E-ISSN: 0975-8232; P-ISSN: 2320-5148



PHARMACEUTICAL SCIENCES



Received on 20 June, 2016; received in revised form, 15 August, 2016; accepted, 27 August, 2016; published 01 December, 2016

METHOD DEVELOPMENT AND VALIDATION FOR THE ESTIMATION OF IDELALISIB IN RABBIT PLASMA BY HPLC

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Keywords:

Idelalisib, Ibrutinib, Anticaner, Rabbit Plasma

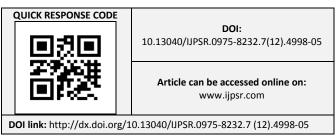
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ABSTRACT: A simple, rapid, sensitive, and accurate high performance liquid chromatography was developed for determination of Idelalisib (IL) in rabbit plasma using Ibrutinib as internal standard (IS). Idelalisib is a phosphoinositide 3-kinase inhibitor indicated in the treatment of chronic lymphocytic leukemia (CLL), relapsed follicular B-cell non-Hodgkin lymphoma (FL), and relapsed small lymphocytic lymphoma (SLL). The analytes and IS were separated on a ODS (250 mm \times 4.6 mm, 5 μ m) column using Mobile phase composition as Buffer and Acetonitrile in the ratio of 85:15 v/v%. The total chromatographic runtime is 10.0 min with retention time for IL and IS at 7.195, and 5.435 min, respectively with a flow rate 1ml/min. The method is validated over a dynamic linear range of 0.02-4 μ g/mL for IL with a correlation coefficient of r2 0.999. The method was validated as per the USFDA guidelines and the results were within the acceptance criteria for selectivity, sensitivity, linearity, precision, accuracy, recovery stability of solution and stability of solution in plasma.

INTRODUCTION: Idelalisib is an oral phosphatidylinositol 3-kinase delta $(PI3K\delta)$ inhibitor ¹. It had clinically significant activity with an acceptable toxicity profile in patients with relapsed or refractory chronic lymphocytic leukemia. follicular B cell non-Hodgkin's lymphoma and small lymphocytic lymphoma ^{2, 3}. Idelalisib has demonstrated activity in indolent B-NHL (iB-NHL) and is approved for use as monotherapy in patients with follicular lymphoma lymphocytic lymphoma and in small combination with rituximab in patients with chronic lymphocytic leukemia 4.



Treatment of lymphoma cells with Idelalisib has been shown to result in inhibition of chemotaxis and adhesion, and reduced cell viability. Chemically Idelalisib is 5-fluoro-3-phenyl-2-[(S)-1-(9H-purin-6-ylamino)-propyl] – 3 H - quinazolin- 4 one, with molecular formula $C_{22}H_{18}FN_7O^{5,6}$.

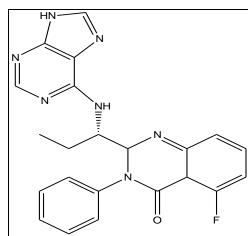


FIG.1: STRUCTURE OF IDELALISIB

Ibrutinib is a bruton tyrosine kinase (BTK) inhibitor effective in patients with chronic lymphocytic leukemia ⁷. Ibrutinib is an orally available small molecule that forms an irreversible covalent bond with the Cys481 of BTK kinase ⁸. Ibrutinib is also indicated for the treatment of patients with Waldenström's Macroglobulinemia (WM). IUPAC Name 1-[(3R)-3-[4-amino-3-(4-phenoxyphenyl)-1H-pyrazolo [3,4-d]pyrimidin-1-yl]piperidin-1-yl]prop-2-en-1-one ⁹.

1.1 Reagents and chemicals:

The pure samples of Idelalisib and Ibrutinib were obtained from Selleckchem.com LLC supplied by Pro lab marketing. HPLC grade Acetonitrile, HPLC grade Methanol and all other chemicals were obtained from Merck chemical division, Mumbai. HPLC grade water obtained from Milli-Q water purification system was used throughout the study.

1.2 Instrumentation:

Chromatography was performed with waters 2695 HPLC provided with high speed auto sampler, column oven, degasser and & 2996 PDA detector to provide a compact and with class Empower-2 software.

1.3 Chromatographic method:

The separation was carried on ODS C18 analytical column (250mm×5mm×4.6 μ m) using mobile phase Buffer and Acetonitrile in the ratio of 85:15%v/v and flow rate is 1 ml/min. The Injection volume was 10 μ l and the run time was 10 min. The temperatures of column and auto sampler were maintained at 30°C and 5°C respectively. The detection was carried at wavelength 270 nm.

1.4 Buffer Preparation:

1ml of 0.1% Perchloric acid was transferred into 1000mL volumetric flask and made the volume to produce 1000mL with water.

1.5 Preparation of Idelalisib Stock solution:

The standard stock solution of Idealisib was prepared at 1mg/mL with mobile phase. The stock is further diluted with mobile phase to obtain 0.1mg/mL solution.

1.6Preparation of Idelalisib Spiking Solutions (4.6 µg/mL to 920µg/mL):

From the above Idelalisib stock solution 2 (0.1mg/mL) take out 0.460ml, 1.380ml, 2.30ml

and from the stock 1(1mg/mL) solution take out 1.150ml, 2.300ml, 4.600ml, 6.900ml and 9.200 ml was pipette and transferred to 8 individual 10 ml volumetric flask and make up the volume up to the mark with mobile phase to produce 4.6 µg/mL, 13.8 µg/mL, 23 µg/mL, 115 µg/mL, 230 µg/mL, 460 µg/mL, 690 µg/mL and 920 µg/mL.

1.6 Extraction procedure:

To $250\mu l$ of drug free plasma $50\mu l$ of internal standard & $10\mu l$ of Idelalisib was added. To the mixture 2 ml of Acetonitrile was added, subjected to cyclomixer for 15 sec. Then vertexed for 2 min and finally centrifuged for 3 min at 3200 rpm speed. After the centrifugation the organic layer was collected and directly injected 10 μL into HPLC.

1.7 Methodology for Analysis:

A thorough and complete method of validation was following the USFDA guidelines. The method was validated for system suitability, auto sampler carryover, specificity and screening of biological matrix, sensitivity, matrix effect, linearity, precision and accuracy, recovery of analyte and internal standard, ruggedness on precision accuracy and linearity, reinjection reproducibility and stability on day zero, long batch, LT at -28°C and LT at -80°C.

System suitability was done by MQC level sample as six homogenous injections and will see the %RSD values for retention time and response of analyte and internal standard. Auto sample carryover was done by ULOQ and LLOQ level and check whether drug is remains or not in system.

Specificity and screening of biological matrix was done by LLOQ level of sample and check it for any interference of blank and sample response. Sensitivity was done by LLOQ level sample for to know the lowest limit of detection and calculate the %mean accuracy and %CV. Matrix effect on analyte quantification with respect to consistency in signal (suppression/ enhancement), the matrix effect was checked in six different lots of Idelalisib plasma three replicates, each at LQC and HQC levels were prepared from these lots of plasma (total 36 QC samples) and checked for the accuracy in terms of % bias in all the QC samples.

E-ISSN: 0975-8232; P-ISSN: 2320-5148

Linearity of the method was determined by analysis of standard plots associated with an 8-point standard calibration curve. Intra-batch and interbatch accuracy and precision was evaluated at five different concentrations levels (LLOQ, LQC, MQC and HQC) in six replicates for both the analytes. Mean values were obtained for calculated drug concentration over these batches. The accuracy and precision was calculated and expressed in terms of % Accuracy and coefficient of variation (% CV), respectively. Recovery of the analytes from the extraction procedure was performed at LQC, MQC, and HQC levels. It was evaluated by comparing peak area of extracted samples (spiked before extraction) to the peak area of un extracted samples (quality control working solutions spiked in extracted plasma). Ruggedness can be done by changing the person to person for linearity, precision and accuracy in the levels of ULOQ, LQC, MQC and HQC.

Stability studies was performed as Zero hours, Long batch, LT at -28°C and LT at -80°C. Day zero having two sample with six replicates of HQC and LQC levels. Long batch have 35replicates of LLOQ, LQC, MQC and HQC level of samples with %Mean accuracy. LT at -28°C and LT at -80°C have HQC and LQC level with % Stability finding by comparison sample and stability sample.

2. RESULTS AND DISCUSSIONS:

3.1 System suitability: Six replicate samples of middle quality control samples of concentration along with internal standards were injected and % CV was calculated. The % CV of the retention time of analyte and IS was found to be $\leq 2.00\%$. The % CV of the peak area ratio of analyte to IS was found to be $\leq 5.00\%$. The results were found to be within limits and are summarized in **Table 1**.

TABLE 1: SYSTEM SUTABILITY DATA

Comple Nome	Analyte			IS	Ausa Dadia
Sample Name	Area	RT (min)	Area	RT (min)	Area Ratio
	57496	7.19	32172	4.31	1.7871
	57495	7.18	32170	4.30	1.7872
MOG	57492	7.17	32168	4.32	1.7872
MQC	57487	7.19	32162	4.30	1.7874
	57484	7.18	32160	4.32	1.7874
	57481	7.20	32161	4.31	1.7873
MEAN		7.185		4.310	1.78729
SD		0.0072		0.0094	0.000115
%CV		0.10		0.22	0.01

The carryover experiment was done to ensure that it does not affect the accuracy and precision. There

was no carryover observed. The results were presented in **Table 2.**

TABLE 2: AUTO SAMPLER CARRYOVER

Sample ID	Peak	Peak Area Drug IS Drug		ryover
Sample 1D	Drug			IS
		Unextracted samples		
RS	0	0	N/A	N/A
AQ ULOQ	221216	48156	0.00	0.00
RS	0	0	0.00	0.00
AQ LLOQ	2956	48152	N/A	N/A
		Extracted samples		
STD Blk	0	0	N/A	N/A
ULOQ	201216	32172	0.00	0.00
STD Blk	0	0	0.00	0.00
LLOQ	1956	32170	N/A	N/A

3.2 Specificity and screening of biological matrix: Response of interfering peaks in standard blank at the retention time of analyte should be

 \leq 20.00 % of that in LLOQ. Response of interfering peaks in standard blank at the retention time of IS should be \leq 5.00 % of that in LLOQ. All the

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samples were found to be free of interference at retention time of analyte in blank samples. The results were shown in **Table 3**.

TABLE 3: SPECIFICITY AND SCREENING OF BIOLOGICAL MATRIX

Sample	Res	Response		rference	- Pass/Fail
Sample	Drug	ISTD	Drug	ISTD	- Pass/Fall
STD Blk1	0	0	0.00	0.00	Pass
LLOQ1	1956	32172	0.00	0.00	rass
STD Blk2	0	0	0.00	0.00	Pass
LLOQ2	1952	32168	0.00	0.00	rass
STD Blk3	0	0	0.00	0.00	Pass
LLOQ3	1991	32160	0.00	0.00	Pass
STD Blk4	0	0	0.00	0.00	D
LLOQ4	1982	32194	0.00	0.00	Pass
STD Blk5	0	0	0.00	0.00	D
LLOQ5	1943	32168	0.00	0.00	Pass
STD Blk6	0	0	0.00	0.00	D
LLOQ6	1968	32173	0.00	0.00	Pass

Sensitivity: The accuracy and precision of Idelalisib at LLOQ level was found to be 7.58 % CV and % Mean accuracy was found to be 100.58. Acceptance Criteria is at least 67 % (4 out of 6) of samples should be within 80.00-120.00 %. % Mean

accuracy should be within 80.00-120.00 %. % CV accuracy should be ≤ 20.00 %. The results comply with acceptance limit and were incorporated in **Table 4**.

TABLE 4: SENSITIVITY DATA

	LLOQ			
Sa1a	Nominal Concentration (µg/mL)			
Sample	0.020			
	Calculated Concentration (µg/mL)			
1	0.018			
2	0.020			
3	0.021			
4	0.022			
5	0.020			
6	0.021			
N	6			
Mean	0.0201			
SD	0.00153			
% CV	7.58			
% Mean Accuracy	100.58			

3.4 Matrix effect: The matrix effect data of HQC and LQC were presented in **Table 5**. The Acceptance Criteria is at least 67 % (2 out of 3) of samples at each level should be within 85.00-115.00 %. At least 80 % (5 out of 6) of the matrix

lot should be within the acceptance criteria. The % mean accuracy of back calculated concentration of LQC and HQC samples prepared from different biological matrix lots should be within 85.00-115.00 %.

TABLE 5: MATRIX EFFECT

G N			7.00
S. No.	Plasma Lot No.	HQC	LQC
		Nominal Concentration (μg/mL)	
		3.000	0.100
		Nominal Concentrat	ion Range (µg/mL)
		(2.550-3.450)	(0.085 - 0.115)
		Calculated Concentr	ation (ng/mL) (n=3)
1	LOT1	2.740	0.097
2	LOT2	2.964	0.101

3	LOT3	2.995	0.102
4	LOT4	2.871	0.095
5	LOT5	3.317	0.105
6	LOT6	3.174	0.102
	Mean	3.0102	0.1001
	SD	0.30948	0.00907
	% CV	10.28	9.06
%	Mean Accuracy	100.34	100.11
N	No. of QC Failed	0	0

3.5 Linearity: The calibration curve was found to be linear at range $0.02 - 4\mu g/mL$ with correlation coefficient (r^2) 0.999. The linear graph is

represented in **Fig.2**. The data of calculated calibration standards are presented in **Table 6**.

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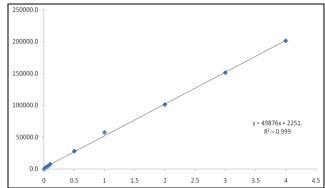


FIG.2: CALIBRATION PLOT FOR CONCENTRATION v/s AREA RATIO

TABLE 6: LINEARITY

S. No.	Conc.	Back Calc	ulated Concentration	on (μg/mL)	Avg.	%CV	% Mean
	(µg/ml)	1	2	3	_		Accuracy
1	0.02	0.018	0.020	0.021	0.0197	7.77	98.33
2	0.06	0.054	0.060	0.068	0.0607	11.58	101.11
3	0.1	0.092	0.102	0.112	0.1020	9.80	102.00
4	0.5	0.468	0.480	0.541	0.4963	7.89	99.27
5	1	0.862	1.125	1.148	1.0450	15.21	104.50
6	2	1.765	1.984	2.247	1.9987	12.07	99.93
7	3	2.580	3.195	3.389	3.0547	13.83	101.82
8	4	3.542	3.860	4.430	3.9440	11.41	98.60

3.6 Precision: The % CV of estimated concentrations for all four level quality control samples with six replicates for analyte was within 8.34 to 12.31%. The % mean accuracy for LLOQ, LOQ, MOQ and HQC was within 97.50% to 103.02%. For inter day precision and accuracy the % CV and accuracy results of all quality control samples were in between 8.36 to 11.44 and 99.61 to

102.40 respectively. The acceptance criteria is that at least 67 % of total QC samples should be \leq 15.00 % and for the LLOQ, should be \leq 20.00 %. % Mean accuracy for LQC, MQC and HQC samples should be within 85.00-115.00 % and for the LLOQ sample should be within 80.00-120.00 %. The data of precision and accuracy is complied in **Table 7**.

TABLE 7: PRECISION INTERDAY AND INTRADAY DATA

	HQC	MQC	LQC	LLOQ
		Nominal Concent	ration (µg/mL)	
	3.000	1.000	0.100	0.020
Day 1 (n=6)				
Mean	3.0907	0.9915	0.1003	0.0203
SD	0.25762	0.11947	0.00918	0.00250
%CV	8.34	12.05	9.15	12.31

% Mean Accuracy	103.02	99.15	100.33	101.67
Day-2 (n=6)				
Mean	3.0660	1.0098	0.0990	0.0200
SD	0.28528	0.12326	0.00787	0.00237
%CV	9.30	12.21	7.95	11.83
% Mean Accuracy	102.20	100.98	99.00	100.00
Day-3 (n=6)				
Mean	3.0593	1.0110	0.0995	0.0195
SD	0.27564	0.12282	0.01003	0.00187
%CV	9.01	12.15	10.09	9.59
% Mean Accuracy	101.98	101.10	99.50	97.50
Between Batch Precision and A	ccuracy			
N	18	18	18	18
Mean	3.0720	1.0041	0.0996	0.0199
SD	0.25690	0.11484	0.00854	0.00215
%CV	8.36	11.44	8.57	10.80
% Mean Accuracy	102.40	100.41	99.61	99.72

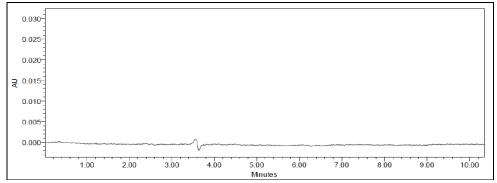


FIG.3: CHROMATOGRAM OF STANDARD BLANK

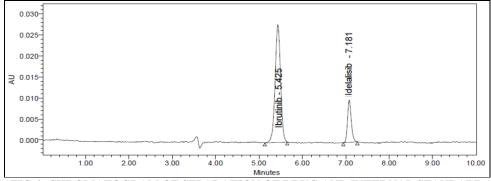


FIG.4: CHROMATOGRAM OF SEPARATION OF DRUG AND INTERNAL STANDARD

3.7 Recovery of analyte: The result of recovery study is given in **Table 8** and **9**. The results are within acceptance limit. The acceptable limit was

% CV of recovery at each QC level and for IS should be \leq 15.00 %. The overall mean recovery % CV for all QC levels should be \leq 20.00 %.

TABLE 8: RECOVERY OF ANALYTE

	НС	QC	MQ	C	LQ	С
Sample	Un extracted	Extracted	Un extracted	Extracted	Un extracted	Extracted
	Response	Response	Response	Response	Response	Response
Mean (n=6)	186199.3	151624.2	63151.8	57828.0	8897.2	7549.3
SD	532.26	491.63	235.17	735.52	69.41	78.20
% CV	0.29	0.32	0.37	1.27	0.78	1.04
% Mean Recovery	81.	43	91.5	57	84.8	35

Overall % Mean Recovery	85.951	
Overall SD	5.1580	
Overall % CV	6.00	

TABLE 9: REOVERY OF INTERNAL STANDARD

S. no.	Un extracted Area Ratio	Extracted Area Ratio
Mean (n=6)	35742.7	32575.7
SD	339.40	481.29
% CV	0.95	1.48
% Mean Recovery	91.	14

3.8 Ruggedness:

The results of ruggedness study for Idelalisib was within acceptance limit. The data is represented in

Table 10. The precision and accuracy values for different columns with different analysts ranged from 5.61% to 12.49% and 98.63% to 104.13%.

E-ISSN: 0975-8232; P-ISSN: 2320-5148

TABLE 10: RUGGEDNESS DATA

LE 10: RUGGEDNESS DATA						
P&A ID	HQC	MQC	LQC	LLOQ		
	Nominal Concentration (µg/mL)					
	3.000	1.000	0.100	0.020		
	Nominal Concentration Range (μg/mL)					
	(2.550 - 3.450)	(0.850 - 1.150)	(0.085 - 0.115)	(0.016 - 0.024)		
	Calculated Concentration (µg/mL)					
Different Column						
Mean (n=6)	2.9863	1.0081	0.1014	0.0200		
SD	0.25744	0.08270	0.00767	0.00200		
% CV	8.62	8.20	7.56	10.00		
% Mean Accuracy	99.54	100.81	101.42	100.00		
Different Analyst						
Mean (n=6)	3.1240	0.9863	0.1050	0.0207		
SD	0.17517	0.6649	0.00623	0.00258		
% CV	5.61	6.74	5.93	12.49		
% Mean Accuracy	104.13	98.63	105.02	103.33		

3.9 Stability studies: Zero hours, Long batch, LT at -28°C and LT at -80°C results of LQC, MQC and HQC were found to more than 95%, which is

within acceptance limit. The results were complied in **Table 11**.

TABLE 11: STABILITY DATA

Sample	Nominal Concentration (µg/mL)	Mean Calculated Concentration (μg/mL) ±SD. (n=6)	% CV		
Stability on day zero					
HQC	3.000	3.1153±0.142	4.57		
LQC	0.100	0.1022 ± 0.008	7.72		
Long batch at -28°C					
HQC	3.000	3.0102±0.155	5.17		
LQC	0.100	0.0979 ± 0.012	12.06		
Long batch at -80°C					
HQC	3.000	3.0753±0.162	5.76		
LQC	0.100	0.0952 ± 0.006	6.76		

CONCLUSION: The objective of this work was to develop a simple, cost-effective, rugged and sensitive method for determination of Idelalisb in plasma by using Ibrutinib as internal standard. The work shows less run time while comparing with

other work articles. The total chromatographic runtime is 10.0 min with retention time for Idelalisib and IS at 7.185, and 4.310 min, respectively. The method is validated over a dynamic linear range of 0.02 - 4 µg/mL for

Idelalisib with a correlation coefficient of r² 0.999. The intra-batch and inter-batch precision (%CV) across five levels (LLOQ, LQC, MQC, HQC, and ULOQ) is less than 13.53. This can be validated according to USFDA guidelines.

ACKNOWLEDGEMENTS: The authors are very much thankful for management of V.V. Institute of Pharmaceutical sciences, and Spectrum Pharma, India, for supporting to do this work.

CONFLICT OF INTEREST: The authors declare that there is no conflict of interests.

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How to cite this article:

Suneetha A and Sharmila D: Method development and validation for the estimation of Idelalisib in rabbit plasma by HPLC. Int J Pharm Sci Res 2016; 7(12): 4998-05.doi: 10.13040/IJPSR.0975-8232.7(12).4998-05.

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