

# RP-HPLC METHOD DEVELOPMENT AND VALIDATION FOR THE DETECTION OF IMPURITIES IN BREXANOLONE PHARMACEUTICAL DOSAGE FORM

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## **Abstract**

*A simple, Accurate, Precise method was developed for the simultaneous estimation of the Erythromycin and Isotretinoin in dosage form. Chromatogram was run through Platisil ODS C18(4.6x250mm,5.0µm) Mobile phase containing Methanol and Orthophosphoric acid taken with in the ratio 70:30 was pumped through column on a flow rate at 1.0 ml/min. Buffer used in this method was Orthophosphoric acid. Ambient Temperature was maintained. Optimized wavelength selected was 245nm. Retention time of Brexanolone is 3.008 min respectively. %RSD of Brexanolone found to be 0.6 respectively. %Recovery was obtained as 100.50 % for Brexanolone respectively. LOD, LOQ values obtained from regression equations of Brexanolone was found to be 2.97 and 9.97 respectively.*

**Key words:** Brexanolone, Chromatogram, RP-HPLC, Method development, validation

## **Introduction**

Although there are many other analytical procedures, such as dissolution testing for drug products or particle size determination for drug substance, these have not been addressed in the initial text on validation of analytical procedures. Validation of these additional analytical procedures are equally important to those listed herein and may be addressed in subsequent documents. Identification tests are intended to ensure the identity of an analyte in a sample. This is normally achieved by comparison of a property of the sample (e.g., spectrum, chromatographic behaviour, chemical reactivity etc) to that of a reference standard. Testing for impurities can be either a quantitative test or a limit test for the impurity in a sample. Either test is intended to accurately reflect the purity characteristics of the sample. Different validation characteristics are required for a quantitative test than for a limit test.

Assay procedures are intended to measure the analyte present in a given sample. In the context of this document, the assay represents a quantitative measurement of the major component in the drug substance. For the drug product, similar validation characteristics also apply when assaying for the active or other selected component(s). [1]The same validation characteristics may also apply to assays associated with other analytical procedures (e.g., dissolution). Related substance by RP-HPLC method development and validation for the detection of impurities in brexanolone drug substance. Very few methods are reported for estimation of both drugs from formulation. We intend to develop RP-HPLC method by simultaneous determination with simple, rapid, greater sensitivity and faster elution.

## Materials and methods

### Method development:

#### Wavelength selection:

UV spectrum of 10 µg / ml Brexanolone in diluents (mobile phase composition) was recorded by scanning in the range of 200 nm to 400 nm. From the UV spectrum wavelength selected as 245. At this wavelength both the drugs show good absorbance.

#### Optimization of Column:

DIKMA Platisil ODSC18 (4.6x250mm,5µm) was found to be ideal as it gave good peak shape and resolution at 1.5 ml/min flow.

### Preparation of buffer and mobile phase:

#### Preparation of 0.1% Ortho phosphoric acid buffer:

Pipetted 1ml of ortho phosphoric acid in 1000 ml HPLC water

#### Preparation of mobile phase:

Mix a mixture of above buffer 300ml (30%), 700ml Methanol (70%) to 1000 ml and degas in ultrasonic water bath for 5 minutes. Filter through 0.45 µ filter under vacuum filtration.

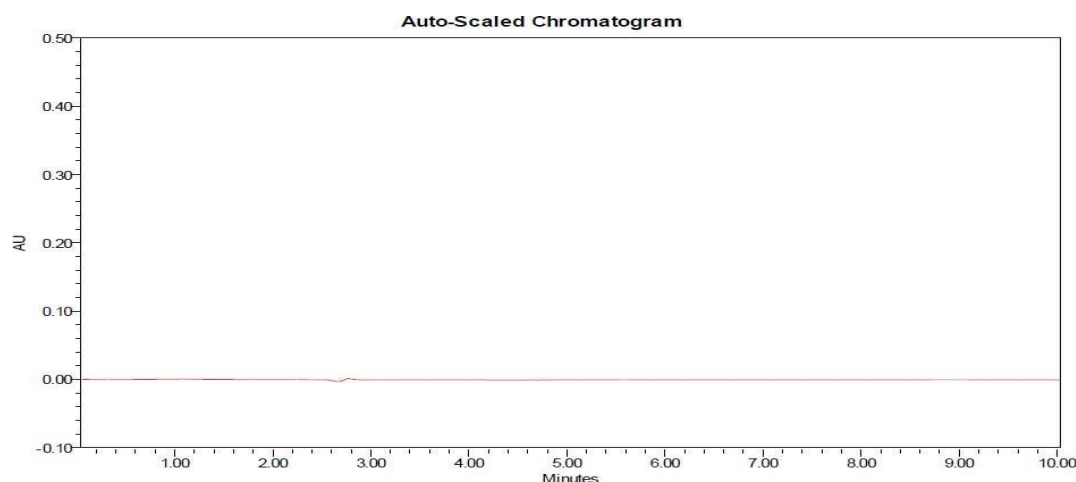
**Diluent Preparation:** Use the Mobile phase as Diluents.

#### Preparation of Brexanolone solution:

##### Preparation of 0.04µg/ml solution:

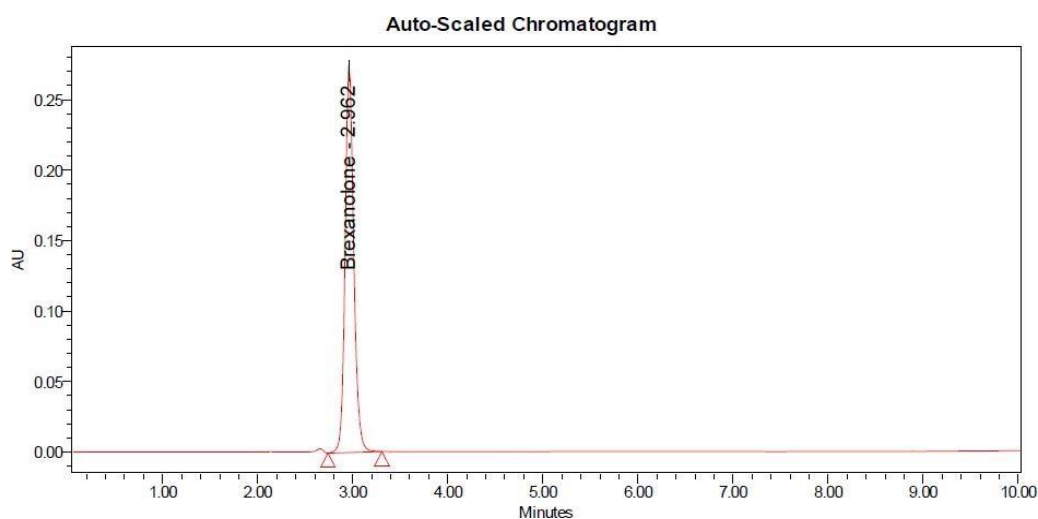
Accurately weigh and transfer 10mg of Brexanolone working standard into a 10ml clean dry volumetric flask add Diluents and sonicate to dissolve it completely and make volume up to the mark with the same solvent. (Stock solution). Further pipette 1ml of the above stock solution into a 10ml volumetric flask and dilute up to the mark with Diluents. Further pipette 3ml of the above stock solution into a 10ml volumetric flask and dilute up to the mark with Diluents. Further pipette 0.5ml of the above stock solution into a 10 ml volumetric flask and dilute up to the mark with Diluent. Further pipette 0.4 ml of the above stock solution into a 10 ml volumetric flask and dilute up to the mark with Diluent.

## Results and discussion

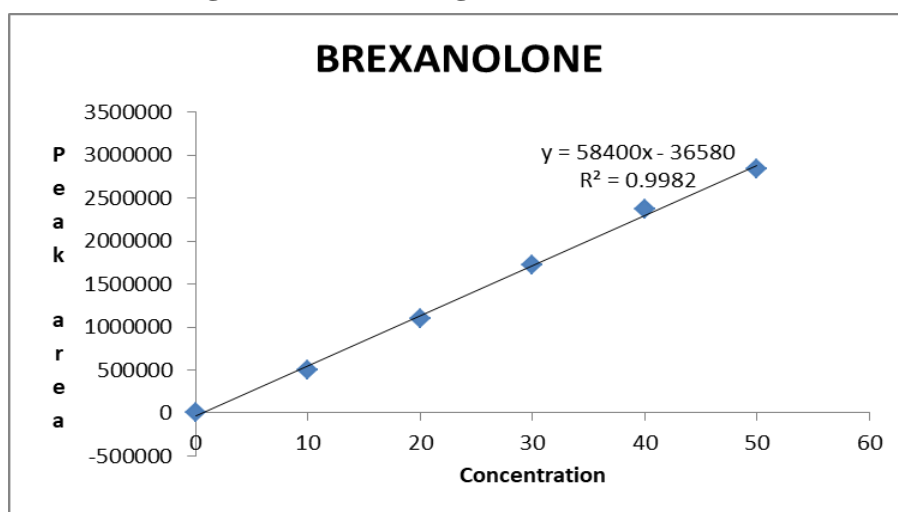


**Figure-1: Chromatogram for system suitability**

Resolution between two drugs must be not less than 2. Theoretical plates must be not less than 2000. Tailing factor must be not more than 2. It was found from above data that all the system suitability parameters for developed method were within the limit.



**Figure-2: Chromatogram for standard**



**Figure-3: Chromatogram of linearity curve**

Correlation coefficient ( $R^2$ ) should not be less than 0.999. The correlation coefficient obtained was 0.999 which is in the acceptance limit.

**Table-1: Results of Precision for Brexanolone**

<b>Injection</b>	<b>Area</b>
Injection-1	1692267
Injection-2	1729767
Injection-3	1692267
Injection-4	1729767
Injection-5	1692267
Injection-6	1729767
<b>Average</b>	1711017.2
<b>Standard Deviation</b>	26516.4
<b>%RSD</b>	1.5

The %RSD for the standard solution is below 1, which is within the limits hence method is precise.

**Table-2: Results of intermediate Precision for Brexanolone**

<b>Injection</b>	<b>Area</b>
Injection-1	1792267
Injection-2	1529767
Injection-3	1692267
Injection-4	1729767
Injection-5	1692267
Injection-6	1719767
<b>Average</b>	1601017
<b>Standard Deviation</b>	31516.4
<b>%RSD</b>	1.9

The % RSD obtained is within the limit, hence the method is rugged.

**Table-3: Accuracy (recovery) data for Brexanolone**

<b>%Concentration(atspecification Level)</b>	<b>Area*</b>	<b>Amount Added(mg)</b>	<b>Amount Found(mg)</b>	<b>% Recovery</b>	<b>Mean Recovery</b>
50%	864883	5	5.07	101.43	100.50
100%	1729767	10	9.92	99.21	
150%	2594650	15	15.13	100.88	

\*Average of three determinations

The percentage recovery was found to be within the limit (97-103%). The results obtained for recovery at 50%, 100%, 150% are within the limits. Hence method is accurate.

**Degradation calculation:**

**Table-4: Degradation calculation**

<b>S.no</b>	<b>Types of forced degradation</b>	<b>Peak Area</b>	<b>% Degradation</b>
1	Standard	1722323	-----
2	Acid	1642002	4.66
3	Base	1647791	4.33
4	Peroxide	1677084	2.63
5	Thermal	1688709	1.95
6	Photo	1646769	4.39

## Summary of the results

<b>Validation Parameters</b>	<b>Result</b>	<b>Limit</b>
Assay(% meanassay)	99.76	90-110%
Linearity	0.999	NMT1
Precision	0.6%	2.0%
Intermediate precision	0.4	2.0%
Accuracy(Percentage recovery)	100.50%	97.0%-103.0%.
LOD	2.97	3
LOQ	9.97	10

The estimation of Brexanolone was done by RP-HPLC. The assay of Brexanolone was performed with tablets and the % assay was found to be 99.76 which shows that the method is useful for routine analysis. The linearity of Brexanolone was found to be linear with a correlation coefficient of 0.999 which shows that the method is capable of producing good sensitivity. The acceptance criteria of precision are RSD should be not more than 2.0% and the method show precision 0.6 for Brexanolone which shows that the method is precise. The acceptance criteria of intermediate precision are RSD should be not more than 2.0% and the method show precision 0.4 for Brexanolone which shows that the method is repeatable when performed in different days also. The accuracy limit is the percentage recovery should be in the range of 97.0% - 103.0%. The total recovery was found to be 100.50% for Brexanolone. [2-4]

## Conclusion

The validation of developed method shows that the accuracy is well within the limit, which shows that the method is capable of showing good accuracy and reproducibility. The acceptance criteria for LOD and LOQ is 3 and 10. The LOD and LOQ for Brexanolone was found to be 2.97 and 9.97. The robustness limit for mobile phase variation and flow rate variation are well within the limit, which shows that the method is having good system suitability and precision under given set of conditions.

## References

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