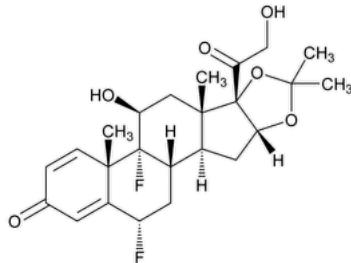


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## Fluocinolone Acetonide



$C_{24}H_{30}F_2O_6$  (anhydrous) 452.49

Dihydrate 488.53

Pregna-1,4-diene-3,20-dione, 6,9-difluoro-11,21-dihydroxy-16,17-[(1-methylethylidene)bis(oxyl)], (6 $\alpha$ ,11 $\beta$ ,16 $\alpha$ )-;

6 $\alpha$ ,9-Difluoro-11 $\beta$ ,16 $\alpha$ ,17,21-tetrahydroxypregna-1,4-diene-3,20-dione, cyclic 16,17-acetal with acetone CAS RN®: 67-73-2; UNII: 0CD5FD6S2M.

### DEFINITION

Fluocinolone Acetonide is anhydrous or contains two molecules of water of hydration. It contains NLT 97.0% and NMT 102.0% of fluocinolone acetonide ( $C_{24}H_{30}F_2O_6$ ), calculated on the dried basis.

### IDENTIFICATION

- A. SPECTROSCOPIC IDENTIFICATION TESTS (197), Infrared Spectroscopy: 197K:** If a difference appears, dissolve portions of both the sample and the USP Reference Standard in ethyl acetate, evaporate to dryness, and repeat the test on the residues.
- 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

**Mobile phase:** Acetonitrile, tetrahydrofuran, and water (13:10:77)

**Diluent:** Acetonitrile and tetrahydrofuran (13:10)

**Standard solution:** 0.2 mg/mL of [USP Fluocinolone Acetonide RS](#), prepared as follows. Transfer a suitable amount of [USP Fluocinolone Acetonide RS](#) to a suitable volumetric flask, and dissolve in a volume of *Diluent* equal to 23% of the flask volume. Dilute with water to volume.

**Sample solution:** 0.2 mg/mL of Fluocinolone Acetonide, prepared as follows. Transfer a suitable amount of Fluocinolone Acetonide to a suitable volumetric flask, and dissolve in a volume of *Diluent* equal to 23% of the flask volume. Dilute with water to volume.

#### Chromatographic system

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

**Mode:** LC

**Detector:** UV 254 nm

**Column:** 4.6-mm  $\times$  10-cm; 5- $\mu$ m packing L1

**Flow rate:** 2.5 mL/min

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

#### System suitability

**Sample:** *Standard solution*

**Suitability requirements**

**Tailing factor:** NMT 1.5

**Relative standard deviation:** NMT 3.0%

#### Analysis

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

Calculate the percentage of fluocinolone acetonide ( $C_{24}H_{30}F_2O_6$ ) in the portion of Fluocinolone Acetonide 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 Fluocinolone Acetonide RS](#) in the *Standard solution* (mg/mL)

$C_u$  = concentration of Fluocinolone Acetonide in the *Sample solution* (mg/mL)

**Acceptance criteria:** 97.0%–102.0% on the dried basis

**Change to read:**

- **ORGANIC IMPURITIES**

Protect the solutions from light throughout the test.

**Mobile phase:** Acetonitrile and water (45:55), prepared as follows. Mix 450 mL of acetonitrile and 500 mL of water, and allow to equilibrate.

Add water to make 1000 mL.

**System suitability solution:** 0.25 mg/mL each of [USP Fluocinolone Acetonide RS](#) and [USP Triamcinolone Acetonide RS](#), prepared as follows.

Transfer suitable amounts of [USP Fluocinolone Acetonide RS](#) and [USP Triamcinolone Acetonide RS](#) to a suitable volumetric flask. Dissolve in 45% of the flask volume of acetonitrile, and dilute with water to volume.

**Standard solution:** 0.025 mg/mL of [USP Fluocinolone Acetonide RS](#) in acetonitrile

**Sample solution:** 2.5 mg/mL of Fluocinolone Acetonide in acetonitrile

**Chromatographic system**

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

**Mode:** LC

**Detector:** UV 238 nm

**Column:** 4.6-mm × 25-cm; 5-μm packing L1

**Flow rate:** 1 mL/min

**Injection volume:** 20 μL

**Run time:** 4 times the retention time of fluocinolone acetonide

**System suitability**

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

[NOTE—The relative retention times for triamcinolone acetonide and fluocinolone acetonide are 0.85 and 1.0, respectively.]

**Suitability requirements**

**Resolution:** NLT 2.0 between triamcinolone acetonide and fluocinolone acetonide, System suitability solution

**Relative standard deviation:** NMT 10.0%, Standard solution

**Analysis**

**Samples:** Standard solution and Sample solution

Calculate the percentage of each impurity in the portion of Fluocinolone Acetonide taken:

$$\text{Result} = (r_u/r_s) \times (C_s/C_u) \times 100$$

$r_u$  = peak response of any impurity from the *Sample solution*

$r_s$  = peak response of fluocinolone acetonide from the *Standard solution*

$C_s$  = concentration of [USP Fluocinolone Acetonide RS](#) in the *Standard solution* (mg/mL)

$C_u$  = concentration of Fluocinolone Acetonide in the *Sample solution* (mg/mL)

**Acceptance criteria**

**Any individual impurity:** NMT 1%; NMT one such peak is greater than 0.5%.

**Total impurities:** NMT 2.5%. Disregard any peak below 0.05% of the peak area of fluocinolone acetonide from the ▲*Sample solution* ▲ (ERR 1-

Jun-2022) .

**SPECIFIC TESTS**

- [OPTICAL ROTATION, Specific Rotation \(781S\)](#)

**Sample solution:** 10 mg/mL in methanol

**Acceptance criteria:** +98° to +108°

- [LOSS ON DRYING \(731\)](#)

**Analysis:** Dry under vacuum at 105° for 3 h.

**Acceptance criteria:** NMT 1.0% for anhydrous Fluocinolone Acetonide; NMT 8.5% for hydrous Fluocinolone Acetonide

**ADDITIONAL REQUIREMENTS**

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

- **LABELING:** Label it to indicate whether it is anhydrous or hydrous.

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

[USP Fluocinolone Acetonide RS](#)

[USP Triamcinolone Acetonide RS](#)

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

Topic/Question	Contact	Expert Committee
FLUOCINOLONE ACETONIDE	<a href="#">Documentary Standards Support</a>	SM52020 Small Molecules 5
REFERENCE STANDARD SUPPORT	RS Technical Services <a href="mailto:RSTECH@usp.org">RSTECH@usp.org</a>	SM52020 Small Molecules 5

**Chromatographic Database Information:** [Chromatographic Database](#)

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