

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:

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

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* ($\mu\text{g/mL}$)

C_u = concentration of Primidone in the *Sample solution* ($\mu\text{g/mL}$)

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 ^a	0.5	0.67	0.1
Phenobarbital	1.4	1.0	0.1
Primidone related compound C ^b	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 ^c	2.8	1.0	0.3
Any unspecified impurity	—	1.0	0.1

^a 2-Ethyl-2-phenylmalonamide (2-ethyl-2-phenylpropanediamide).

^b 2-Phenylbutyramide.

^c 5-Ethyl-5-phenyl-2-(1-phenylpropyl)dihydropyrimidine-4,6(1H,5H)-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.

$\text{C}_{10}\text{H}_{13}\text{NO}$ 163.22

Topic/Question	Contact	Expert Committee
PRIMIDONE	Documentary Standards Support	SM42020 Small Molecules 4
REFERENCE STANDARD SUPPORT	RS Technical Services RSTECH@usp.org	SM42020 Small Molecules 4

Chromatographic Database Information: [Chromatographic Database](#)

Most Recently Appeared In:

Pharmacopeial Forum: Volume No. PF 35(3)

Current DocID: GUID-E8398E86-B7AB-4A12-9666-1D751FBC5527_2_en-US

DOI: https://doi.org/10.31003/USPNF_M69110_02_01

DOI ref: [ka9el](#)

OFFICIAL