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# Original Article



Correlation of results between validated in-house analysis method with new pharmacopeia monograph for analysis of Sitagliptin Phosphate API

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## Abstract

Having validated analysis methods for medicinal ingredients is attractive for pharmaceutical companies. When a new molecule is introduced to the market, there is not any pharmacopeial analysis method for that. After publishing official methods, the correlation between validated in-house methods and the official one could establish the value of the in-house method. Sitagliptin phosphate is a new antidiabetic pharmaceutical ingredient and many pharmaceutical companies are trying to manufacture high-quality dosage forms using this agent. In the present study, a full validated in-house method for analysis of sitagliptin phosphate Active Pharmaceutical Ingredient (API) is presented and the method is compared with newly published United State Pharmacopeia (USP) monograph. Results show that the in-house method is correlated with the USP method with regard to assay study and even could separate and detect more probable impurities in the sample. In brief, a full analytical method validation based on USP general chapter (<1225>) was done on the developed analysis method and a calibration curve was plotted successfully with a reasonable R2 equal to 0.9993 and the equation of the curve was Y = 3.4588X + 30.099. Then precision, accuracy, and robustness were studied. The mobile phases, column, column temperature, sample preparation of the solvent, as well as detector wavelength, are different in two methods. It seems that the validated method could be a valuable alternative method for USP method depending on users facilities for analysis. It seems with presenting of this method, more pharmaceutical research centers will be able to analyze sitagliptin with a high degree of assurance.

**Keywords:** Sitagliptin Phosphate, Analytical Method Validation, Assay, Impurities, HPLC, Pharmacopeial analytical method.

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## Introduction

Sitagliptin Phosphate, previously identified as MK-0431 and marketed under the trade name Januvia, is an oral antihyperglycemic drug. Sitagliptin was approved by the U.S. Food and Drug Administration (FDA) on October 17,

2006. Sitagliptin works to competitively inhibit the enzyme dipeptidyl peptidase 4 (DPP-4). This enzyme breaks down the incretins GLP-1 and GIP, gastrointestinal hormones released in response to a meal. By preventing GLP-1 and

HPLC Column		Mobile phase composition	composition	Flow rate		Column temp.	emp.
USP Method	In-house method	USP Method	In-house method	USP Method	In-house method	USP Method	In-house method
4.6-mm*15-cm;5-μ packing L 10	4.6-mm*15- cm;5-μ pack- ing L 1	Mobile phase: (Acetonitrile: buffer) (15:85) Buffer: 1.36 g/L monobasic potas- sium phosphate, adjusted with phosphoric acid to a pH of 2.0	Mobile phase A: Dissolve 11.2 ml /L triethylamine and add 5 ml phosphoric acid, adjusted with phosphoric acid to a pH of 3.0 Mobile phase B: (Acetonitrile: Mobile phase A) (70:30)	1 ml/ min	1 ml/ min	30 ° C	Room tempera- ture
Solvent		Sitagliptin peak retention time	tention time	Detector		Injection volume	olume
USP Method	In-house method	USP Method	In-house method	USP Method	In-house method	USP Method	In-house method
(Acetonitrile:dilute phosphoric acid) (5:95) Dilute phosphoric acid: 1ml/L phos- phoric acid	Mobile phase B	7-8 min	11-13 min	UV 205 nm	UV 220 nm	20 µL	20 µL

#### Table 1: in-house analysis method in comparison to USP method (USP 2006)

Assay				Related				Turnition	CONO		
La Pto	Com DT	Ctd amon	Com anoo	Impurities RT	τ		Std RT	Ampurtues area	arca		Std area
IN me		ou area	Dalli area	1	5	3	I	1	5	3	
11.12	11.72	414.03	432.06	12.47	14.20	16.61	13.77	5213	1627	1064	75443
11.12	11.18	410.58	433.44	12.45	14.19	16.61	13.77	5224	1620	1059	73979
11.10	11.15	408.64	436.76	12.46	14.18	16.56	13.73	5224	1611	1082	75762
Average		411.08	434.09	Average		-	-	5220.33	1619.33	1068.33	75061.33
RSD		0.66	0.56	RSD				0.12	0.50	1.13	1.27
Result		77 ± 1.28	72 ± 1.84	Each impurity: 0.007 %	ity: 0.007 %			Total impu	Total impurity: 0.01 %		
Sample	e 1 (QC-ST	G-95003)	Sample 1 (QC-STG-95003) ( USP method )	( po							
Assay				Related							
Ta Pts	Com DT	Ctd area	Com area	Impurities RT	T	Std RT		AIII purtues area	arca	Std area	
				1	7			1	5		
11.35	11.36	4866281	4605156	5.723	9.303	11.032		331	420	8149	
11.35	11.34	4885179	4613642	5.691	9.247	11.040		331	420	8276	
11.32	11.31	4921541	4614962	5.703	9.284	11.043		330	417	8265	
Average		4891000	4611253	Average				331	419	8230	
RSD		0.11	0.12	RSD				0.17	0.41	0.65	
Result		99.12%		Each impurity: 0.004 %	ity: 0.004 %			Total: 0.007 %	17 %		

## Table 2: Analysis data for sample 1 using in-house and USP method

	Sample	; 2 (QC	S-STG-9	Sample 2 (QC-STG-95004) ( in-house method )	in-hou	se met	( poq															
	Assay				Related	_						Turnitio	0040									
		Sam	Std	Sam	Impuri	ties RT					Std RT	and mutan	o al ca					Std area				
		RT	area	area	T	7	3	4	w	9		1	5	3	4	N	9					
		10.57	501.28	571.48	12.37	12.47	13.03	14.20	14.75	16.61	13.79	5213	1627	1064	1778	16468	755	73395				
		10.58	499.79	571.26	12.38	12.47	13.03	14.20	l	16.62	13.78	5224	1620	1059	1773	16249	756	73789				
$\varepsilon$ $11.08$ $49.98$ Average $1619.33$ $1619.33$ $1619.33$ $0.66$ $0.25$ <b>RSD</b> $0.12$ $0.012$ $0.001$ $0.0101$ $0.0101$ $0.0101$ $0.0101$ $0.0101$ $0.0101$ $0.001$ $0.001$ $0.001$		10.55	498.85	579.63	12.35	12.43	13.00	14.17	l	16.59	13.78	5224	1611	1082	1790	16633	778	72799				
$ \left  \begin{array}{c c c c c c c c } \hline 0.66 & 0.25 & \textbf{RSD} \\ \hline 100.26 \ \& D & \textbf{Each impurity} 0.02 \ \& & \textbf{Total impurities} 0.0 \\ \hline 100.26 \ \& & \textbf{Each impurity} 0.02 \ \& & \textbf{Total impurities} 0.0 \\ \hline 100.26 \ \& & \textbf{Each impurity} 0.02 \ \& & \textbf{Total impurities} 0.0 \\ \hline 1 & \textbf{Total impurities} $	Average		411.08	499.98	Averag	e						5220.33	1619.33	1068.33	1783	16450	763	75061.33				
100.26 %Each impurity: 0.02 %Total impurities: 0.0le 2 (QC-STG-95004) ( USP method )Related )Total impurities: 0.0le 2 (QC-STG-95004) ( USP method )Related )Related (II.032)lo 3 m RTStd areaRTRthriteslo 3333746.834242.289.189lo 83333746.834242.289.189lo 83333744.384242.289.189lo 83333744.384243.739.126lo 83333740.754243.739.126lo 83333740.754243.739.126lo 83333743.994243.769.143lo 83333743.994243.759.143lo 83333743.994243.76Averagelo 83333743.994243.76Averagelo 9.98%0.04RSD	RSD		0.66	0.25	RSD							0.12	0.50	1.13	0.34	1.17	1.70	1.27				
Ide 2 (QC-STG-95004) ( USP method )         Related         Related         F       Related       Related $\Gamma$ Sam RT       Std area       Bam area $0$ 10.8333       3746.83       4242.28       9.189       11.032 $0$ 10.8333       3746.83       4242.28       9.189       11.032 $0$ 10.8333       3744.38       4243.73       9.1266       11.040 $3$ 10.8333       3740.75       4245.27       9.143       11.040 $3$ 10.8333       3740.75       4245.27       9.143       11.040 $3$ 10.8333       3740.75       4245.27       9.143       11.043 $3$ 0.08       0.04       Rotrage       10.03 % $3$ 0.08       0.04       RSD $203 \%$	Result		100.26	%	Each in	npurity:						Total imp	urities: 0.6	04 %			-					
Related $\Gamma$ RelatedRelated $\Gamma$ Sam RT $\frac{Impurities}{RT}$ $\frac{Impurities}{RT}$ 010.83333746.834242.289.189110.83333746.834243.739.126010.83333740.754243.739.126110.83333740.754245.279.143310.83333740.754245.779.143 $2$ 10.83333740.754245.759.143 $2$ 00.04 $2.43.76$ $Average$ Be0.08O0.04BCOOOOOOOOOOOOOOOOOOOOOOOOOOOOOOOOOOOOO <td <="" colspan="4" th=""><th>Sample</th><th>32 (QC</th><th>S-STG-9</th><th>95004) (</th><th>USP n</th><th>nethod</th><th></th><th></th><th></th><th></th><th></th><th></th><th></th><th></th><th></th><th></th><th></th><th></th></td>	<th>Sample</th> <th>32 (QC</th> <th>S-STG-9</th> <th>95004) (</th> <th>USP n</th> <th>nethod</th> <th></th>				Sample	32 (QC	S-STG-9	95004) (	USP n	nethod												
$ \begin{array}{ c c c c c } \hline \mbox{H} \$	Assay								Related	F												
1       1       1         1       10.8333       3746.83       4242.28       9.189       11.032         1       10.8333       3744.38       4243.73       9.126       11.040         1       10.8333       3740.75       4243.73       9.126       11.040         1       10.8333       3740.75       4245.27       9.143       11.043         e       3743.99       4245.27       9.143       11.043         e       3743.99       4243.76       Average         0.08       0.04       RSD $RSD$	Std RT		Sam RT	r	Std are		Samar	63	Impuri RT	ities	Std RT			Impuriti	es area		Std area	Sa				
010.83333746.834242.289.18911.032010.83333744.384243.739.12611.040310.83333740.754245.279.14311.043310.83333743.994243.768.14311.043ge $2143.99$ 4243.76Average $11.043$ $11.043$ 90.8% $0.04$ $BD$ $BD$ $BD$						ł		}	1					1								
0 $10.8333$ $3744.38$ $4243.73$ $9.126$ $11.040$ 3 $10.8333$ $3740.75$ $4245.27$ $9.143$ $11.043$ $3740.75$ $4245.27$ $9.143$ $11.043$ $2743.99$ $4243.76$ $Average$ $3743.99$ $0.04$ $Average$ $9.08\%$ $0.04$ $BD$	10.8500		10.8333		3746.83	~	4242.28	~	9.189		11.032			333			8149					
3     10.8333     3740.75     4245.27     9.143     11.043       Be       State	10.8500		10.8333		3744.38	~	4243.73	~	9.126		11.040			344			8276					
ge     3743.99     4243.76     Average       0.08     0.04     RSD       99.98%     Each impurity: 0.003 %	10.8333		10.8333		3740.75		4245.27	2	9.143		11.043			334			8265					
0.08         0.04         RSD           99.98%         Each impurity: 0.003 %	Average				3743.99		4243.76		Averag	še				337			8230					
99.98% Each impurity: 0.003 %	RSD				0.08		0.04		RSD					1.80			0.65					
	Result				99.98%				Each ir	mpurity	: 0.003 9	%		Total im	purities:	0.003 %						

## Table 3: Analysis data for sample 2 using in-house and USP method

Assay				Related									
	E S	F7D		Related				Std RT	Impurities area	s area			Std area
TA DIC		old area	Sam area	1	7	3	4	T	1	5	e	4	1
13.27	13.27	618596	672304	12.55	12.93	12.43	16.67	38 ± 1.42	7891	2675	1427	1005	71240
13.27	13.27	625496	677553	12.56	12.93	14.23	16.67	$46 \pm 1.84$	7941	2605	1464	983	71465
13.28	13.26	621567	673514	12.54	12.92	14.24	16.60	54 ± 1.27	7960	2636	1422	066	71824
Average	د <u>ه</u> -	621916	6744573	Average					7919	2638	1420	686	71509
RSD		0.56	0.41	RSD					0.46	1.33	0.5	1.62	0.41
Result		99.79%		Each im	Each impurity: 0.01 %	.01 %			Total imp	Total impurity: 0.02 %	%	_	_
Sampl	Sample 3(QC-STG-95005) (USP me	G-95005) (	(USP meth	thod)									
Assay				Related					1				
La Pto	DT	Ct.J amon		Impurities RT	ies RT			Std RT	unpurtues area	s area			Std area
TV mc		ou area	Sall area	1		7		I	1		7		1
11.02	11.07	4123.84	4638.12	4.475		9.134		11.032	236		448		8149
11.03	11.07	4122.36	4669.38	4.468		9.116		11.040	239		462		8276
11.03	11.07	4116.41	4670.04	4.466		9.134		11.043	234		451		82654
Average	. (1)	4120.87	4659.18	Average					236		454		8230
RSD		0.10	0.39	RSD					1.06		1.62		0.65
Result		99.19%		Each im	Each impurity: 0.004 %	.004 %			Total imp	Total impurities: 0.007 %	07 %		

## Table 4: Analysis data for sample 3 using in-house and USP method

GIP inactivation, they are able to increase the secretion of insulin and suppress the release of glucagon by the alpha cells of the pancreas (Herman et al., 2006 and Herman et al., 2005). Although many companies synthesized the Sitagliptin API and others formulated that as pharmaceutical dosage form but after 11 years from discovery of this molecule a Pharmacopeial analytical method was published for it in USP 39. Prior to the USP method, a full validated analysis method was developed by our research team which is presented in this manuscript. Also, we compared the results of both pharmacopeial and in-house methods when we tested three different batches of sitagliptin API.

#### **Materials and Methods**

#### In-house method validation:

The developed method is presented in Table 1 in comparison with USP monograph. A full analytical method validation based on USP general chapter (<1225>) was done on the developed analysis method. In brief, a calibration curve was plotted successfully with a reasonable R2 equal to 0.9993 and the equation of the curve was Y = 3.4588X+30.099. The range of linearity was 1-1000 ppm. Precision was studied for two different concentrations, 10 and 100 ppm, RSD was 0.62 and 0.44% ( $\leq 2$ ). The accuracy of the method was investigated for the same concentrations as precision and error percentages were 0.55 and 1.2%, which are completely reasonable. Robustness was studied by changing mobile phase flow rate, pH and temperature and the effect of changing these factors on tailing factor were investigated. Limit of Detection (LOD) and Limit of Quantitation (LOQ) of the method were 0.75 and 2.5 ppm, respectively.

## **Comparison of the assay and related substances test results using In-house and USP method:** Three different batches of sitagliptin were

analyzed by both methods and results were compared with each other. Results are described in Table 2-4.

#### **Results and discussion**

The results of analysis of three samples with two methods are presented in Tables 2-4. Results of the assay for all samples are very similar. Generally, in-house method could detect more impurities in most of the samples. Although previous validated HPLC and UV spectroscopy methods were published for analysis of sitagliptin using different analysis conditions (Lavanya et al., 2013, Ravisankar et al., 2015 and Tarkase et al., 2013), there is not any published data in which results were compared to USP monograph.

## Conclusion

An HPLC method was found to be simple, accurate, precise, linear, robust and specific for quantitative estimation of Sitagliptin phosphate in bulk API. Then the correlation between results of this method with newly published USP monograph was investigated. As the column, column temperature, mobile phase composition, sample preparation solvent and detector wavelength are different in two methods, it seems that in-house method could be a valuable alternative for the USP method which could be performed when the user facilities are fit with that.

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