</sup>	2.3	12.5	1.0
Any other individual impurity	—	—	0.4
Total impurities	—	—	7.0

- <sup>a</sup> (3*R*,4*S*,5*S*,6*R*,7*R*,9*R*,11*R*,12*R*,13*S*,14*R*)-4-[(2,6-Dideoxy-3-*C*-methyl-3-*O*-methyl- $\alpha$ -*L*-*ribo*-hexopyranosyl)oxy]-14-ethyl-7,12,13-trihydroxy-3,5,7,9,11,13-hexamethyl-6-[(3,4,6-trideoxy-3-dimethylamino- $\beta$ -*D*-*xylo*-hexopyranosyl)oxy]oxacyclotetradecane-2,10-dione *N*-oxide.
- <sup>b</sup> (3*R*,4*S*,5*S*,6*R*,7*R*,9*R*,11*R*,12*R*,13*S*,14*R*)-4-[(2,6-Dideoxy-3-*C*-methyl-3-*O*-methyl- $\alpha$ -*L*-*ribo*-hexopyranosyl)oxy]-14-ethyl-7,12,13-trihydroxy-3-hydroxymethyl-5,7,9,11,13-pentamethyl-6-[(3,4,6-trideoxy-3-dimethylamino- $\beta$ -*D*-*xylo*-hexopyranosyl)oxy]oxacyclotetradecane-2,10-dione.
- <sup>c</sup> (3*R*,4*S*,5*S*,6*R*,7*R*,9*R*,11*R*,12*R*,13*S*,14*R*)-4-[(2,6-Dideoxy-3-*C*-methyl-3-*O*-methyl- $\alpha$ -*L*-*ribo*-hexopyranosyl)oxy]-14-ethyl-7,12,13-trihydroxy-3,5,7,9,11,13-hexamethyl-6-[(3,4,6-trideoxy-3-methylamino- $\beta$ -*D*-*xylo*-hexopyranosyl)oxy]oxacyclotetradecane-2,10-dione.
- <sup>d</sup> (3*R*,4*S*,5*S*,6*R*,7*R*,9*R*,11*R*,12*R*,13*S*,14*R*)-4-[(2,6-Dideoxy-3-*C*-methyl- $\alpha$ -*L*-*ribo*-hexopyranosyl)oxy]-14-ethyl-7,12,13-trihydroxy-3,5,7,9,11,13-hexamethyl-6-[(3,4,6-trideoxy-3-formylmethylamino- $\beta$ -*D*-*xylo*-hexopyranosyl)oxy]oxacyclotetradecane-2,10-dione.
- <sup>e</sup> (2*S*,4*aR*,4'*R*,5'*S*,6'*S*,7*R*,8*S*,9*R*,10*R*,12*R*,14*R*,15*R*,16*S*,16*aS*)-7-Ethyl-5',8,9,14-tetrahydroxy-4'-methoxy-4',6',8,10,12,14,16-heptamethyl-15-{[3,4,6-trideoxy-3-(dimethylamino)- $\beta$ -*D*-*xylo*-hexopyranosyl]oxy}hexadecahydrospiro[5*H*,11*H*-1,3-dioxino[5,4-*c*]oxacyclotetradecin-2,2'-pyrane]-5,11-dione.
- <sup>f</sup> (1*S*,2*R*,3*R*,4*S*,5*R*,8*R*,9*S*,10*S*,11*R*,12*R*,14*R*)-9-[(2,6-Dideoxy-3-*C*-methyl-3-*O*-methyl- $\alpha$ -*L*-*ribo*-hexopyranosyl)oxy]-5-ethyl-3-hydroxy-2,4,8,10,12,14-hexamethyl-11-[(3,4,6-trideoxy-3-(dimethylamino)- $\beta$ -*D*-*xylo*-hexopyranosyl)oxy]-6,15,16-trioxatricyclo[10.2.1.1<sup>1,4</sup>]hexadecan-7-one.
- <sup>g</sup> (2*R*,3*R*,6*R*,7*S*,8*S*,9*R*,10*R*)-7-[(2,6-Dideoxy-3-*C*-methyl-3-*O*-methyl- $\alpha$ -*L*-*ribo*-hexopyranosyl)oxy]-3-[(1*R*,2*R*)-1,2-dihydroxy-1-methylbutyl]-2,6,8,10,12-pentamethyl-9-[(3,4,6-trideoxy-3-(dimethylamino)- $\beta$ -*D*-*xylo*-hexopyranosyl)oxy]-4,13-dioxabicyclo[8.2.1]tridec-1(12)-en-5-one.
- <sup>h</sup> (2*R*,3*R*,4*S*,5*R*,8*R*,9*S*,10*S*,11*R*,12*R*)-9-[(2,6-Dideoxy-3-*C*-methyl-3-*O*-methyl- $\alpha$ -*L*-*ribo*-hexopyranosyl)oxy]-5-ethyl-3,4-dihydroxy-2,4,8,10,12,14-hexamethyl-11-[(3,4,6-trideoxy-3-(dimethylamino)- $\beta$ -*D*-*xylo*-hexopyranosyl)oxy]-6,15-dioxabicyclo[10.2.1]pentadec-1(14)-en-7-one.

▲ (USP 1-Dec-2024)

**SPECIFIC TESTS**

Delete the following:

- ▲ [OPTICAL ROTATION \(781S\)](#), [Procedures](#), [Specific Rotation](#) ▲ (USP 1-Dec-2024)

Change to read:

- [WATER DETERMINATION \(921\)](#), [Method I](#)

**Sample solution:** Use ▲ (USP 1-Dec-2024) [methanol](#) containing 10% of [imidazole](#) in place of [methanol](#) in the titration vessel.

**Acceptance criteria:** NMT ▲ 6.5% ▲ (USP 1-Dec-2024)

- [CRYSTALLINITY \(695\)](#): Meets the requirements

**ADDITIONAL REQUIREMENTS**

Change to read:

- **PACKAGING AND STORAGE:** ▲ Preserve in hermetic containers, and protect from light. ▲ (USP 1-Dec-2024)

Change to read:

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

[USP Erythromycin RS](#)

[USP Erythromycin B RS](#)

▲ (USP 1-Dec-2024)

[USP Erythromycin C RS](#)

- ▲ [USP Erythromycin System Suitability Mixture RS](#) ▲ (USP 1-Dec-2024)

Topic/Question	Contact	Expert Committee
ERYTHROMYCIN	<a href="#">Julie Zhang</a> Associate Science & Standards Liaison	BIO42020 Biologics Monographs 4 - Antibiotics

Chromatographic Database Information: [Chromatographic Database](#)

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