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**Research article** 

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## **RP-HPLC** method development and validation of Rilpivirine

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## ABSTRACT

A simple, accurate, rapid, and stability-indicating RP-HPLC method for a Rilpivirine has been developed and subsequently validated in commercial tablets. The proposed HPLC method utilizes Develosil ODS HG-5 RP C18,  $5\mu$ m, 15cmx4.6mm and mobile phase consisting of ACN : Acetate buffer (pH=4.0) = 65:35 (v/v) at a flow rate of 1.0 ml/min. Quantitation was achieved with UV detection at 260nm. The method was validated in terms of accuracy, precision, linearity, limits of detection, limits of quantitation, and robustness. This optimized method has been successively applied to pharmaceutical formulation and no interference from the tablet excipients was found. Rilpivirine drug products were subjected to acid, base, neutral hydrolysis, oxidation, dry heat, and photolytic stress conditions and the stressed samples were analyzed by the proposed method. As the proposed RP-HPLC method could effectively separate the drugs from its degradation products, it can be employed as stability-indicating method for the determination of instability of these drugs in bulk and pharmaceutical dosage form.

Keywords: HPLC, Method development, Validation, Reverse Phase, Rilpivirine.

## **INTRODUCTION**

The number of drugs introduced into the market is increasing every year. These drugs may be either new entities or partial structural modification of the existing one [1]. Very often there is a time lag from the date of introduction of a drug into the market to the date of its inclusion in pharmacopoeias. This happens because of the possible uncertainties in the continuous and wider usage of these drugs, reports of new toxicities (resulting in their withdrawal from the market), development of patient resistance and introduction of better drugs by competitors. Under these conditions, standards and analytical procedures for these drugs may not be available in the pharmacopoeias. It becomes necessary, therefore to develop newer analytical methods for such drugs [2-5].

Chromatographic techniques [6-8] are dynamic processes wherein two mutually immiscible phases are brought into contact; one phase is stationary and the other mobile phase. A liquid mobile phase is pumped under pressure through a stainless steel column containing particles of stationary phase with a diameter of  $3-10 \ \mu\text{m}$ . The analyte is loaded onto the head of the column via a loop valve and separation of a mixture occurs according to the relative lengths of time spent by its components in the stationary phase. Components with the least affinity for the stationary phase emerge or elute first whereas the components with greater affinity for stationary phase elute last. Monitoring of the column effluent can be carried out with a variety of detectors. The aim of the proposed method is to develop simple and accurate methods for the determination of Rilpivirine by RP-HPLC method in pharmaceutical dosage forms. This new method was successfully developed and validated as per ICH guidelines [9-10], can be utilized for the validation of Rilpivirine in pharmaceutical dosage forms.

Rilpivirine is non-nucleoside reverse transcriptase inhibitor (NNRTI) which is used for the treatment of HIV-1 infections in treatmentnaive patients. It is a diarylpyrimidine, a class of molecules that resemble pyrimidine nucleotides found in DNA. Because of its flexible chemical structure, resistance of rilpivirine is less likely to develop than other NNRTI's. FDA approved on May 20, 2011.



| IUPAC Name       | 4-{[4-({4-[(1E)-2-cyanoeth-1-en-1-yl]-2,6dimethylphenyl}amino)pyrimidin-2-yl]amino}<br>benzonitrile |
|------------------|---|
| Chemical Formula | C22H18N6  |

## **MATERIALS AND METHODS**

Table 2.1: List of chemicals and equipemnets

| Chemicals                | Equipments           |
|--------------------------|----------------------|
| Ammonium acetate         | Analytical Balance   |
| Dimethyl Sulfoxide(DMSO) | Sonicator            |
| HPLC grade Water         | HPLC                 |
|                          | UV-spectrophotometer |

## Standard & sample preparation for UVspectrophotometer analysis

25 mg of Rilpivirine standard was transferred into 25 ml volumetric flask, dissolved in mobile phase & make up to volume with mobile phase. Further dilution was done by transferring 4 ml of the above solution into a 10ml volumetric flask and make up to volume with mobile phase.

The standard & sample stock solutions were prepared separately by dissolving standard & sample in a solvent in mobile phase diluting with the same solvent.(After optimization of all conditions) for UV analysis. It scanned in the UV spectrum in the range of 200 to 400nm. This has been performed to know the maxima of Rilpivirine, so that the same wave number can be utilized in HPLC UV detector for estimating the Rilpivirine. While scanning the Rilpivirine solution we observed the absorption maxima was 260 nm. The UV spectrum has been recorded on Elico, corp. make UV – Vis spectophotometer model UV-2450. The scanned UV spectrum is attached in



Fig -2.1.1: UV spectrum of Rilpivirine

#### **Mobile Phase Preparation**

The mobile phase used in this analysis consists of a mixture of acetate Buffer (pH adjusted to 4.2 with Glacial acetic acid) and Acetonitrile in a ratio of 40:60.

400 ml of this buffer solution was added and properly mixed with 600 ml of acetronitrile and a homogenous solution is achieved. This mobile phase was filled and sonicated for 15 minutes before using in the experiment.

# Sample & Standard Preparation for the Analysis

25 mg of Rilpivirine standard was transferred into 25 ml volumetric flask, dissolved in DMSO & make up to volume with mobile phase. Further dilution was done by transferring 4 ml of the above solution into a 10ml volumetric flask and make up to volume with mobile phase.

#### **Method Development**

## Trails

| Table no. 2.4.1.1:- Different trails used in method development |  |   |  |  |  |
|---|--|---|--|--|--|
| Mobile Phase  | Flow Wave  |   | Observation  | Result   |  |
|   | Rate   | length  |  |  |  |
| ACN: Water = 60:  | 1.0  | 260   | Did nt get any Peaks   | Method   |  |
| 40  | ml/min   | nm  |  | rejected   |  |
| ACN: water  | 1.0  | 260   | Pseudo peaks   | Method   |  |
| = 70 : 30   | ml/min   | nm  | interfering main peak  | rejected   |  |
| ACN: phosphate  | 1.0 ml/  | 260 nm  | Low response &   | Method   |  |
| buffer = 50 : 50  | min  |   | Broad Peak   | rejected   |  |
| ACN : phosphate   | 1.0  | 260 nm  | Peak broadening  | Method   |  |
| buffer (pH= $2.2$ ) =   | ml/min   |   | C  | rejected   |  |
| 40:60   |  |   |  |  |  |
| ACN : Acetate buffer  | 1.0  | 260 nm  | Nice peak  | Method   |  |
| (pH=4.0) = 65:35  | ml/min   |   |  | accepted   |  |
|   | Mobile Phase<br>ACN : Water = 60 :<br>40<br>ACN: water<br>= 70 : 30<br>ACN: phosphate<br>buffer = 50 : 50<br>ACN : phosphate<br>buffer (pH=2.2) =<br>40:60<br>ACN : Acetate buffer | Mobile Phase         Flow<br>Rate           ACN : Water = $60$ :         1.0 $40$ ml/min           ACN: water         1.0           = 70 : 30         ml/min           ACN: phosphate         1.0 ml/           buffer = $50 : 50$ min           ACN : phosphate         1.0           buffer (pH=2.2) =         ml/min           40:60         1.0 | Mobile PhaseFlow<br>RateWave<br>lengthACN : Water = 60 : $1.0$ $260$ $40$ ml/minnmACN: water $1.0$ $260$ $= 70 : 30$ ml/minnmACN: phosphate $1.0$ $260$ nmbuffer = $50 : 50$ min $260$ nmACN : phosphate $1.0$ $ml/$ ACN : phosphate $1.0$ $260$ nmbuffer (pH=2.2) =ml/min $40:60$ ACN : Acetate buffer $1.0$ $260$ nm | Mobile PhaseFlow<br>RateWave<br>lengthObservationACN : Water = 60 :1.0260Did nt get any Peaks $40$ ml/minnmDid nt get any Peaks $40$ ml/minnmACN: water $= 70 : 30$ 1.0260Pseudo peaks<br>interfering main peakACN: phosphate1.0ml/minbuffer = 50 : 501.0260 nmLow response &<br>Broad PeakACN : phosphate1.0260 nmPeak broadeningbuffer (pH=2.2) =ml/min260 nmPeak broadeningACN : Acetate buffer1.0260 nmNice peak |  |

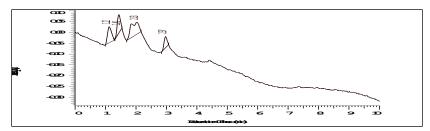


Fig -2.4.1.1:- Chromatogram of Trial-1

| Table 2.4.1.2 Results of Trial-1 |      |                           |      |                |
|----------------------------------|------|---------------------------|------|----------------|
| Sl. No                           | Rt   | <b>Theoretical Plates</b> | Area | Tailing factor |
| 1                                | 1.12 | 2153                      | 2125 | 1.91           |
| 2                                | 1.44 | 2586                      | 2823 | 1.97           |
| 3                                | 2.03 | 2453                      | 2456 | 1.92           |
| 4                                | 2.97 | 2354                      | 1956 | 1.94           |

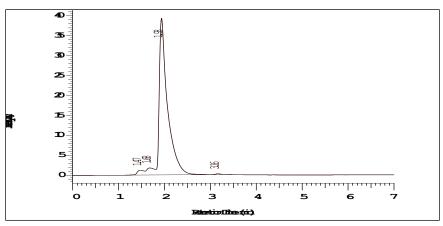


Fig - 2.4.1.2:- Chromatogram of Trial-2

| Table 2.4.3.: Results of Trial-2 |      |                           |      |                |  |
|----------------------------------|------|---------------------------|------|----------------|--|
| Sl. No                           | Rt   | <b>Theoretical Plates</b> | Area | Tailing factor |  |
| 1                                | 1.47 | 2365                      | 2546 | 1.94           |  |
| 2                                | 1.68 | 2658                      | 2752 | 1.68           |  |
| 3                                | 1.93 | 4745                      | 3862 | 0.89           |  |

1358

1.76

4

3.16

2214

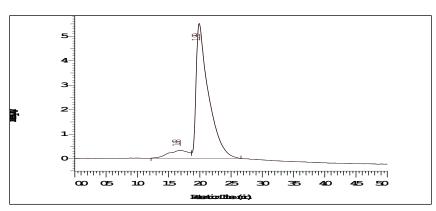


Fig- 2.4.1.3:- Chromatogram of Trial-3

| Table 2.4.1.4: Results of Trial-3 |      |                           |      |                |
|-----------------------------------|------|---------------------------|------|----------------|
| Sl. No                            | Rt   | <b>Theoretical Plates</b> | Area | Tailing factor |
| 1                                 | 1.68 | 2568                      | 3254 | 1.91           |
| 2                                 | 1.99 | 2764                      | 6587 | 0.95           |

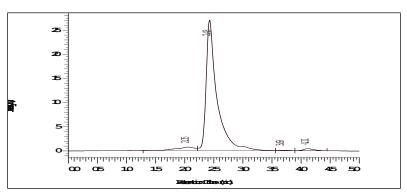


Fig- 2.4.1.4:- Chromatogram of Trial-4 Table 2.4.1.5:Results of Trial-4

| Sl. No | Rt   | <b>Theoretical Plates</b> | Area | Tailing factor |
|--------|------|---------------------------|------|----------------|
| 1      | 1.47 | 2365                      | 2365 | 1.84           |
| 2      | 1.68 | 6681                      | 5462 | 0.74           |
| 3      | 1.93 | 2847                      | 2684 | 1.28           |
| 4      | 3.16 | 2354                      | 2364 | 1.59           |

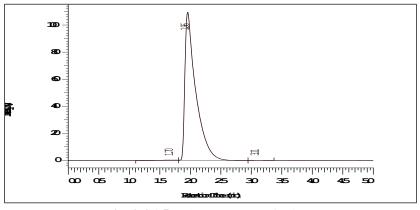


Fig- 2.4.1.5:- Chromatogram of Trial-5

| Ta    | Table 2.4.1.6:Results of Trial-5 |      |         |  |
|-------|----------------------------------|------|---------|--|
| S. no | Name                             | RT   | Area    |  |
| 1     | Rilpivirine                      | 1.96 | 1678995 |  |

| <b>Optimized Chromatographic Conditions</b> |                         | Injection volume   | : 20µl                             |  |
|---|-------------------------|--------------------|------------------------------------|--|
| Column                                      | : Develosil ODS HG-5 RP | Run time           | : 05minutes<br>: 25 <sup>0</sup> C |  |
| 150mm x 4.6mm 5µm particle size             |                         | Column temperature |                                    |  |
| Flow Rate                                   | : 1.0ml/minute          | Sampler cooler     | :Ambient                           |  |
| Wave length                                 | : 260nm                 |                    |                                    |  |

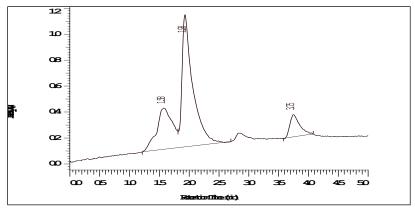


Fig -2.4.2.1:- Chromatogram for blank

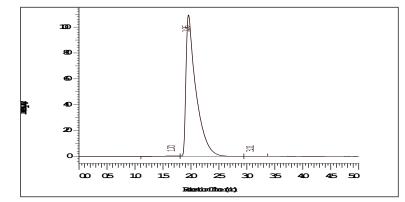


Fig-2.4.2.2: HPLC spectrum of Rilpivirine (40 ppm) in optimized conditions (RT 1.96 min.)

| Table no 2.4.2.1: Peak results of optimised chromatogram |
|--|
|--|

| S. no | Name        | RT   | Area    |
|-------|-------------|------|---------|
| 1     | Rilpivirine | 1.96 | 1678995 |

#### **Evaluation of system suitability**

Perform the blank run by injecting  $5\mu$ l of diluent and ensure that there is no interference with the main peak retention time. Inject  $5\mu$ l of the standard solution for two times into the chromatograph and measure the peak responses. Calculate the % RSD for replicate injections of the standard solution and it should be less than 2.0%.

#### Procedure

 $5\mu$ l of placebo solution is injected and ensured that there is no interference. Then  $5\mu$ l of sample solution is injected into the chromatograph and recorded the chromatograms. Now peak area responses are measured and the average peak responses are taken.

## Calculations

Content of the drug = At\As x Sc\Tc x LC\100 x Potency Where,

At = Avg area responses of Rilpivirine obtained from the Sample preparation As = Avg area responses of Rilpivirine obtained from the Standard preparation Sc = Working standard concentration Tc = Test sample concentration LC = Label claim

## RESULTS

#### **Forced Degradation Studies**

Following protocol was strictly adhered to for forced degradation of Rilpivirine Active Pharmaceutical Ingredient (API). The API (Rilpivirine) was subjected to stress conditions in various ways to observe the rate and extent of degradation that is likely to occur in the course of storage and/or after administration to body. This is one type of accelerated stability studies that helps us determining the fate of the drug that is likely to happen after along time storage, within a very short time as compare to the real time or long term stability testing.

The various degradation pathways studied are acid hydrolysis, basic hydrolysis, thermal degradation and oxidative degradation.

#### Acid hydrolysis

An accurately weighed 25 mg. of pure drug was transferred to a clean & dry 25 ml volumetric flask. To which 0.1 N Hydrochloric acid was added & make up to the mark & kept for 24 hrs. from that 0.1 ml was taken in to a 10 ml volumetric flask & make up to the mark with mobile phase, then injected into the HPLC system against a blank of HCl (after all optimized conditions).

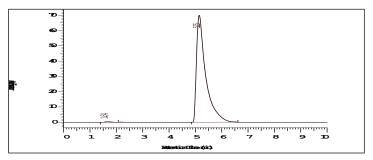


Fig -3.2.1: Chromatogram showing degradation for Rilpivirine in 0.1 N HCl

Table-3.2.1: Results of acid hydrolysis of Rilpivirine in 0.1 N HCL

| Sl. No | Rt   | Theoretical Plates | Area | Tailing factor |
|--------|------|--------------------|------|----------------|
| 1      | 1.64 | 1745               | 2548 | 1.59           |
| 2      | 5.15 | 2934               | 5687 | 0.17           |

#### **Basic Hydrolysis**

An accurately weighed 10 mg. of pure drug was transferred to a clean & dry 10 ml volumetric flask. To which 0.1 N Sodium hydroxide was added & make up to the mark & kept for 24 hrs. from that 4s ml was taken in to a 10 ml volumetric flask & make up to the mark with mobile phase, then injected into the HPLC system against a blank of NaOH (after all optimized conditions).

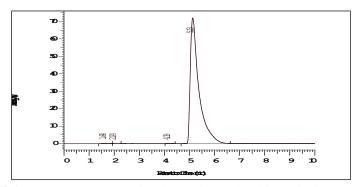


Fig -3.3.1: Chromatogram showing degradation related impurity in 0.1 N NaOH

Table-3.3.2: Results of basic hydrolysis of Rilpivirine in 0.1 N NaOH

| Sl. No | Rt   | <b>Theoretical Plates</b> | Area | Tailing factor |
|--------|------|---------------------------|------|----------------|
| 1      | 1.64 | 1685                      | 2365 | 1.68           |
| 2      | 2.04 | 2156                      | 2658 | 1.29           |
| 3      | 4.19 | 2654                      | 2864 | 1.75           |
| 4      | 5.15 | 5698                      | 4985 | 0.59           |

#### **Thermal Degradation**

An accurately weighed 10 mg. of pure drug was transferred to a clean & dry 100 ml volumetric flask, make up to the mark with mobile phase & was maintained at 50 0C. for 24 hrs then injected into the HPLC system against a blank of mobile phase (after all optimized conditions).

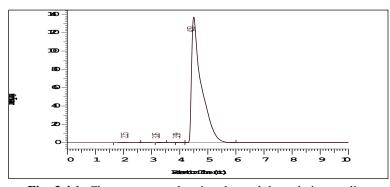


Fig -3.4.1: Chromatogram showing thermal degradation studies

| Table-3.4.2: Results of thermal degradation of Rilpivirine |      |                    |      |                |  |  |
|--|------|--------------------|------|----------------|--|--|
| Sl. No   | Rt   | Theoretical Plates | Area | Tailing factor |  |  |
| 1  | 2.15 | 2416               | 2365 | 1.37           |  |  |
| 2  | 3.26 | 2568               | 2484 | 1.28           |  |  |
| 3  | 3.99 | 2654               | 2657 | 1.49           |  |  |
| 4  | 4.50 | 6598               | 6254 | 0.73           |  |  |

#### **Photolytic degradation**

Approximately 10 mg. of pure drug was taken in a clean & dry Petridis. It was kept in a UV cabinet at 254 nm wavelength for 24 hours without interruption. Accurately weighed 1 mg. of the UV exposed drug was transferred to a clean & dry 10 ml. volumetric flask. First the UV exposed drug was dissolved in methanol & make up to the mark than injected into the HPLC system against a blank of mobile phase (after all optimized conditions).

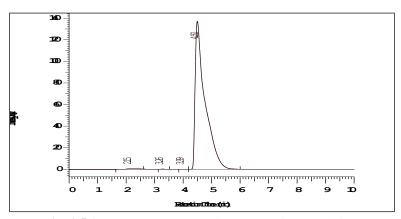


Fig -3.5.1: Chromatogram showing photolytic degradation.

| Table-3.5.1: Results of pho | tolytic degrad | ation of Rilpivirine |
|-----------------------------|----------------|----------------------|
|-----------------------------|----------------|----------------------|

| Sl. No | Rt   | Theoretical Plates | Area | Tailing factor |
|--------|------|--------------------|------|----------------|
| 1      | 2.15 | 2145               | 2365 | 1.38           |
| 2      | 3.26 | 2365               | 2654 | 1.49           |
| 3      | 3.99 | 2564               | 2846 | 1.53           |
| 4      | 4.50 | 6954               | 5942 | 0.81           |

## Oxidation with (3%) h<sub>2</sub>o<sub>2</sub>

Accurately weighed 10 mg. of pure drug was taken in a clean & dry 100 ml. volumetric flask. 30 ml. of 3% H2O2 and a little methanol was added to

it to make it soluble & then kept as such in dark for 24 hours. Final volume was made up to 100 ml. using water to prepare 100 ppm solution. The above sample was injected into the HPLC system.

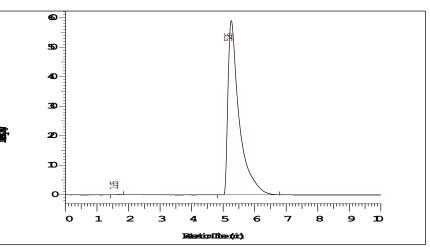


Fig -3.6.1: Chromatogram showing oxidative degradation.

Table-3.6.1: Results of oxidative degradation of Rilpivirine

| Sl. No | Rt   | Theoretical Plates | Area | Tailing factor |
|--------|------|--------------------|------|----------------|
| 1      | 2.15 | 2365               | 2564 | 1.85           |
| 2      | 3.26 | 5687               | 4685 | 0.73           |

#### **Results of degradation studies**

The results of the stress studies indicated the specificity of the method that has been developed.

Rilpivirine was stable in photolytic & temperature stress conditions. The result of forced degradation studies are given in the following table-15.

| Table-3.7.1: F | Results of forc | e degradation | studies | of Rilpivi | rine API. |
|----------------|-----------------|---------------|---------|------------|-----------|
|                |                 |               |         |            |           |

| Stress condition                        | Time   | Assay of active | Assay of degraded | Mass Balance |
|---|--------|-----------------|-------------------|--------------|
|   |        | substance       | products          | (%)          |
| Acid Hydrolysis (0.1 M                  | 24Hrs. | 40.73           | 59.27             | 100          |
| HCl)                                    |        |                 |                   |              |
| Basic Hydrolysis (0.I M                 | 24Hrs. | 80.93           | 19.07             | 100          |
| NaOH)                                   |        |                 |                   |              |
| Thermal Degradation (50 <sup>0</sup> C) | 24Hrs. | 99.35           |                   | 99.35        |
| UV (254nm)                              | 24Hrs. | 98.31           |                   | 99.31        |
| 3 % Hydrogen peroxide                   | 24Hrs. | 91.37           | 08.46             | 99.83        |

## **METHOD VALIDATION**

#### Accuracy: Recovery study

To determine the accuracy of the proposed method, recovery studies were carried out by adding different amounts (80%, 100%, and 120%) of pure drug of RILPIVIRINE were taken and added to the pre-analysed formulation of concentration  $10\mu g/ml$ . From that percentage recovery values were calculated. The results were shown in table-16.

| STD   |         |   | Spike-1 |         |         |            |
|-------|---------|---|---------|---------|---------|------------|
| Conc. | AUC     |   | Conc    | AUC     | Diff    | % Recovery |
| 8     | 506881  | 1 | 18      | 1905481 | 1398600 | 99.14396   |
| 8     | 506881  | 2 | 18      | 1900430 | 1393549 | 98.7859    |
| 8     | 506881  | 3 | 18      | 1902331 | 1395450 | 98.92066   |
|       |         |   |         |         |         | 98.95017   |
|       |         |   |         |         |         | 0.180843   |
|       |         |   |         |         |         | 0.182762   |
| STD   |         |   | Spike-2 |         |         |            |
| Conc. | AUC     |   | Conc    | AUC     | Diff    | % Recover  |
| 10    | 1426346 | 1 | 20      | 2832999 | 1406653 | 99.71482   |
| 10    | 1426346 | 2 | 20      | 2834395 | 1408049 | 99.81378   |
| 10    | 1426346 | 3 | 20      | 2833499 | 1407153 | 99.75026   |
|       |         |   |         |         |         | 99.75962   |
|       |         |   |         |         |         | 0.050139   |
|       |         |   |         |         |         | 0.05026    |
| STD   |         |   | Spike-3 |         |         |            |
| Conc. | AUC     |   | Conc    | AUC     | Diff    | % Recover  |
| 12    | 1999858 | 1 | 22      | 3401595 | 1401737 | 99.36633   |
| 12    | 1999858 | 2 | 22      | 3400499 | 1400641 | 99.28864   |
| 12    | 1999858 | 3 | 22      | 3403358 | 1403500 | 99.49131   |
|       |         |   |         |         |         | 99.38209   |
|       |         |   |         |         |         | 0.10225    |
|       |         |   |         |         |         | 0.102885   |

Table-3.8.1: Accuracy Readings

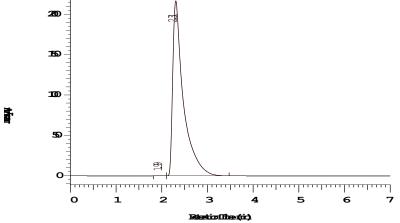


Fig -3.8.1:-Chromatogram for accuracy-1 Spike-1

 Table-3.8.2: Accuracy Readings of accuracy-1 Spike-1

|   |      | Theoretical Plates |      | <i>y</i> 1 |
|---|------|--------------------|------|------------|
| 1 | 1.99 | 2156               | 2654 | 1.59       |
| 2 | 2.31 | 4512               | 3984 | 0.73       |

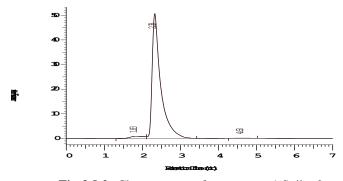
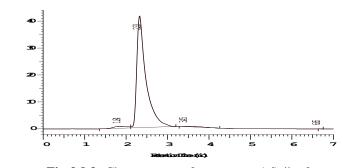


Fig-3.8.2:-Chromatogram for accuracy-1 Spike-2

Table-3.8.3: Accuracy Readings of accuracy-1 Spike-2

| Sl. No | Rt   | <b>Theoretical Plates</b> | Area | Tailing factor |
|--------|------|---------------------------|------|----------------|
| 1      | 1.83 | 2641                      | 2365 | 1.65           |
| 2      | 2.33 | 6954                      | 5654 | 1.97           |
| 3      | 4.63 | 2021                      | 2345 | 0.36           |



Ŧ

Fig-3.8.3:-Chromatogram for accuracy-1 Spike-3

| Table-3.6.4. Accuracy Readings of accuracy-1 Spike-5 |      |                    |      |                |  |  |
|--|------|--------------------|------|----------------|--|--|
| Sl. No   | Rt   | Theoretical Plates | Area | Tailing factor |  |  |
| 1  | 1.84 | 2364               | 2325 | 1.28           |  |  |
| 2  | 2.33 | 5461               | 4564 | 0.36           |  |  |
| 3  | 3.45 | 2021               | 2684 | 1.59           |  |  |
| 4  | 6.65 | 2465               | 2643 | 1.49           |  |  |

Table-3.8.4: Accuracy Readings of accuracy-1 Spike-3

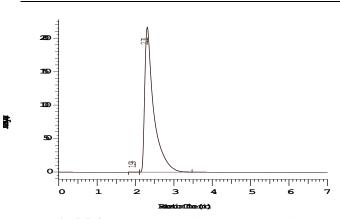


Fig-3.8.4:-Chromatogram for accuracy-2 Spike-1

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Table-3.8.5: Accuracy Readings of accuracy-2 Spike-1

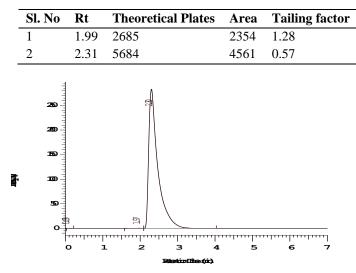


Fig-3.8.5:-Chromatogram for accuracy-2 Spike-2

**Table-3.8.6:** Accuracy Readings of accuracy-2 Spike-2

| Sl. No | Rt   | <b>Theoretical Plates</b> | Area | Tailing factor |
|--------|------|---------------------------|------|----------------|
| 1      | 0.09 | 2657                      | 2365 | 1.28           |
| 2      | 1.97 | 2634                      | 2546 | 1.37           |
| 3      | 2.30 | 5478                      | 4857 | 0.76           |
|        |      |                           |      |                |

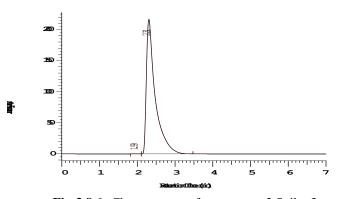


Fig-3.8.6:-Chromatogram for accuracy-2 Spike-3

| Table-3.8.7: Accuracy Readings of accuracy-2 Spik |
|---|
|---|

| Sl. No | Rt   | <b>Theoretical Plates</b> | Aı   | rea | Tailing factor |
|--------|------|---------------------------|------|-----|----------------|
| 1      | 1.99 | 2654                      | 2364 |     | 1.59           |
| 2      | 2.31 | 6587                      | 5698 |     | 0.76           |

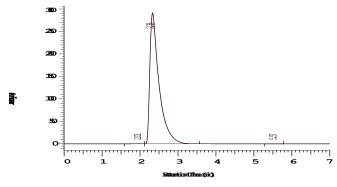


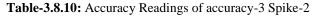
Fig-3.8.7:-Chromatogram for accuracy-3 Spike-1

Table-3.8.8: Accuracy Readings of accuracy-3 Spike-1

| 1 401                      | Tuble-5.6.6. Recuracy Readings of accuracy 5 Spike 1 |                           |      |                |  |
|----------------------------|--|---------------------------|------|----------------|--|
| Sl. No                     | Rt   | <b>Theoretical Plates</b> | Area | Tailing factor |  |
| 1                          | 2.01   | 2354                      | 2546 | 1.59           |  |
| 2                          | 2.33   | 6356                      | 5762 | 0.71           |  |
| 3                          | 5.57   | 2021                      | 2014 | 1.43           |  |
| 30<br>30<br>20<br>10<br>10 | ուսերերերերեր  | 臣君                        |      |                |  |

۳G) С  $\begin{bmatrix} 0 & 1 & 2 & 3 & 4 & 5 & 6 & 7 \end{bmatrix}$ 

Fig-3.8.9:-Chromatogram for accuracy-3 Spike-2



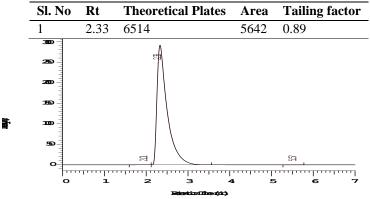


Fig-3.8.10:-Chromatogram for accuracy-3 Spike-3

| Sl. No | Rt   | <b>Theoretical Plates</b> | Area | Tailing factor |
|--------|------|---------------------------|------|----------------|
| 1      | 2.01 | 2461                      | 2645 | 1.27           |
| 2      | 2.33 | 7546                      | 6421 | 1.49           |
| 3      | 5.57 | 3164                      | 3024 | 0.53           |

## Precision

## **Repeatability**

The precision of each method was ascertained separately from the peak areas & retention times

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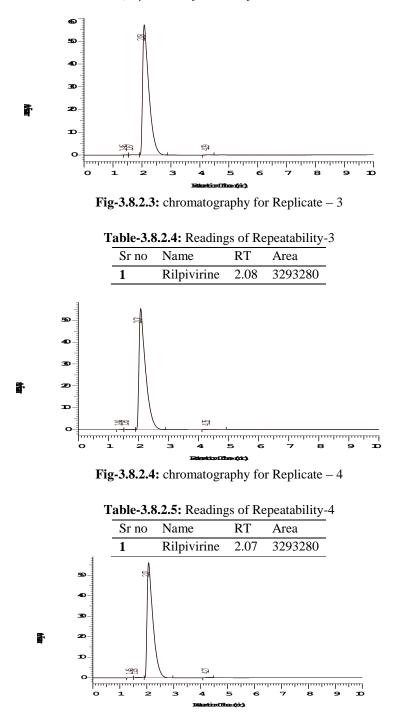
obtained by actual determination of five replicates of a fixed amount of drug. Rilpivirine (API). The percent relative standard deviations were calculated for Rilpivirine are presented in the table-26.

| HPLC         | Injection                       | n R                                    | Retentior                          | 1 Time             | Area     |
|--------------|---------------------------------|--|------------------------------------|--------------------|----------|
| Replica      | tes of Ri                       | lpivirine                              |                                    |                    |          |
| Replica      | te – 1                          | 2                                      | .08                                |                    | 833769   |
| Replicat     | te – 2                          | 2                                      | .08                                |                    | 835768   |
| Replicat     | te – 3                          | 2                                      | .08                                |                    | 855929   |
| Replicat     | te – 4                          | 2                                      | .07                                |                    | 833458   |
| Replicat     | te – 5                          | 2                                      | .07                                |                    | 848232   |
| Average      | e                               | 2                                      | .076                               |                    | 841431.2 |
| Standar      | d Deviati                       | ion 0                                  | .0054772                           | 226                | 10133.98 |
| % RSD        |                                 | 0                                      | .2638355                           | 529                | 1.204374 |
|              |                                 | <u></u>                                |                                    |                    |          |
| •<br>•<br>Fi | -                               | : chromatogr<br>3.2.2: Reading         | gs of Rep                          | -                  |          |
| •<br>•<br>Fi | g-3.8.2.1                       | : chromatogr                           | aphy for                           | -                  |          |
| •<br>•<br>Fi | g-3.8.2.1<br>Fable-3.8          | : chromatogr<br>3.2.2: Reading         | aphy for<br>gs of Rep              | peatabilit         | ty-1     |
| •<br>•<br>Fi | g-3.8.2.1<br>Fable-3.8<br>Sr no | : chromatogr<br>3.2.2: Reading<br>Name | aphy for<br>gs of Rep<br><b>RT</b> | peatabilit<br>Area | ty-1     |

**Fig-3.8.2.2:** chromatography for Replicate – 2

| Table-3.8.2.3: | Readings ( | of Repeata | bility-2 |
|----------------|------------|------------|----------|
|----------------|------------|------------|----------|

| Sr no | Name        | RT   | Area    |
|-------|-------------|------|---------|
| 1     | Rilpivirine | 2.08 | 3293280 |







| Sr no | Name        | RT   | Area    |
|-------|-------------|------|---------|
| 1     | Rilpivirine | 2.07 | 3293280 |

#### Intra-assay & inter-assay

The intra & inter day variation of the method was carried out & the high values of mean assay & low values of standard deviation & % RSD (% RSD < 2%) within a day & day to day variations for Rilpivirine revealed that the proposed method is precise.

| Conc. Of Rilpivirine (API) (µg/ml) | <b>Observed Conc.</b> | Of Rilpivirine | e (µg/ml) by the pr | oposed method |  |
|------------------------------------|-----------------------|----------------|---------------------|---------------|--|
|                                    | Intra-Day Inter-Day   |                | Intra-Day Inter-Day |               |  |
|                                    | Mean (n=6)            | % RSD          | Mean (n=6)          | % RSD         |  |
| 10                                 | 9.93                  | 0.14           | 10.32               | 0.95          |  |
| 30                                 | 30.63                 | 0.78           | 30.14               | 0.16          |  |
| 100                                | 99.21                 | 0.96           | 99.78               | 0.73          |  |

Table-3.8.2.7: Results of intra-assay & inter-assay

## Linearity & Range

The calibration curve showed good linearity in the range of 0-25 µg/ml, for Rilpivirine (API) with correlation coefficient (r<sup>2</sup>) of 0.994 (Fig-39). A typical calibration curve has the regression equation of y = 6780x + 65596 for Rilpivirine.

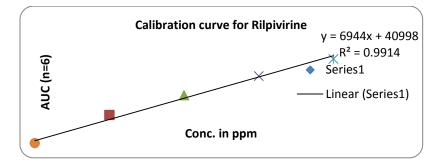


Fig-3.8.3.1: Calibration curve of Rilpivirine (API).

| Table-3.8.3.1: Linearity Results |           |  |  |
|----------------------------------|-----------|--|--|
| CONC. in ppm                     | AUC (n=6) |  |  |
| 0                                | 0         |  |  |
| 50                               | 460548    |  |  |
| 100                              | 783769    |  |  |
| 150                              | 1096795   |  |  |
| 200                              | 1376884   |  |  |

| Fig-3.8.3.2: | Chromatogram | for | 50ppm |
|--------------|--------------|-----|-------|
|--------------|--------------|-----|-------|

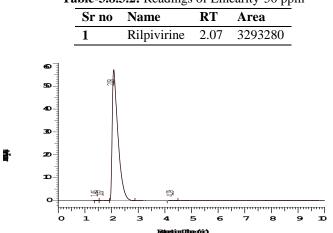


Table-3.8.3.2: Readings of Linearity-50 ppm

Fig-3.8.3.3: Chromatogram for 100ppm

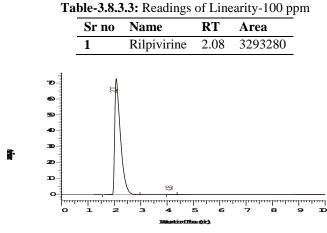


Fig-3.8.3.4: Chromatogram for 150ppm

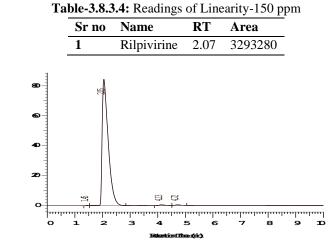


Fig-3.8.3.5: Chromatogram for 200ppm

|  | Table-3. | 8.3.5: | Readings | of Lin | earity-20 | ) ppm |
|--|----------|--------|----------|--------|-----------|-------|
|--|----------|--------|----------|--------|-----------|-------|

| Sr no | Name        | RT   | Area    |
|-------|-------------|------|---------|
| 1     | Rilpivirine | 2.05 | 3293280 |

## **Method Robustness**

Influence of small changes in chromatographic conditions such as change in flow rate ( $\pm$  0.1ml/min), Temperature ( $\pm 2^{0}$ C), Wavelength of detection ( $\pm 2$ nm) & acetonitrile content in mobile

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phase ( $\pm 2\%$ ) studied to determine the robustness of the method are also in favour of (Table-38, % RSD < 2%) the developed RP-HPLC method for the analysis of Rilpivirine(API).

| Table-5.8.4.1: Result of method root | istness test |
|--------------------------------------|--------------|
| Change in parameter                  | % RSD        |
| Flow (1.1 ml/min)                    | 0.06         |
| Flow (0.9 ml/min)                    | 0.04         |
| Temperature $(27^{0}C)$              | 0.08         |
| Temperature $(23^{\circ}C)$          | 0.11         |
| Wavelength of Detection (202 nm)     | 0.03         |
| Wavelength of detection (209 nm)     | 0.02         |

## LOD & LOQ

The Minimum concentration level at which the analyte can be reliable detected (LOD) & quantified (LOQ) were found to be 0.03 & 0.09  $\mu$ g/ml respectively.

#### **System Suitability Parameter**

System suitability testing is an integral part of many analytical procedures. The tests are based on

the concept that the equipment, electronics, analytical operations and samples to be analyzed constitute an integral system that can be evaluated as such. Following system suitability test parameters were established. The data are shown in Table-39.

| Table-3.8.6.1: | Data of System | Suitability Parameter |
|----------------|----------------|-----------------------|
|----------------|----------------|-----------------------|

| S.No. | Parameter         | Limit      | Result           |
|-------|-------------------|------------|------------------|
| 1     | Resolution        | Rs > 2     | 9.15             |
| 2     | Asymmetry         | $T \leq 2$ | Rilpivirine=0.12 |
| 3     | Theoretical plate | N > 2000   | Rilpivirine=3246 |

#### SPECIFICITY

## **Preparation and running of Rilpivirine**

The performance test of the method has been conducted on market sample. As per the label claim, each tablet contains 50mg of Rilpivirine. To estimate this powder of the tablet equivalent to 25mg of Rilpivirine has been dissolved in 25 ml of the mobile phase. Further dilution was done by taking 1ml of this solution in 10ml volumetric flask, dissolve and making up the volume upto the mark with mobile phase by which 100ppm solution was prepared. Again same process is repeated to make 10ppm from 100ppm solution. To extract the drug in the solution, it has been sonicated for 5 minutes followed by cyclo-mixing for 5 minutes. Resulting solution was filtered by using Millipore syringe filter (0.45 micron). Resulting clear solution was injected in HPLC in duplicate as per the above mentioned HPLC method. Chromatogram obtained for the injection is shown below with Rt of 2.69 mins.

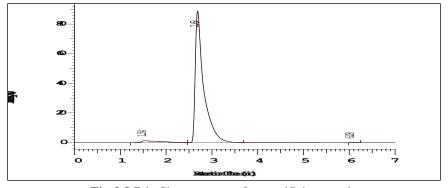


Fig-3.8.7.1: Chromatogram for specificity sample

Table-3.8.7.1: Results for specificity sample

| Sl. No | Rt   | <b>Theoretical Plates</b> | Area | Tailing factor |
|--------|------|---------------------------|------|----------------|
| 1      | 1.54 | 2143                      | 2354 | 1.28           |
| 2      | 2.69 | 5658                      | 4689 | 1.74           |
| 3      | 6.08 | 2654                      | 2541 | 0.49           |

## Assay of Rilpivirine in dosage form

## Estimation of rilpivirine in tablet dosage form rilpivirine 25 mg

Twenty tablets were taken and the I.P. method was followed to determine the average weight. Above weighed tablets were finally powdered and triturated well. A quantity of powder equivalent to 100 mg of drugs were transferred to 100 ml volumetric flask, and 70 ml of Hplc grade methanol was added and solution was sonicated for 15 minutes, there after volume was made up to 100 ml with same solvent. Then 10 ml of the above

## ASSAY

Where:

AT = Peak Area of Test obtained with test preparation AS = Peak Area of Standard obtained with standard preparation WS = Weight of working standard taken in mg WT = Weight of sample taken in mg solution was diluted to 100 ml with hplc grade methanol. The solution was filtered through a membrane filter (0.45  $\Box$  m) and sonicated to degas. From this stock solution (3.5 ml) was transferred to five different 10 ml volumetric flasks and volume was made up to 10 ml with same solvent system.

The solution prepared was injected in five replicates into the HPLC system and the observations were recorded.

A duplicate injection of the standard solution was also injected into the HPLC system and the peak areas were recorded. The data are shown in Table-41.

DS = Dilution of Standard solution DT = Dilution of sample solution P = Percentage purity of working standard Assay was performed as described in previous chapter. Results obtained are tabulated below:

| Brand name of  | Labelled amount of Drug | Mean (±SD) amount (mg) found by | Mean (± SD)     |
|----------------|-------------------------|---------------------------------|-----------------|
| tablets        | ( <b>mg</b> )           | the proposed method (n=6)       | Assay $(n = 6)$ |
| EDURANT        | 25                      | 25.31 (±0.06)                   | 100.07 (±0.48)  |
| (Janssen Inc.) |                         |                                 |                 |

The assay of EDURANT tablets containing Rilpivirine was found to be 100.07 %.

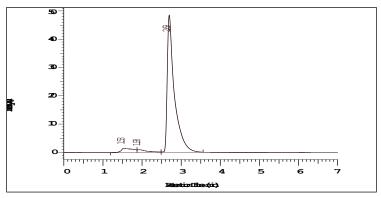
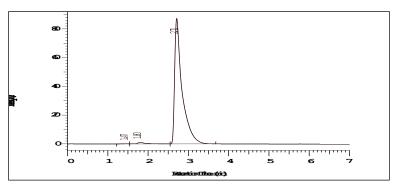


Fig-3.8.8.1: Chromatogram for assay sample-1

Table-3.8.8.1: Assay chromatogram results of Rilpivirine sample-1

| Sl. No | Rt   | <b>Theoretical Plates</b> | Area | Tailing factor |
|--------|------|---------------------------|------|----------------|
| 1      | 1.53 | 2153                      | 2133 | 1.91           |
| 2      | 1.93 | 2586                      | 2852 | 1.97           |
| 3      | 2.69 | 5642                      | 5293 | 0.98           |



**Fig-3.8.8.2:** Chromatogram for assay sample-2 **Table-3.8.8.2:** Assay chromatogram results of Rilpivirine sample-2

| Sl. No | Rt   | <b>Theoretical Plates</b> | Area | Tailing factor |
|--------|------|---------------------------|------|----------------|
| 1      | 1.47 | 2654                      | 2635 | 1.57           |
| 2      | 1.80 | 2745                      | 2843 | 1.29           |
| 3      | 2.71 | 6846                      | 5684 | 0.37           |

## **DISCUSSION AND CONCLUSION**

To develop a precise, linear, specific & suitable stability indicating RP-HPLC method for analysis chromatographic of Rilpivirine, different conditions were applied & the results observed are presented in previous chapters. Isocratic elution is simple, requires only one pump & flat baseline separation for easy and reproducible results. So, it was preferred for the current study over gradient elution. In case of RP-HPLC various columns are available, but here develosil C<sub>18</sub>, 5µm, 150 x 4.6 mm i.d. column was preferred because using this column peak shape, resolution and absorbance were good. Mobile phase & diluents for preparation of various samples were finalized after studying the solubility of API in different solvents of our disposal (methanol, acetonitrile, dichloromethane, water, 0.1N NaOH, 0.1NHCl).

The drug was found to be highly soluble in acetonitrile & soluble in dichloromethane and methanol. Drug was soluble in water. Using these

solvents with appropriate composition newer methods can be developed and validated. Detection wavelength was selected after scanning the standard solution of drug over 200 to 400nm. From the U.V spectrum of Rilpivirine it is evident that most of the HPLC work can be accomplished in the wavelength range of 210-2500 nm conveniently. Further, a flow rate of 1 ml/min & an injection volume of 20  $\mu$ l were found to be the best analysis.

The result shows the developed method is yet another suitable method for assay and purity which can help in the analysis of Rilpivirine in different formulations. A sensitive & selective RP-HPLC method has been developed & validated for the analysis of Rilpivirine API.

Further the proposed RP-HPLC method has excellent sensitivity, precision and reproducibility. The result shows the developed method is yet another suitable method for assay, purity which can help in the analysis of Rilpivirine in different formulations.

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