# RP-HPLC METHOD DEVELOPMENT FOR THE ESTIMATION OF L-ASPARAGINASE FOR INJECTION 10000 IU

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## Abstract:

L-asparaginase is an enzyme used primarily in the treatment of certain types of cancer, particularly acute lymphoblastic leukemia (ALL). The primary aim of this study is to develop a new method for the quantification of L-Asparaginase for injection 10000IU using Reverse phase-High Performance Liquid Chromatography (RP-HPLC) after an extensive literature survey. Method was achieved with Welchrom XB-C18, 250 x 4.6mm 5 $\Box$ m column with run time 15 min at a flow rate 0.6ml/min. Retention time was obtained at 2.782 min. Detector wavelength was set at 225nm. The estimated percent of L- Asparaginase in its marketed formulation was found to be 101.67%. Consequently, the author aimed to develop a robust assay method capable of quantifying the drugs in their commercial pharmaceutical dosage form. This method is designed to be simple, specific, precise, and economical, featuring a shorter run time that makes it ideal for routine quality control applications.

Key words: L-Asparaginase, Oncoginase for Injection 10000 IU, RP-HPLC, & Acute lymphoblastic leukemia.

## 1. Introduction

L-asparaginase is an enzyme used primarily in the treatment of certain types of cancer, particularly acute lymphoblastic leukemia (ALL). It works by breaking down asparagine, an amino acid that some cancer cells rely on for growth. By depleting asparagine levels, L-asparaginase can inhibit the growth of these cancer cells. L-asparaginase is usually administered as an injection, either intramuscularly or intravenously. It is freely soluble in water with several hours to 3-4 days half life. IUPAC name is E-coli L-Asparaginase amidohydrolase and chemical formula is C1377H2208N382O442S17. It is mainly used in

combination with other chemotherapy agents for treating ALL and sometimes for other malignancies. Common side effects may include allergic reactions, pancreatitis, liver function abnormalities, and effects on blood clotting. Patients may also experience nausea, vomiting, or changes in blood sugar levels [1-23].

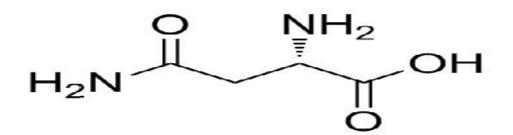


Figure 1. Chemical structure of L-Asparaginase

This Exhaustive literature review [24-37] examines the physicochemical properties of L- Asparaginase and discusses various analytical methodologies used for its quantification. This foundational work supports the validation [38] of a novel reverse-phase high-performance liquid chromatography (RP-HPLC) method for quantifying L-Asparaginase

#### 2. Experimental part:

2.1 Chemicals, Reagents and	Solvents:
Table 1.	Chemicals, Reagents & Standards

Chemicals & Solvents Standards	Make	Grade	Batch No.	Assay
Triethyl amine	Sigma Aldrich	HPLC	614381027L U	99.5%
Acetonitrile	Merck	HPLC	A24B025	≥99.90%
Orthophosphoric acid	Merck	AR	622940712G W	85%w/w
0.9% NaCl	Merck	AR	8489801717	≥99.98%
L-Asparaginase Working Standards	Sigma Aldrich	HPLC	WS-O63	98%
L-Asparaginase injection 10000IU	Chandra Bhag Pharma	gatHIS	LAI2401B	-

S.No	Instrument	Make, Model & Details	Identification No.
1.	HPLC	Waters e2695/Alliance Series	ARD/LC/2021009
2.	HPLC-Column	Welchrom XB-C18, 250 x 4.6mm 5 m	CLL/CCC/64/23
3.	Analytical Balance	Sartorius	ARD/BAL/201806
4.	pH Meter	Thermo Scientific:ORIONSTAR A215	ARD/PHM/202103
5.	Sonicator	PCi Ultrasonic bath chiller Model	ARD/SC/2021005
6.	Shakers and Mixers	Thermo Scientific - MaxQ <sup>™</sup> 4450	ARD/MIX/TFS/09

## 2.2 Analytical Instrumentation & Equipments: Table 2. Analytical Instrumentation & Equipments

**Chromatographic Conditions:** 

Table 5. Chromatographic Conditions				
Column	Welchrom XB-C18, 250x4.6 mm, 5 □m			
Flow rate	0.6ml/min			
Wavelength (λmax)	225 nm			
Run time	15 minutes			
Column Temp. & Sample Temp.	25°C & 5°C			
Injection Volume	20 🗆 L			
Elution	Isocratic			
Diluent	Buffer:Acetonitrile 70:30			

## Table 3. Chromatographic Conditions

2.3 Sample, Standards & Solutions Preparations 10% (v/v) diluted Orthophosphoric acid:

5.9 mL of Orthophosphoric acid was taken in a volumetric flask of 50mL and made till the mark with water.

**Mobile phase A (Buffer):** 2 mL of Triethylamine was taken in 2000mL of water and mixed well. The pH was adjusted to 7.02 with 10% orthophosphoric acid. The mobile phase was filtered with  $0.45\mu$  filter & sonicated for 5 minutes.

Mobile Phase B: Acetonitrile and water was made in 95:5 ratio.

**Diluent:** Mobile phase A & Acetonitrile was mixed in 70:30 ratio.

**Solutions:** 

i) **Standard Stock Solution-1:** Weighed accurately 60.47 mg of L-Asparaginase standard and transferred into volumetric flask of 100mL. 50ml of buffer was added further and shaken vigorously, then 5 mL of Acetonitrile was added and swirled to dissolve. 20 mL of Buffer was added along with 5 mL of acetonitrile again and made upto the mark with buffer.

**ii) Standard-1**: From the above standard stock solution-1, pipetted out 3 mL and transferred into 50 mL of volumetric flask & made till the volume with the diluent.

**iii) Standard stock-2:** 60.52mg of L-Asparaginase was weighed accurately & transferred to volumetric flask of 100ml. Then 50 ml of buffer and added 5 mL of Acetonitrile was added & shaken vigorously. The solution was swirled to dissolve and added 20 ml of Buffer and 5 mL of Acetonitrile again and made till the volume with buffer.

iv) Standard-2: From the above standard stock solution-2, pipetted out 3 ml and transferred into volumetric flask of 50ml. The final volume was made up with diluent.

v) Sample stock-1: Two vials were flipped off the seal and reconstituted each vial with 5 mL of readymade 0.9% Sodium chloride saline solution. The vial was shaken vigorously and transferred the solution carefully into 100 ml volumetric flask. The procedure was repeated for three times with buffer, it was ensured that the vials are clear and added 50 mL of buffer further and shaken vigorously. To this, 10 mL of Acetonitrile was added and volume was made to final with the buffer.

vi) Sample solution-1: From the above Sample Stock Solution-1, 3mL was pipetted and transferred into volumetric flask of 100 mL. The final volume was made with the diluents.

vii) Sample stock-2: 2 vials were taken & flipped off the seal, and reconstituted with 5mL of readymade 0.9% Sodium chloride saline solution. Then they were shaken vigorously and transferred the solution carefully into 100 ml volumetric flask. The procedure was repeated for three times with buffer, it was ensured that the vials were clear and then, 50 mL of buffer was added and shaken vigorously.10 mL of Acetonitrile was added to it and the final volume was made up with buffer.

**viii)** Sample solution-2: From the above Sample Stock Solution-2, 3mL was pipette into 100ml volumetric flask and the volume was made upto mark with the diluent.

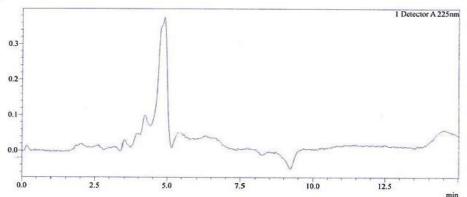
## 3. Results and Discussion

## 3.1 Data summary of specificity & system suitability

 Table 4. Specificity & System Suitability Summary Data

Specificity & System s	uitability		
		Results	Acceptance criteria
Specificity			·
Parameter	Standard Solution		
Identification	&		RT obtained with the sample solution should
Retention Time (RT	Г)L-Asparaginase1 & 2	2.752 &	be comparable with the standard solution.
Conformation.		2.735	
	Sample Solution		
	L-Asparaginasefor in 10000IU	j2.731	
	÷	Single point	
Peak purity index		threshold	
Standard Solution			Peak purity should pass. Peak purity index
Peak Purity	1.0000	0.9998	should be greater than single point threshold.
Sample Solution			

Peak Purity	0.99999	0.99998		
Blank andPlacebo Solution.	L-Asparaginase	NIL		Blank and placebo solution must not elute at the retention time of the analyte peak.
Interference				
System suitability				·
% RSD of standard respectively	solution 1 & 2	0.349 0.733	&	% RSD should be NMT 2.0%
% RSD of the L-ASPase	sample solution	0.149		% RSD for RT should be NMT1.0%
The Number of theoret peak in L-ASPase standa	1	1423 1194	&	The theoretical plates for main peak in standard solution should be NLT 1000
The tailing factor for r standard solution 1& 2	nain peak in L-ASF	1.197 1.293	&	The tailing factor for main peak in standard solution should be NMT 0.8 and 2.0
mV		Chromatog LC_26_1304	gram 102_led	1 Detector A 225nm



**Figure 2. Chromatogram for Blank** 

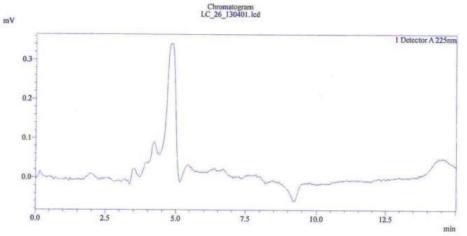


Figure 3. Chromatogram for Placebo

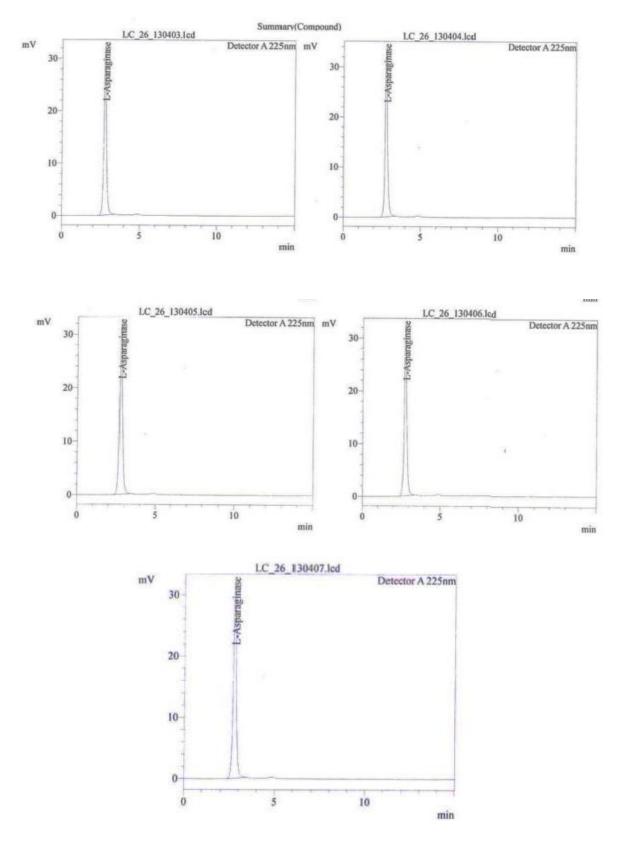
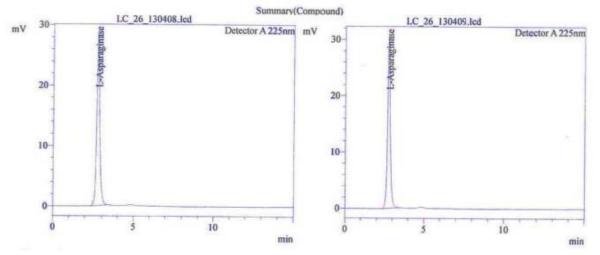
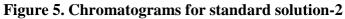


Figure 4. Chromatograms for standard solution-1





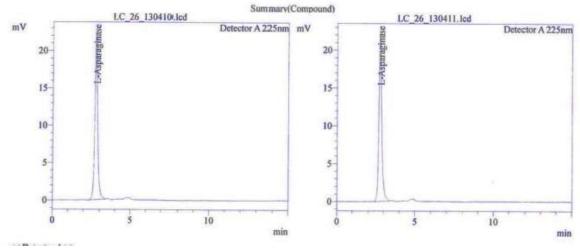


Figure 6. Sample Solutions 1&2 chromatograms

Solutions	RT	Area	Tailing factor	Theoretical plates
Standard solution injection-1	2.748	366860	1.208	1387
Standard solution injection-2	2.754	368024	1.213	1549
Standard solution injection-3	2.753	368874	1.189	1362
Standard solution injection-4	2.753	365676	1.186	1417
Standard solution injection-5	2.750	368331	1.187	1402
Mean	2.752	367553	1.197	1423
Standard deviation	0.003	1282	0.013	73
% Relative Standard deviation	0.091	0.349	1.065	5.143
Bracketing Standard solution	2.738	371988	1.297	1171

Table 4. System Suitability of Standard Solution-1

Solutions	RT	Area	Tailing factor	Theoretical plates
Standard solution injection-1	2.727	372349	1.362	1090
Standard solution injection-2	2.744	368507	1.224	1298
Mean	2.735	370428	1.293	1194
Standard deviation	0.012	2716	0.098	147
% Relative Standard deviation	0.438	0.733	7.587	12.312
	1			

#### Table 5. System Suitability of Standard Solution-2

## Table 6. L-Asparaginase Sample Solution Specificity

Solutions	RT	Area	Tailing factor	Theoretical plates
Sample solution injection-1	2.734	285870	1.334	1160
Sample solution injection-2	2.729	286475	1.460	1060
Mean	2.731	286173	1.397	1110
Standard deviation	0.003	427	0.089	70
% Relative Standard deviation	0.127	0.149	6.392	6.326

## **3.2 Precision**

Precision describes the closeness between serial measurements that were obtained from multiple sampling conditions from the homogeneous sample.

## **3.2.1 System precision:**

System precision describes the instrument performance under the prescribed chromatographic parameters. System precision studies were carried out by single preparation of standard solution and injecting the sample under the same conditions with six determinations.

## 3.2.2 Method Precision:

Method Precision studies were carried out with test solution of six preparations and injecting the sample under the same chromatographic conditions.

	cision summary	or L-Asparaginase			
System precision	Results	Acceptance limits			
% RSD of standard solution	0.582	% RSD of the standard for six replicate			
(6 replicates)		injections should be NMT 2.0%			
No. of theoretical plates for main		Theoritical plates should be NLT 1000			
peak in standard solution	1381				
Tailing factor	1.213	The tailing factor for main peak in standard solution should be			
		NMT 0.8 and 2.0			
Method precision					
Calculated %RSD for % Assay		%RSD should be NMT 2.0 with all the			
content L-ASPase from six sample	1.468	individual values within limit.			
preparations					

Table 7. Precision summary of L-Asparaginase

#### 4. Proposed procedures for marketed Pharmaceutical Formulation:

The Marketed Formulation (Oncoginase 10000IU) was analysed separately by injecting the standard & sample solutions of 10µ L volume into the HPLC. The quantity of the drug found in the formulation was calculated by comparison of the peak area of the standard and sample.

Table 8. Results of Assay in dosage form						
Formulation Batch No Label claim(mg) % Assay* Est am						
inase injection10000IU	LSI2406CB	Oncoginase 10000IU	101.67	10167.37 IU		

Tabla 8	<b>Results</b> (	of A c	ow in	docodo	form
I abic o.	ICSUITS (	JI AS	say m	uusage	IUIII

\*= Average assay % of 3 replicate injections of individual batch.

### Conclusion

The precision results reflected minimal variability, underscoring the method's reliability for routine analysis. Key parameters such as specificity and precision were within acceptable limits, confirming the method's efficacy in quantifying L-Asparaginase accurately. Overall, the HPLC method provided a comprehensive analytical tool for the quality control of Oncoginase for injection 10000 IU ensuring that pharmaceutical products meet regulatory standards.

### **Conflict of Interest**

Authors declared that there was no conflict of interest.

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