

HPLC Method Description for Identity, Assay and Related Substances of PNDa01 and PNDa06-HCI

| Project | Pyronaridine_INV-054926 |
|----------------|---------------------------------|
| Compound | PNDa01 and PNDa06-HCI |
| Purpose | Method Description |
| Category | Methods |
| Substance Type | Intermediate |
| Report ID | INV_054926_HPLC_M1R Version 1.0 |

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Distribution

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1. Objective

This method 'INV_054926_HPLC_M1R' for intermediates PNDa01 and PNDa06-HCl of Pyronaridine (INV-054926) project is developed by HPLC. The parameters of the reversed phase HPLC method suitable for identity, assay, and related substances of PNDa01 and PNDa06-HCl shall be described in this document.

Related reports:

INV_054926_HPLC_M1: HPLC Method Description for In-process Control of Intermediates INV_054926_HPLC_V1: HPLC Method Limited Validation for PNDa01 and PNDa06-HCl (non-GMP)

2. Summary and conclusion

The method is applied for the HPLC testing of PNDa01 and PNDa06-HCl (identity, assay and related substances). This method is based on reversed phase liquid chromatography with UV detection and gradient elution using a Waters Atlantis T3, 3µm, 150 x 4.6mm HPLC column. **Table 1** Structure, Retention time and RRT of PNDa01 and its related substances

| Compound | Compound Structure | | RRT |
|-------------------------------------|-----------------------------|--------------|----------|
| 6-methoxy-3- aminepyridine (SM2) | NH ₂ | ca. 6.8 min | ca. 0.35 |
| PNDa01 impurity 2 | a01 impurity 2 са. 17.4 min | | ca. 0.90 |
| 2,4-Dichlorobenzoic acid (SM1) | СІСІОН | ca. 18.0 min | ca. 0.94 |
| PNDa01 impurity 1 | СІ ОН | ca. 18.2 min | ca. 0.95 |
| PNDa01 | Da01 | | 1.00 |
| PNDa01 impurity 3 | | ca. 20.5 min | ca. 1.07 |



| Compound | Structure | Retention time (RT) | RRT |
|------------------------|------------------|---------------------|----------|
| PNDa06-HCI | | ca. 8.1 min | 1.00 |
| Acetaminophen (SM5) | OH HN OCH3 | ca. 8.5 min | ca. 1.04 |

 Table 2
 Structure, Retention time and RRT of PNDa06-HCl and its related substances

Example chromatograms and extracted HPLC-PDA spectra of PNDa01 and PNDa06-HCl are given in Section 4.

Specificity, LOQ, Linearity, Accuracy, Repeatability was performed and reported in report 'INV_054926_HPLC_V1'.

3. Experimental

Equivalent equipment or grade of materials can be used.

3.1. HPLC

- HPLC System: Quaternary pump module (e.g.: Waters Alliance 2695)
 PDA detector (e.g.: Waters Alliance 2998)
 Auto sampler (e.g.: Waters Alliance 2695)
 - Column oven (e.g.: Waters Alliance 2695)
- Empower-control and integration software or equivalent
- Column: Waters Atlantis T3, 3µm, 150 x 4.6mm
- Flow rate: 1.0 mL/min
- Elution: Gradient mode
- Run time: 30.0 min
- Detection: 254 nm for PNDa01
- 278 nm for PNDa06-HCI

10 µL

- Injection:
- Column temp.: 35°C ± 5°C
- Auto sampler temp.: Room temperature
- Mobile phase (see section 3.3.2):
 - **A:** 0.1% TFA in Water
 - **B:** 0.1% TFA in Acetonitrile
- Diluent:

DMSO/Acetonitrile (50: 50 v/v) for PNDa01

Water/Acetonitrile (80: 20 v/v) for PNDa06-HCI

- Needle wash: Water/ Acetonitrile (50:50 v/v)
- Equilibration time: 7.0 min
- Gradient:

Table 3Gradient Table

| Time (min) | % A | % B |
|------------|-----------|------|
| 0.0 | 100.0 | 0.0 |
| 1.0 | 100.0 0.0 | |
| 13.0 | 65.0 | 35.0 |
| 20.0 | 1.0 | 99.0 |
| 25.0 | 1.0 | 99.0 |
| 25.1 | 100.0 | 0.0 |
| 30.0 | 100.0 | 0.0 |

3.2. Equipment and reagents

• Balance:

•

- e.g.: Mettler Toledo XP56
- Acetonitrile: HPLC grade, e.g.: Merck LiChrosolv
- Water: HPLC grade, e.g.: from Milipore ultra-pure water system
- TFA: HPLC grade, e.g.: Sigmer-Aldrich
- DMSO: HPLC grade, e.g.: Sigmer-Aldrich
- Glassware: 10, 50-mL volumetric flasks, 1L graduated cylinders
- Pipette: e.g.: 1.0 mL Pipette

3.3. Solutions

3.3.1. Diluent

Different volumes can be prepared as soon as the solvent ratio is the same.

PNDa01: DMSO/Acetonitrile (50:50 v/v).

PNDa06-HCI: Water/Acetonitrile (80: 20 v/v).

3.3.2. Mobile phase preparation

Preparation is described for a volume of 1 liter. Different volumes can be prepared as soon as the solvent ratio is the same.

Mobile phase A (0.1% TFA in Water):

In a suitable container, add 1000 mL of water and 1 mL of TFA. Mix well.

Mobile phase B (0.1% TFA in Acetonitrile):

In a suitable container, add 1000 mL of acetonitrile and 1 mL of TFA. Mix well.

3.3.3. Solution preparations

Other volumes and weigh-ins might be used as long as the final concentration remains the same. Min. weight of used balance must be considered during sample preparation.

3.3.3.1. Standard solutions

PNDa01:

Standard Solution 1 & 2 (conc.: 0.2 mg/ml):

Accurately weigh approx. 10 mg of PNDa01 reference standard into a 50-mL volumetric flask.

Dissolve and dilute to volume with diluent. Mix well.

Prepare in duplicate if needed.

Standard Solution 3 (0.05%, corresponding to 0.0001 mg/ml):

Transfer 0.5 mL of Standard Solution 1 into a 50 mL volumetric flask. Fill up to volume with diluent and mix well.

Transfer 0.5 mL of above solution into a 10 mL volumetric flask. Fill up to volume with diluent and mix.

PNDa06-HCI:

Standard Solution 1 & 2 (conc.: 2.0 mg/ml):

Accurately weigh approx. 20 mg of PNDa06-HCl reference standard into a 10-mL volumetric flask. Dissolve and dilute to volume with diluent. Mix well.

Prepare in duplicate if needed.

Standard Solution 3 (0.05%, corresponding to 0.001 mg/ml):

Transfer 0.5 mL of Standard Solution 1 into a 50 mL volumetric flask. Fill up to volume with diluent and mix well.

Transfer 0.5 mL of above solution into a 10 mL volumetric flask. Fill up to volume with diluent and mix.

3.3.3.2. Sample solutions

Number of sample preparations depends on the samples under analysis.

PNDa01:

Accurately weigh approx. 10 mg of PNDa01 sample into a 50-mL volumetric flask. Dissolve and dilute to volume with diluent. Mix well.

PNDa06-HCI:

Accurately weigh approx. 20 mg of PNDa06-HCl sample into a 10-mL volumetric flask. Dissolve and dilute to volume with diluent. Mix well.

3.4. Proposed injection sequence and system suitability test

| Table 4 | Proposed injection sequence and SST criteria | |
|---------|--|--|
| | | |

| Sample name | No. of injections ^[1] | SST acceptance criteria |
|------------------------------------|----------------------------------|--|
| Blank (diluent) |] + N ^[]] | No interference between the blank peaks and the components of interest |
| Standard Solution 3 (0.05%) | 1 | S/N ≥ 10 |
| Standard Solution 2 | 6 | %RSD (main peak area) ≤ 2 % |
| Standard Solution 1 | 1 | Recovery: 98% - 102% (6 injection Std 2 to be used as reference) |
| Sample solution prep.1 | 1 | N/A |
| Sample solution prep.2 | 1 | for Identification purpose just 1 sample preparation is required |
| Standard Solution 2 ^[2] | 1 | Recovery: 98% - 102% (6 injection Std 2 to be used as reference) |

^[1] Additional blanks may be run until an acceptable baseline is obtained. - ^[2] For multiple sample analysis, 1 injection of standard solution 2 is recommended every 6 sample preparation injections.

3.5. Calculation and Reporting

Calculations should be performed individually for each sample weighing. Only then should the calculation of the average result be performed.

3.5.1. Identification by HPLC

The main peak retention time of standard injections should not differ by more than 5% from the main peak retention time of the sample injections.

3.5.2. Purity by HPLC in %area

Calculate the purity using the following formula, for each sample preparation: 100 - %(area) Total impurities

Calculate the average of the 2 individual preparations by: (P1+P2) /2

Where:

Pi = PNDa01 or PNDa06-HCl purity % (area)

3.5.3. Related substances by HPLC: Total impurities in %area

Sum of the % area of all impurities (Report only the peaks for which the % area is not less than 0.05%).

3.5.4. Related substances by HPLC: Individual impurities in %area (by RRT)

Report all individual impurities \geq 0.05% (area) by their RRT. Analyses with two sample weighings: (A1+A2)/2 Where:

Ai = impurity peak % (area)

In case of the specified impurities, report:

- For impurity content below LOQ concentration, report "Less than 0.05 % (area)";

- If impurity is not detected, report "Not detected".

3.5.5. Assay by HPLC in %w/w

PNDa01:

Assay_{PNDa01} (%w/w) =
$$\frac{\text{Area}_{\text{sam}} \times W_{\text{std}} \times V_{\text{sam}}}{\text{Area}_{\text{std}} \times W_{\text{sam}} \times V_{\text{std}}} \times P_{\text{std}}$$

Where:

| Area _{sam} = | PNDa01 peak area obtained in the sample chromatogram |
|-----------------------|--|
| Area std= | Average PNDa01 peak area obtained for the 6 standard injections (std 2) |
| W std = | Standard weight (Std 2) |
| W sam = | Sample weight (mg) |
| V sam = | Volume of the flask used in the sample preparation |
| V std = | Volume of the flask used in the standard preparation (Std 2) |
| P _{std} = | Potency determined for the standard in used. Use value from CoA for PNDa01 |

PNDa06-HCI:

Assay free base (%w/w) =
$$\frac{\text{Area}_{\text{sam}} \times W_{\text{std}} \times V_{\text{sam}}}{\text{Area}_{\text{std}} \times W_{\text{sam}} \times V_{\text{std}}} \times P_{\text{std}}$$

Where:

| Area _{sam} = Area _{std} = | PNDa06 peak area obtained in the sample chromatogram Average PNDa06 peak area obtained for the 6 standard injections (std 2) |
|--|---|
| W std = | Standard weight (Std 2) |
| W s _{am} = | Sample weight (mg) |
| V _{sam} = | Volume of the flask used in the sample preparation |
| V std = | Volume of the flask used in the standard preparation (Std 2) |
| P _{std} = | Potency determined for the standard in used. Use value (free base) from CoA |
| | for PNDa06-HCI |
| Assay _{salt} | $(\%w/w) = Assay_{free base} \times \frac{MW_{salt}}{MW_{free base}}$ |

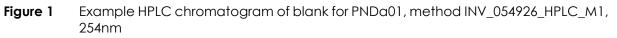
Where:

Assay free base = Potency determined for the free base of PNDa06 sample

MW salt = Molecular weight of PNDa06-HCl salt (390.35 g/mol)

MW free base = Molecular weight of PNDa06 free base (317.43 g/mol)

4. Figures



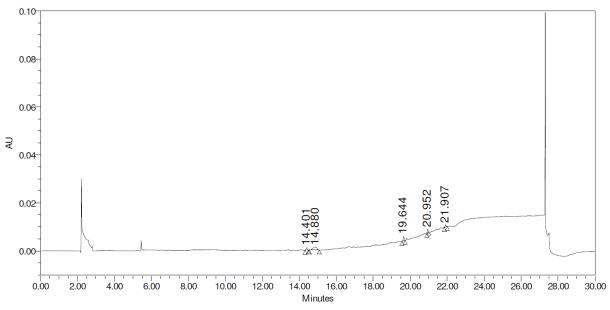
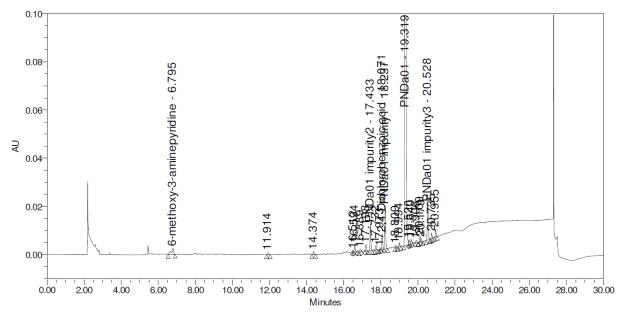
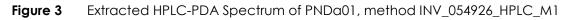


Figure 2 Example HPLC chromatogram of PNDa01 batch PHTANWARL-528-1, method INV_054926_HPLC_M1, 254nm





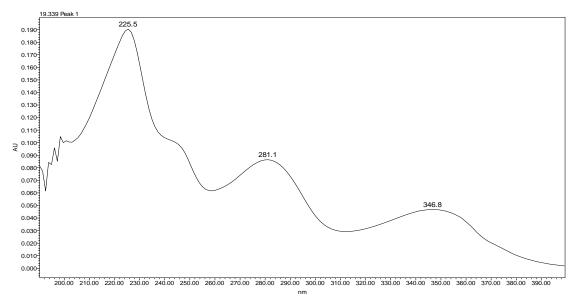
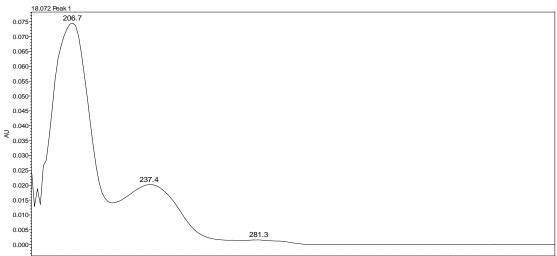
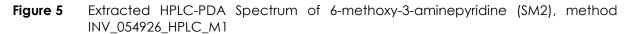
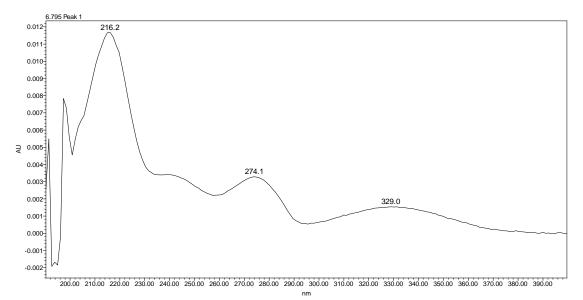


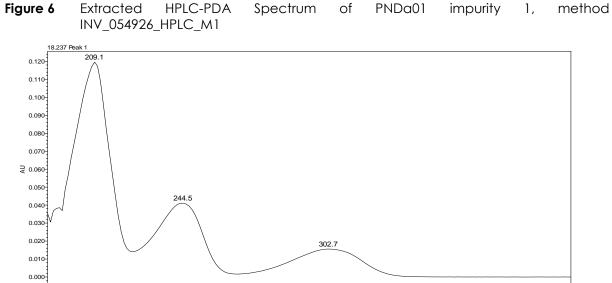
Figure 4 Extracted HPLC-PDA Spectrum of 2,4-Dichlorobenzoic acid (SM1), method INV_054926_HPLC_M1



200.00 210.00 220.00 230.00 240.00 250.00 260.00 270.00 280.00 290.00 300.00 310.00 320.00 330.00 340.00 350.00 360.00 370.00 380.00 390.00







200.00 210.00 220.00 230.00 240.00 250.00 260.00 270.00 280.00 290.00 300.00 310.00 320.00 330.00 340.00 360.00 360.00 370.00 380.00 390.00



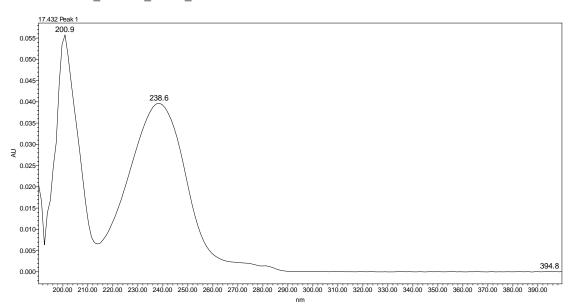
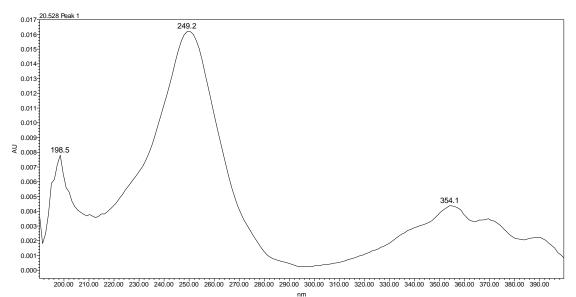


Figure 8 Extracted HPLC-PDA Spectrum of PNDa01 impurity 3, method INV_054926_HPLC_M1



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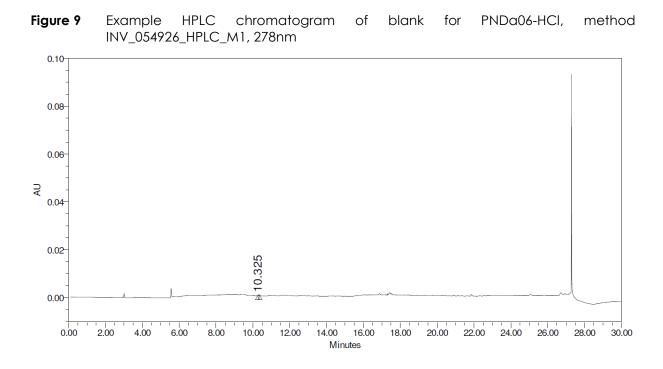
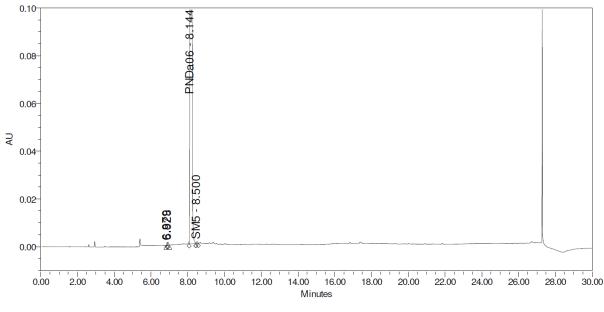


Figure 10 Example HPLC chromatogram of PNDa06-HCl batch PHTANWARL-528-1, method INV_054926_HPLC_M1, 278nm



| Name | Retention Time, min | RRT | Area | % Area |
|---------------------|---------------------|------|---------|--------|
| | 6.879 | 0.84 | 3443 | 0.07 |
| | 6.928 | 0.85 | 3577 | 0.07 |
| PNDa06 | 8.144 | 1.00 | 5067326 | 99.78 |
| Acetaminophen (SM5) | 8.500 | 1.04 | 4279 | 0.08 |

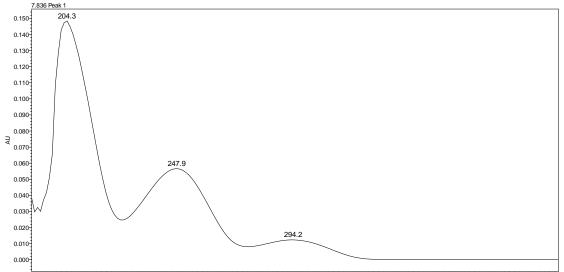


Figure 11 Extracted HPLC-PDA Spectrum of PNDa06-HCl, method INV_054926_HPLC_M1

