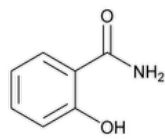


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Salicylamide



$C_7H_7NO_2$ 137.14
Benzamide, 2-hydroxy-;
2-Hydroxybenzamide CAS RN®: 65-45-2; UNII: EM8BM710ZC.

DEFINITION
Salicylamide contains NLT 98.0% and NMT 102.0% of salicylamide ($C_7H_7NO_2$), calculated on the anhydrous 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 Assay.

ASSAY

- **PROCEDURE**
Solution A: 0.1% Formic acid in water
Solution B: Acetonitrile
Mobile phase: See [Table 1](#).

Table 1

| Time (min) | Solution A (%) | Solution B (%) |
|------------|----------------|----------------|
| 0.0 | 92 | 8 |
| 6.0 | 55 | 45 |
| 6.5 | 5 | 95 |
| 8.0 | 5 | 95 |
| 8.01 | 92 | 8 |
| 12.0 | 92 | 8 |

Diluent: Acetonitrile and water (50:50)
System suitability solution: 1.0 mg/mL of [USP Salicylamide RS](#) and 0.001 mg/mL of [USP Salicylic Acid Related Compound B RS](#) in *Diluent*
Standard solution: 0.2 mg/mL of [USP Salicylamide RS](#) in *Diluent*
Sample solution: 0.2 mg/mL of Salicylamide in *Diluent*. Pass through a suitable filter of 0.2-µm pore size.
Chromatographic system
(See [Chromatography \(621\), System Suitability.](#))
Mode: LC

Detector: UV 300 nm**Column:** 2.1-mm × 10-cm; 1.8-μm packing L1**Column temperature:** 35°**Flow rate:** 0.3 mL/min**Injection volume:** 1 μL**System suitability****Samples:** *System suitability solution* and *Standard solution***Suitability requirements****Resolution:** NLT 1.5 between salicylic acid related compound B and salicylamide, *System suitability solution***Relative standard deviation:** NMT 0.73%, *Standard solution***Analysis****Samples:** *Standard solution* and *Sample solution*Calculate the percentage of salicylamide ($C_7H_7NO_2$) in the portion of Salicylamide 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 Salicylamide RS](#) in the *Standard solution* (mg/mL) C_U = concentration of Salicylamide in the *Sample solution* (mg/mL)**Acceptance criteria:** 98.0%–102.0% on the anhydrous basis**IMPURITIES**• [RESIDUE ON IGNITION \(281\)](#): NMT 0.1%• **ORGANIC IMPURITIES****Solution A, Solution B, Mobile phase, Diluent, System suitability solution, and Chromatographic system:** Proceed as directed in the Assay.**Standard solution:** 0.001 mg/mL each of [USP Salicylamide RS](#) and [USP Salicylic Acid RS](#) in *Diluent***Sample solution:** 1.0 mg/mL of Salicylamide in *Diluent*. Pass through a suitable filter of 0.2-μm pore size.**System suitability****Samples:** *System suitability solution* and *Standard solution***Suitability requirements****Resolution:** NLT 1.5 between salicylic acid related compound B and salicylamide, *System suitability solution***Relative standard deviation:** NMT 2.8%, *Standard solution***Analysis****Samples:** *Standard solution* and *Sample solution*

Calculate the percentage of salicylic acid in the portion of Salicylamide taken:

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

 r_U = peak response of the salicylic acid peak from the *Sample solution* r_S = peak response of the salicylic acid peak from the *Standard solution* C_S = concentration of [USP Salicylic Acid RS](#) in the *Standard solution* (mg/mL) C_U = concentration of Salicylamide in the *Sample solution* (mg/mL)

Calculate the percentage of each unspecified impurity in the portion of Salicylamide taken:

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

 r_U = peak response of each unspecified impurity from the *Sample solution* r_S = peak response of salicylamide from the *Standard solution* C_S = concentration of [USP Salicylamide RS](#) in the *Standard solution* (mg/mL)

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

Acceptance criteria: See [Table 2](#). Disregard any impurity peak less than 0.05%.

Table 2

| Name | Relative Retention Time | Acceptance Criteria, NMT (%) |
|--|-------------------------|------------------------------|
| Salicylic acid related compound B ^a | 0.93 | — |
| Salicylamide | 1.00 | — |
| Salicylic acid | 1.33 | 0.1 |
| Individual unspecified impurity | — | 0.10 |
| Total impurities | — | 1 |

^a For identification only.

SPECIFIC TESTS

- **WATER DETERMINATION, *Method I* (921):** NMT 0.5%

ADDITIONAL REQUIREMENTS

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

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

[USP Salicylamide RS](#)

[USP Salicylic Acid RS](#)

[USP Salicylic Acid Related Compound B RS](#)

4-Hydroxyisophthalic acid.

$C_8H_6O_5$ 182.13

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

| Topic/Question | Contact | Expert Committee |
|----------------------------|---|---------------------------|
| SALICYLAMIDE | Documentary Standards Support | SM22020 Small Molecules 2 |
| REFERENCE STANDARD SUPPORT | RS Technical Services RSTECH@usp.org | SM22020 Small Molecules 2 |

Chromatographic Database Information: [Chromatographic Database](#)

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