

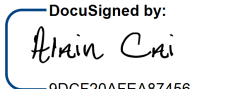
# HPLC Method Description for Identity, Assay and Related Substances of TGF-001


<b>Project</b>	<b>Pyronaridine_INV-054926</b>
<b>Compound</b>	<b>TGF-001</b>
<b>Purpose</b>	<b>Method Description</b>
<b>Category</b>	<b>Methods</b>
<b>Substance Type</b>	<b>Drug substance</b>
<b>Report ID</b>	<b>INV_054926_HPLC_M4 Version 1.0</b>

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## Distribution

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## 1. Objective

This method 'INV\_054926\_HPLC\_M4' for API TGF-001 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 TGF-001 shall be described in this document.

### Related reports:

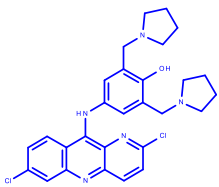
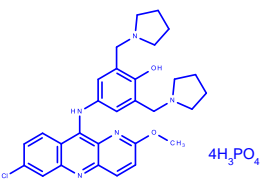
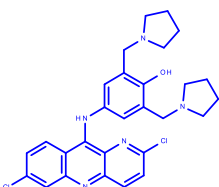
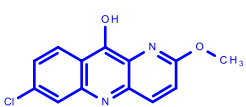
INV\_054926\_HPLC\_M1: HPLC Method Description for In-process Control of Intermediates

INV\_054926\_HPLC\_V4: HPLC Method Limited Validation for TGF-001 (non-GMP)

## 2. Summary and conclusion

The method is applied for the HPLC testing of TGF-001 (identity, assay and related substances). This method is based on reversed phase liquid chromatography with UV detection and gradient elution using a Waters X-Bridge C18, 150\*4.6mm, 3.5µm HPLC column.

**Table 1** Structure, Retention time and RRT of TGF-001 and its related substances

Compound	Structure	Retention time (RT)	RRT	RRF [1]
DIA		ca. 10.4 min	ca. 0.86	1.00
TGF-001		ca. 12.1 min	<b>1.00</b>	-
DIN		ca. 25.3 min	ca. 2.09	1.00
TGF-001 impurity 1		ca. 29.9 min	ca. 2.47	1.00 [2]

[1] RRF value was tested in report 'INV\_054926\_HPLC\_V4'. - [2] Use 1.00, in case there is no RRF determined.

Example chromatograms and extracted HPLC-PDA spectrum of TGF-001 are given in Section 4.

Autosampler stability, Specificity, LOD, LOQ, Linearity, Accuracy and Repeatability were performed and reported in report 'INV\_054926\_HPLC\_V4'.

### 3. Experimental

Equivalent equipment or grade of materials can be used.

#### 3.1. HPLC

- HPLC System: e.g.: Agilent 1260 Infinity II system  
Open Lab CDS-control and integration software or equivalent
- Column: Waters X-Bridge C18, 150\*4.6mm, 3.5µm PN: 186003034
- Flow rate: 1.0 mL/min
- Elution: Gradient mode
- Run time: 40.0 min
- Detection: 278 nm
- Injection: 10 µL
- Column temp.: 35°C ± 2°C
- Auto sampler temp.: Room temperature
- Mobile phase (see section 3.3.2):
  - **A:** 0.1% H<sub>3</sub>PO<sub>4</sub> in Water
  - **B:** Acetonitrile
- Diluent: 0.1% H<sub>3</sub>PO<sub>4</sub> in Water/Acetonitrile (80/20 v/v)
- Needle wash: Water/ Acetonitrile (50:50 v/v)
- Equilibration time: 10.0 min
- Gradient:

**Table 2** Gradient Table

Time (min)	% A	% B
0.0	98	2
3.0	98	2
5.0	92	8
25.0	85	15
32.0	15	85
37.0	15	85
37.1	98	2
40.0	98	2

#### 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
- Phosphoric acid: 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.

0.1% H<sub>3</sub>PO<sub>4</sub> in Water/Acetonitrile (80/20 v/v):

Combine 800 mL of water and 200 mL of acetonitrile into a suitable container, and then add 1 mL of Phosphoric acid. Mix well.

### 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% H<sub>3</sub>PO<sub>4</sub> in Water):

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

#### Mobile phase B (Acetonitrile):

Acetonitrile.

### 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. All TGF-001 Materials must be weighted out in a humidity-controlled environment ( $\leq 10\%$ ).*

#### 3.3.3.1. Standard solutions

##### Standard Solution 1 & 2 (conc.: 0.3 mg/ml):

Accurately weigh approx. 15 mg of TGF-001 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 0.00005 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

Accurately weigh approx. 15 mg of TGF-001 sample into a 50-mL volumetric flask. Dissolve and dilute to volume with diluent. Mix well.

*Number of sample preparations depends on the samples under analysis.*

### 3.4. Proposed injection sequence and system suitability test

**Table 3** Proposed injection sequence and SST criteria

Sample name	No. of injections <sup>[1]</sup>	SST acceptance criteria
Blank (diluent)	1 + N <sup>[1]</sup>	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 <sup>[2]</sup>	1	Recovery: 98% - 102% (6 injection Std 2 to be used as reference)

<sup>[1]</sup> Additional blanks may be run until an acceptable baseline is obtained. - <sup>[2]</sup> 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:  
 $100 - \%(area) \text{ Total impurities}$

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

Where:

$P_i = \text{TGF-001 purity \% (area)}$

#### 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 ≥ 0.05% (area) by their RRT.

Analyses with two sample weighings:  $(A1+A2)/2$

Where:

$A_i = \text{impurity peak \% (area)}$

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

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

Where:

Area <sub>sam</sub> = TGF-001 peak area obtained in the sample chromatogram  
 Area <sub>std</sub> = Average TGF-001 peak area obtained for the 6 standard injections (std 2)  
 W <sub>std</sub> = Standard weight (Std 2)  
 W <sub>Sam</sub> = Sample weight (mg)  
 V <sub>sam</sub> = Volume of the flask used in the sample preparation  
 V <sub>std</sub> = Volume of the flask used in the standard preparation (Std 2)  
 P <sub>std</sub> = Potency determined for the standard in used. Use value (free base) from CoA for TGF-001

$$\text{Assay}_{\text{salt}} (\%w/w) = \text{Assay}_{\text{free base}} \times \frac{MW_{\text{salt}}}{MW_{\text{free base}}}$$

Where:

Assay <sub>free base</sub> = Potency determined for the free base of TGF-001 sample  
 MW <sub>salt</sub> = Molecular weight of TGF-001 salt (910.04 g/mol)  
 MW <sub>free base</sub> = Molecular weight of TGF-001 free base (518.05 g/mol)

### 3.5.6. Related substances by HPLC: Total impurities in %w/w

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

### 3.5.7. Related substances by HPLC: Individual impurities in %w/w

$$\text{Content}_{\text{impurity}} (\%w/w) = \frac{\text{Area}_{\text{imp. sam.}} \times W_{\text{std}} \times V_{\text{sam.}}}{\text{Area}_{\text{TGF-001 std}} \times W_{\text{sam.}} \times V_{\text{std}}} \times \text{RRF}_{\text{imp.}} \times P_{\text{TGF-001 std (free base)}}$$

Where:

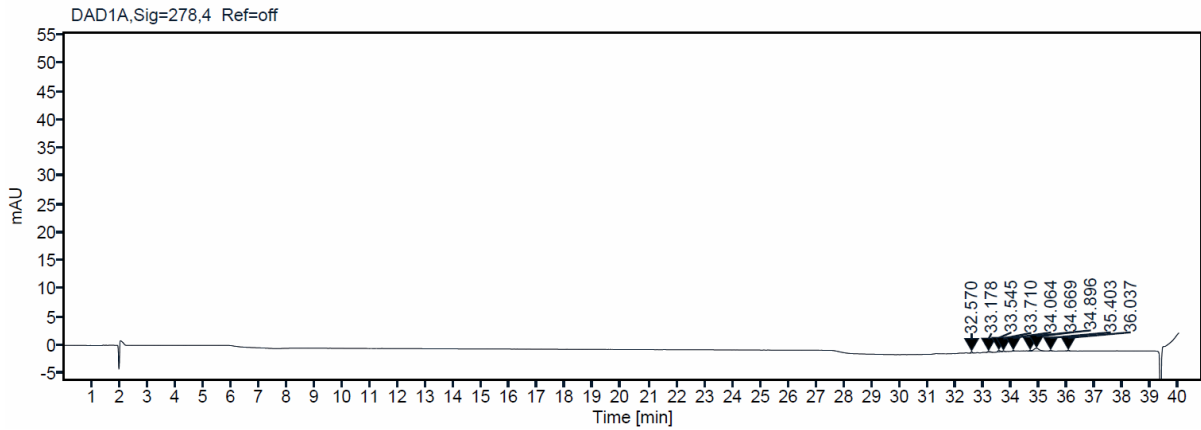
Area <sub>imp. sam.</sub> = Impurity peak area obtained in the sample chromatogram  
 Area <sub>TGF-001 std</sub> = Average TGF-001 peak area obtained for the 6 standard injections (std 2)  
 W <sub>std</sub> = Standard weight (Std 2)  
 W <sub>sam.</sub> = Sample weight (mg)  
 V <sub>sam.</sub> = Volume of the flask used in the sample preparation  
 V <sub>std</sub> = Volume of the flask used in the standard preparation (Std 2)  
 RRF <sub>imp.</sub> = Relative response factor determined for the impurity.  
 (For values see Table 1. Use 1.00, in case there is no RRF determined)  
 P <sub>TGF-001 std (free base)</sub> = Potency determined for the standard in used. Use value from the CoA for TGF-001 (free base).

- For impurity content below LOQ concentration, report "Less than 0.05 %(w/w)" (ICH reportable threshold);

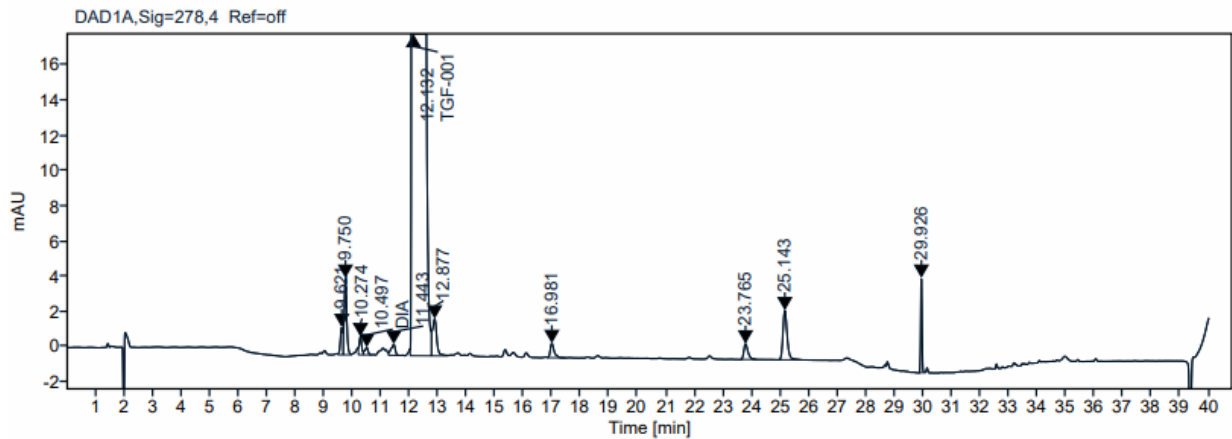
- If impurity is not detected, report "Not detected".

#### 4. Figures

**Figure 1** Example HPLC chromatogram of blank, method INV\_054926\_HPLC\_M4, 278nm

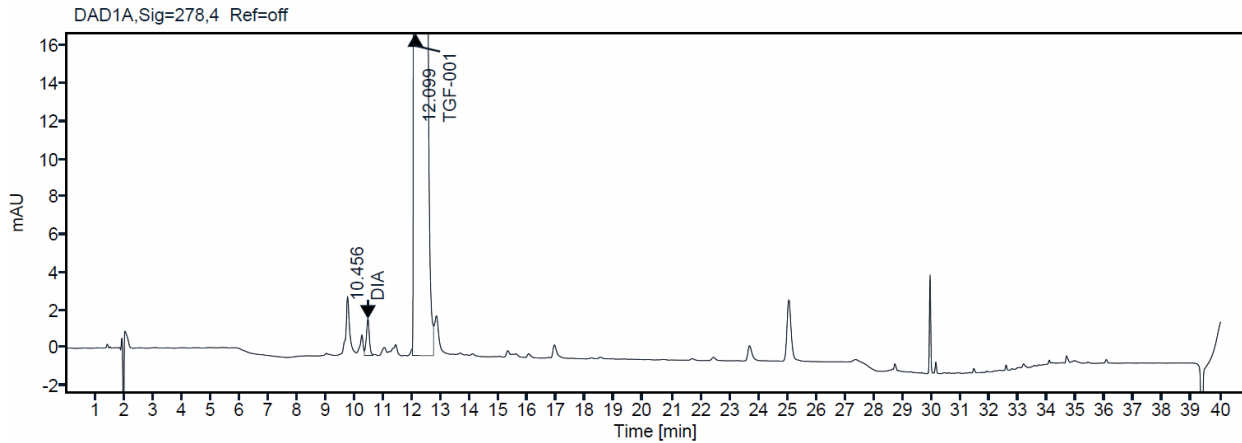


**Figure 2** Example HPLC chromatogram of TGF-001 batch 100144-200502-REF, method INV\_054926\_HPLC\_M4, 278nm

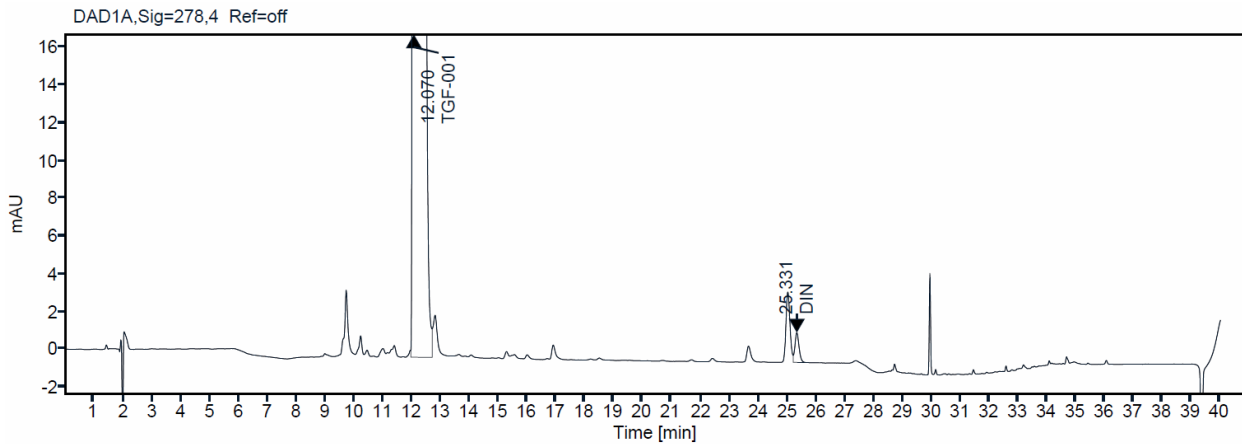


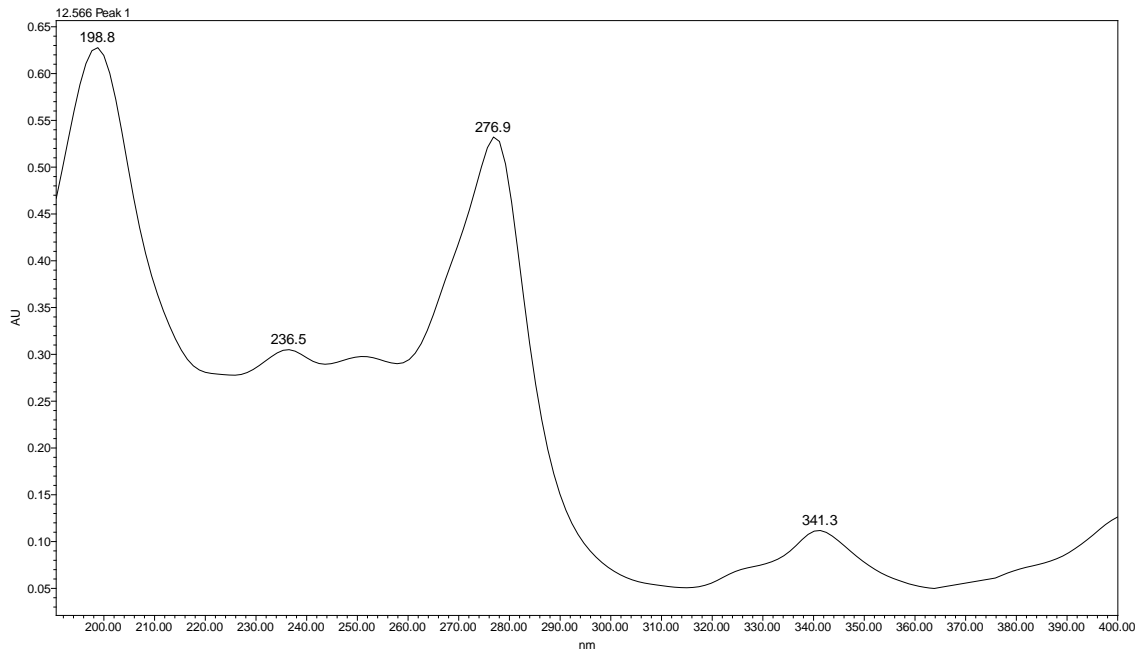
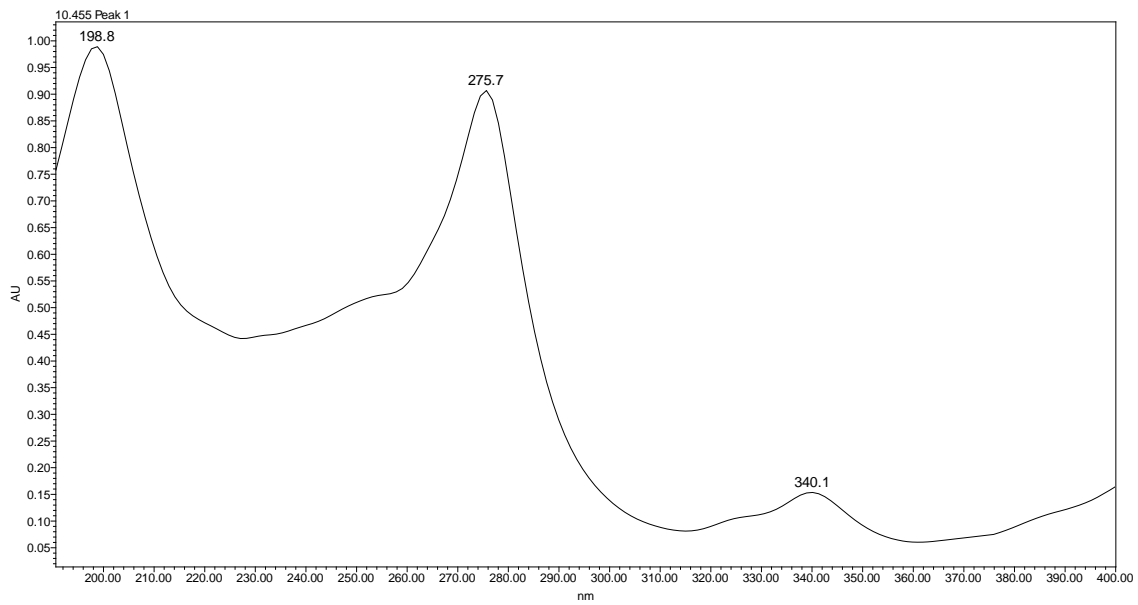


**Figure 3** Example HPLC chromatogram of TGF-001 batch 100144-200502-REF spiked with impurity DIA batch PHTRACKD-394-REF, 278 nm



**Figure 4** Example HPLC chromatogram of TGF-001 batch 100144-200502-REF spiked with impurity DIN batch PHTRACKD-389-REF, 278 nm



**Figure 5** Extracted HPLC-PDA Spectrum of TGF-001, method INV\_054926\_HPLC\_M4**Figure 6** Extracted HPLC-PDA Spectrum of impurity DIA, method INV\_054926\_HPLC\_M4

**Figure 7** Extracted HPLC-PDA Spectrum of impurity DIN, method INV\_054926\_HPLC\_M4

