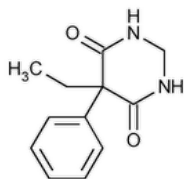


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## Primidone



$C_{12}H_{14}N_2O_2$  218.25

4,6(1*H*,5*H*)-Pyrimidinedione, 5-ethyldihydro-5-phenyl-;

5-Ethyldihydro-5-phenyl-4,6(1*H*,5*H*)-pyrimidinedione CAS RN®: 125-33-7; UNII: 13AFD7670Q.

### DEFINITION

Primidone contains NLT 98.0% and NMT 102.0% of  $C_{12}H_{14}N_2O_2$ , calculated on the dried basis.

### IDENTIFICATION

#### Change to read:

- **A.** ▲ [SPECTROSCOPIC IDENTIFICATION TESTS \(197\)](#), [Infrared Spectroscopy: 197K](#) ▲ (CN 1-MAY-2020) : If a difference appears, dissolve portions of both the sample and the Reference Standard in alcohol, evaporate the solutions to dryness, and repeat the tests.
- **B.** The retention time of the major peak in the *Sample solution* corresponds to that of the *Standard solution*, as obtained in the *Assay*.

### ASSAY

#### PROCEDURE

**Solution A:** 6.8 g/L of monobasic potassium phosphate

**Mobile phase:** Methanol, tetrahydrofuran, and *Solution A* (35:0.5:65)

**Diluent:** Methanol and water (35:65)

**Standard stock solution:** 0.5 mg/mL of [USP Primidone RS](#) in methanol

**Standard solution:** 0.05 mg/mL of [USP Primidone RS](#) in *Diluent*, prepared from the *Standard stock solution*

**Sample stock solution:** 0.5 mg/mL of Primidone in methanol

**Sample solution:** 0.05 mg/mL of Primidone from the *Sample stock solution* diluted with *Diluent*

#### Chromatographic system

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

**Mode:** LC

**Detector:** UV 220 nm

**Column:** 4.6-mm × 10-cm; 3-μm packing L1

**Column temperature:** 35°

**Flow rate:** 1.3 mL/min

**Injection size:** 20 μL

#### System suitability

**Sample:** *Standard solution*

#### Suitability requirements

**Column efficiency:** NLT 3000 theoretical plates

**Tailing factor:** NMT 2.0

**Relative standard deviation:** NMT 2.0%

#### Analysis

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

Calculate the percentage of  $C_{12}H_{14}N_2O_2$  in the portion of Primidone 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 Primidone RS](#) in the *Standard solution* (mg/mL)

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

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

## IMPURITIES

### INORGANIC IMPURITIES

- **RESIDUE ON IGNITION (281):** NMT 0.2%

### ORGANIC IMPURITIES

#### • PROCEDURE

**Solution A:** 1.36 g/L of monobasic potassium phosphate

**Solution B:** Methanol

**Mobile phase:** See the gradient table below.

Time (min)	Solution A (%)	Solution B (%)
0	75	25
1.0	75	25
6.0	40	60
8.0	40	60
8.5	75	25
10.0	75	25

**Standard stock solution:** 1000 µg/mL of [USP Primidone RS](#) in methanol

**Standard solution:** 1 µg/mL of [USP Primidone RS](#) in methanol, prepared from the *Standard stock solution* in methanol

**System suitability solution:** 1000 µg/mL of [USP Primidone RS](#) and 10 µg/mL each of [USP Phenobarbital RS](#) and [USP Primidone Related Compound C RS](#). Prepare by diluting weighed quantities of [USP Phenobarbital RS](#) and [USP Primidone Related Compound C RS](#) with the *Standard stock solution*.

**Sample solution:** 1000 µg/mL of Primidone in methanol

### Chromatographic system

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

**Mode:** LC

**Detector:** UV 215 nm

**Column:** 4.6-mm × 10-cm; monolithic packing L1

**Flow rate:** 3.2 mL/min

**Injection size:** 10 µL

### System suitability

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

#### Suitability requirements

**Resolution:** NLT 2.5 between phenobarbital and primidone related compound C, *System suitability solution*

**Tailing factor:** NMT 2.0 for primidone, *Standard solution*

**Relative standard deviation:** NMT 5.0% for primidone, *Standard solution*

### Analysis

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

Identify the impurities using the relative retention times listed in [Impurity Table 1](#). Calculate the percentage of each impurity in the portion of sample taken:

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

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

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

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

$C_U$  = concentration of Primidone in the *Sample solution* (µg/mL)

$F$  = relative response factor (see [Impurity Table 1](#))

#### Acceptance criteria

**Individual impurities:** See [Impurity Table 1](#).

**Total impurities:** NMT 0.5%

[NOTE—Disregard impurity peaks below 0.05%.]

**Impurity Table 1**

Name	Relative Retention Time	Relative Response Factor	Acceptance Criteria, NMT (%)
Primidone related compound <a href="#">A<sup>a</sup></a>	0.5	0.67	0.1
Phenobarbital	1.4	1.0	0.1
Primidone related compound <a href="#">C<sup>b</sup></a>	1.6	0.67	0.1
2-Cyano-2-phenylbutyramide	1.8	0.71	0.1
2-Phenylbutyric acid	2.0	0.77	0.1
Phenylpropylprimidone <a href="#">C<sup>c</sup></a>	2.8	1.0	0.3
Any unspecified impurity	—	1.0	0.1

<sup>a</sup> 2-Ethyl-2-phenylmalonamide (2-ethyl-2-phenylpropanediamide).

<sup>b</sup> 2-Phenylbutyramide.

<sup>c</sup> 5-Ethyl-5-phenyl-2-(1-phenylpropyl) dihydropyrimidine-4,6(1*H*,5*H*)-dione.

#### SPECIFIC TESTS

- **Loss on Drying (731):** Dry a sample at 105° for 2 h: it loses NMT 0.5% of its weight.

#### ADDITIONAL REQUIREMENTS

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

- **USP REFERENCE STANDARDS (11).**

[USP Phenobarbital RS](#)

[USP Primidone RS](#)

[USP Primidone Related Compound C RS](#)

2-Phenylbutyramide.

$C_{10}H_{13}NO$  163.22

Topic/Question	Contact	Expert Committee
PRIMIDONE	<a href="#">Documentary Standards Support</a>	SM42020 Small Molecules 4
REFERENCE STANDARD SUPPORT	RS Technical Services <a href="mailto:RSTECH@usp.org">RSTECH@usp.org</a>	SM42020 Small Molecules 4

Chromatographic Database Information: [Chromatographic Database](#)

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