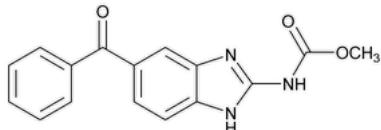


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Mebendazole



$C_{16}H_{13}N_3O_3$ 295.29

Carbamic acid, (5-benzoyl-1H-benzimidazol-2-yl), methyl ester;
Methyl 5-benzoyl-2-benzimidazolcarbamate CAS RN®: 31431-39-7; UNII: 81G6I5V05I.

DEFINITION

Mebendazole contains NLT 98.0% and NMT 102.0% of mebendazole ($C_{16}H_{13}N_3O_3$), calculated on the dried basis.

IDENTIFICATION

Change to read:

- A. ▲ [SPECTROSCOPIC IDENTIFICATION TESTS \(197\), Infrared Spectroscopy: 197K](#) ▲ (CN 1-MAY-2020)
- 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

Solution A: 7.5 g/L of ammonium acetate

Solution B: Acetonitrile

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

Table 1

Time (min)	Solution A (%)	Solution B (%)
0	80	20
15	70	30
20	10	90
25	10	90
26	80	20
30	80	20

Diluent: Acetonitrile and water (50:50)

System suitability solution: 50 µg/mL of [USP Mebendazole RS](#) and 2.5 µg/mL of [USP Mebendazole Related Compound D RS](#) in *Diluent*.

Sonicate in *Diluent* using 80% of the final volume at 45°–50° for 10 min. Dilute with *Diluent* to final volume.

Standard solution: 0.05 mg/mL of [USP Mebendazole RS](#) in *Diluent*. Sonicate in *Diluent* using 80% of the final volume at 45°–50° for 10 min.

Dilute with *Diluent* to final volume.

Sample solution: 0.05 mg/mL of Mebendazole in *Diluent*. Sonicate in *Diluent* using 80% of the final volume at 45°–50° for 10 min. Dilute with *Diluent* to final volume.

Chromatographic system

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

Mode: LC

Detector: UV 250 nm

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

Column temperature: 40°**Flow rate:** 1.2 mL/min**Injection volume:** 10 µL**System suitability****Samples:** System suitability solution and Standard solution

[NOTE—The relative retention times of mebendazole and mebendazole related compound D are 1.0 and 1.2, respectively.]

Suitability requirements**Resolution:** NLT 5.0 between mebendazole and mebendazole related compound D, *System suitability solution***Tailing factor:** NMT 2.0, *Standard solution***Relative standard deviation:** NMT 0.73%, *Standard solution***Analysis****Samples:** Standard solution and Sample solutionCalculate the percentage of mebendazole ($C_{16}H_{13}N_3O_3$) in the portion of Mebendazole taken:

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

 r_U = peak response of mebendazole from the *Sample solution* r_S = peak response of mebendazole from the *Standard solution* C_S = concentration of [USP Mebendazole RS](#) in the *Standard solution* (mg/mL) C_U = concentration of Mebendazole in the *Sample solution* (mg/mL)**Acceptance criteria:** 98.0%–102.0% on the dried basis**IMPURITIES**• [RESIDUE ON IGNITION \(281\)](#): NMT 0.1%• **ORGANIC IMPURITIES****Solution A, Solution B, Mobile phase, Diluent, System suitability solution, Sample solution, and Chromatographic system:** Prepare as directed in the Assay.**Standard solution:** 2.5 µg/mL of [USP Mebendazole RS](#) in *Diluent***System suitability****Sample:** System suitability solution**Suitability requirements****Resolution:** NLT 5.0 between the mebendazole and mebendazole related compound D peaks**Relative standard deviation:** NMT 1.0% for the mebendazole peak and NMT 5.0% for the mebendazole related compound D peak**Analysis****Samples:** Sample solution and Standard solution

Calculate the percentage of each impurity in the portion of Mebendazole taken:

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

 r_U = peak response of each impurity from the *Sample solution* r_S = peak response of mebendazole from the *Standard solution* C_S = concentration of [USP Mebendazole RS](#) in the *Standard solution* (mg/mL) C_U = concentration of Mebendazole in the *Sample solution* (mg/mL) F = relative response factor (see [Table 2](#))**Acceptance criteria:** See [Table 2](#). Disregard peaks less than 0.05%.**Table 2**

Name	Relative Retention Time	Relative Response Factor	Acceptance Criteria, NMT (%)
2-Amino mebendazole ^a	0.46	1.0	0.25
2-Hydroxy mebendazole ^b	0.53	1.0	0.25

Name	Relative Retention Time	Relative Response Factor	Acceptance Criteria, NMT (%)
2-Amino-1-methyl mebendazole ^c	0.67	1.0	0.25
Mebendazole	1.0	—	—
Mebendazole related compound D	1.1	1.0	0.25
Ethyl mebendazole ^d	1.3	1.0	0.25
Toluoyl mebendazole ^e	1.4	1.0	0.25
Mebendazole dimer ^f	1.6	0.71	0.5
Any other unspecified impurity	—	1.0	0.10
Total impurities	—	—	1.0

^a 2-Amino-5-benzoylbenzimidazole.^b 5-Benzoyl-2-hydroxybenzimidazole.^c 2-Amino-5-benzoyl-1-methylbenzimidazole.^d Ethyl (5-benzoyl-1*H*-benzimidazol-2-yl)carbamate.^e Methyl 5-(4-toluoyl)-1*H*-benzimidazol-2-ylcarbamate.^f 1,3-Bis(5-benzoylbenzimidazol-2-yl)urea.**SPECIFIC TESTS****• LOSS ON DRYING (731)****Analysis:** Dry at 105° for 4 h.**Acceptance criteria:** NMT 0.5%**ADDITIONAL REQUIREMENTS****• PACKAGING AND STORAGE:** Preserve in well-closed containers.**• USP REFERENCE STANDARDS (11)**[USP Mebendazole RS](#)[USP Mebendazole Related Compound D RS](#)

Methyl 5-benzoyl-1-methylbenzimidazol-2-ylcarbamate.

 $C_{17}H_{15}N_3O_3$ 309.32**Auxiliary Information** - Please [check for your question in the FAQs](#) before contacting USP.

Topic/Question	Contact	Expert Committee
MEBENDAZOLE	Documentary Standards Support	SM12020 Small Molecules 1

Chromatographic Database Information: [Chromatographic Database](#)**Most Recently Appeared In:**

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