

# Quality-by-Design Based Development and Validation of Stability Indicating Method by UPLC Method for Impurities of Simvastatin from Drug and Pharmaceutical Dosage Form

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Desai *et al.*: QbD based UPLC method development for Simvastatin impurities

A rapid, robust and accurate ultra-high performance liquid chromatographic method was developed and validated for determination of impurities of simvastatin in drug and its pharmaceutical formulation. A systematic Quality-by-design approach was used for method development with the Fusion AE™ software, to screen and optimize the column, mobile phase, column temperature, gradient time and other chromatographic conditions. The optimized method uses Waters Acquity Charged Surface Hybrid, Octadecylsilane C18 (1.7µm×2.1 mm×100 mm) column with gradient elution. Orthophosphoric acid pH was adjusted to 4.5 with triethylamine and acetonitrile with a ratio of (80:20) was used as mobile phase A, whereas methanol:acetonitrile (20:80) as mobile phase B, with a flow rate of 0.35 ml/min. Ultra Violet detector was programmed at 238 nm with a runtime of 13 min, wherein all the impurities were well resolved and were separated from main peak of simvastatin. The method was validated for accuracy, repeatability, reproducibility and robustness. All the validation parameters were in acceptable range. The linearity, limit of detection and limit of quantitation was established for all the impurities and for simvastatin. The method was also applied suitably for determining the degradation products of simvastatin using stress degradation studies

**Key words:** Simvastatin, ultra-performance liquid chromatography, quality-by-design, stress degradation

The conventional approach for analytical method development involves changing one factor at a time (OFAT), commonly known as OFAT technique. The OFAT is very time consuming and lengthy process which generates voluminous data, resulting in a tedious approach. Also, once the method is developed, the method still may need additional efforts when validated<sup>[1]</sup>. The modern pharmaceutical analysis and regulatory scenario, demands to use novel chemometric tools which control many variables simultaneously and helps to provide desired results with minimum experimental trials. This can be achieved by implementing Quality-by-Design (QbD) approach in analytical method development. In International Conference on Harmonization (ICH) guideline Q8(R2) for Pharmaceutical development, QbD is defined as “A systematic approach to development that begins with predefined objectives and emphasizes product and

process understanding and process control, based on sound science and quality risk management”<sup>[1,2]</sup>. The elements of QbD can be extended for analytical method development. The Analytical QbD utilizes statistical modeling and design of experiments (DoE) to arrive at a method operable design region (design space), the robust area, where the developed method provides the desired results<sup>[3]</sup>.

Simvastatin (SMV), chemically designated as -(+)-{1S, 3R, 7S, 8S, 8aR)-1,2,3,7,8,8a-hexahydro-3,7-dimethyl-8-[2-(2R,4R)-tetrahydro-4-hydroxy-6-oxo-2H-pyran-

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2-yl]-1-naphthyl-2, 2-dimethyl butanoate (fig. 1). It is a synthetic lipid lowering agent, which acts by inhibiting 3-hydroxy-3-methyl-glutaryl-coenzyme A (HMG CoA) reductase and is widely used for the treatment of hypercholesterolemia. It has significantly reduced the morbidity and mortality related to the coronary heart disease. SMV is a prodrug and after oral administration it is converted into  $\beta$  hydroxy acid of SMV, which is a potent inhibitor of HMG CoA reductase, a key enzyme used in the biosynthesis of cholesterol in liver<sup>[4]</sup>.

SMV is official in United States Pharmacopoeia (USP), British Pharmacopoeia (BP) and European Pharmacopoeia (Ph. Eur.), whereas SMV Tablets are official in BP and USP. SMV Impurity A, B, C, D, E, F and G (fig. 2) are specified in SMV Tablet monograph in BP as well as in USP. The analytical method followed for related substance test in SMV Tablets in both Pharmacopoeia is high performance liquid chromatography (HPLC) method of analysis<sup>[5-7]</sup>. The literature shows few methods for estimation of SMV along with its impurities are reported in the literature such as HPLC with Ultra-Violet detection<sup>[8-12]</sup>. The degradation behavior studies using ultra-high performance liquid chromatography (UHPLC)/Mass spectrometry (MS) is reported<sup>[1]</sup>. There are various stability indicating methods available for determination of SMV along with the other drugs and its degradation products<sup>[13-17]</sup>. But these methods are either non-specific with respect to Pharmacopoeial impurities or having very long run time.

The analytical method development by using the QbD approach is a recent trend and very few manuscripts are published where the methods are developed using the analytical QbD. Also, various regulatory agencies are requesting that along with the product development, even the analytical methods shall be developed using

the QbD approach. Hence, the present work was aimed at development of a specific, robust and validated method for estimation of SMV and its impurities using QbD approach and DoE principles. SMV is chosen as a model drug because of its great scientific interest due to low solubility and proven instability.

## MATERIALS AND METHODS

SMV and its impurities (SMV Impurity A, B, C, D, E, F and G) were obtained as gift sample from Ipca Laboratories Ltd., Mumbai, India. Acetonitrile (HPLC grade), methanol (HPLC grade), triethylamine (TEA) (for Chromatography) and orthophosphoric acid (OPA) 88 % (Emparta® American Chemical Society (ACS)) was purchased from Merck Life Sciences Pvt. Ltd., Mumbai, India. Milli Q water, obtained from Millipore (Massachusetts, USA), was used in preparation of solutions. Buffers and all other chemicals were of analytical grade.

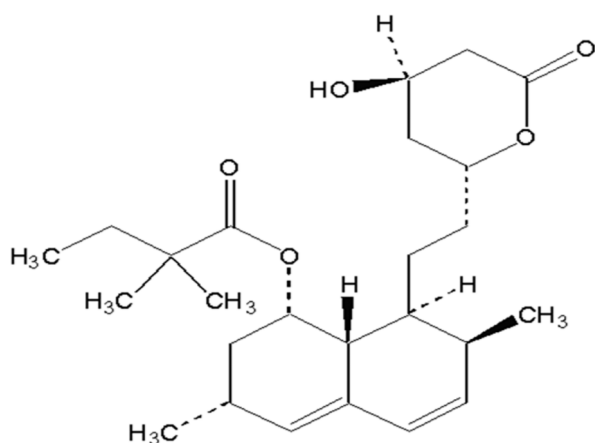
### Equipment:

Experiments were performed on an Acquity ultra performance liquid chromatography (UPLC) H-Class System (Waters Corporation, Milford, Massachusetts) equipped with an autosampler, a quaternary gradient pump, a temperature controlled column compartment, photo-diode array detector and column selection valve. The UPLC system was also equipped with column switch. Instrument control and data acquisition were performed on Empower 2 Chromatography Software (Waters). The method development experiments were designed using Fusion AE™ software (S-Matrix Corporation, Eureka, California (CA)).

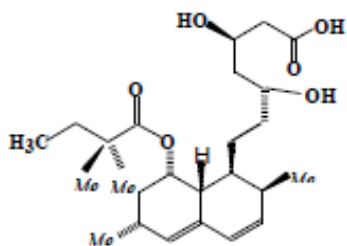
### Method:

The method development studies were carried out in two phases. The first phase consists of the screening experiments wherein columns of different selectivity were used to find the best column which can offer the greatest possible separation of SMV and its impurities. Along with columns, aqueous mobile phases at different pH, organic modifiers, and gradient time were also evaluated using systematic QbD design generated using Fusion AE™ software to obtain best and most robust separation with shortest run time.

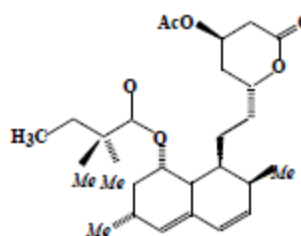
In the second phase the best column, aqueous mobile phase with pH 4.5 and organic solvent type were used for further method optimization to achieve the best chromatographic conditions. These selected parameters were kept constant and other chromatographic



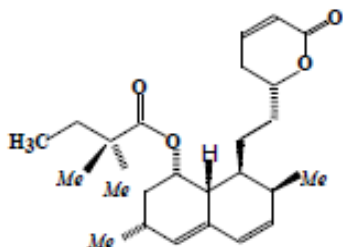
**Fig. 1: Chemical structure of SMV**



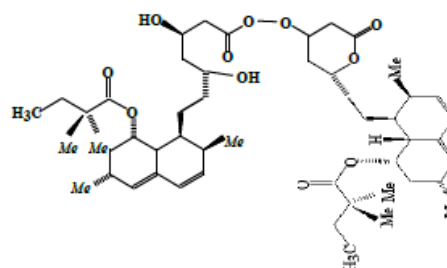
Impurity A



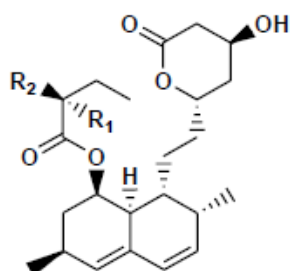
Impurity B



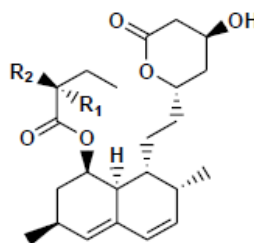
Impurity C



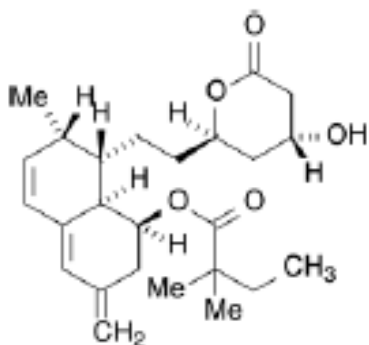
Impurity D

R1 = CH<sub>3</sub>, R2 = H

Impurity E

R1 = H, R2 = CH<sub>3</sub>

Impurity F



Impurity G

Fig. 2: Impurities of SMV

parameters such as flow rate, gradient time and column oven temperature were studied to characterize their response effects and establish method performance region (design space).

**Preparation of buffer and diluent:** To 950 ml of purified water, 5.0 ml of OPA was added. The pH of the solution was adjusted to 4.5 with TEA and volume was made upto 1000 ml with purified. Diluent is prepared

by mixing 400 ml of the above buffer with 600 ml of acetonitrile. The same buffer is used for mobile phase A preparation.

**Preparation of standard solution:** A stock solution was prepared by dissolving accurately weighed 50 mg of SMV to obtain 0.5 mg/ml solution using above mentioned diluent. The stock solution was further diluted with diluent to give 0.5 µg/ml solution.

**Standard preparation:** Tablets powder equivalent to 50 mg of SMV is weighed and transferred to 100 ml volumetric flask. About 70 ml of diluent was added to the flask and sonication was done for about 20 min with intermittent shaking. Flask was cooled to room temperature and then filtered solution was used for analysis.

#### **Method validation:**

Method was validated on the following criteria: specificity, linearity, Limit of detection (LOD) and limit of quantitation (LOQ), precision, accuracy, robustness and stress degradation studies based on ICH guide lines (ICH Q2 (R1)).

**Specificity:** Specificity is the ability to unequivocally assess the analyte in the presence of other potential components, typically including impurities, degradants, matrix, etc. In order to determine the specificity of the method, the following solutions were injected: Diluent, placebo preparation, sample solution and individual impurity solution (Impurity A, B, C, D, E, F and G).

**Linearity:** In order to verify the linearity of the detector, a minimum of six concentration levels of each impurities corresponding to 5 % to 150 % of the specified value was performed. For Impurity A, Impurity C and Impurity D, 0.10, 0.20, 0.50, 1.00, 2.00 and 3.00 µg/ml; for Impurity B and Impurity G, 0.05, 0.10, 0.20, 0.50, 1.00 and 1.50 µg/ml; for Impurity E and F, 0.25, 0.50, 1.00, 2.50, 5.00 and 7.50 µg/ml; whereas, for SMV 0.025, 0.05, 0.10, 0.25, 0.50 and 0.75 µg/ml were the concentration levels prepared and injected for linearity constructions.

**Detection and quantitation limits:** Limit of detection (LOD) and limit of quantitation (LOQ) were obtained from signal to noise ratio. The detection limit was defined as the lowest concentration level resulting in a peak area of three times the baseline noise. The quantitation limit was defined as the lowest concentration level that

provided a peak area with a signal-to-noise ratio higher than 10.

**Accuracy of the method:** The accuracy experiment was performed over the range of 50 %, 100 % and 150 % of the specification level of each impurity. Three samples at each concentration level were prepared and injected.

**System precision:** System Precision was determined from six replicate injections of the SMV standard solution (0.5 µg/ml). The acceptance criterion was  $\pm 2$  % of coefficient of variation (% CV) for retention times and peak areas for both drugs.

**Method precision (Repeatability):** The repeatability of the method was evaluated by the analysis of impurities in the sample solution. Six sample preparations were performed and each one was injected once.

**Intermediate precision:** In order to evaluate the intermediate precision of the UPLC method, related compounds test, was analyzed on a different day, using a different preparation of mobile phase, a different UPLC system, a different column as well as a different analyst performing the analysis. The same batch of the tablets was used in this study for a comparison purpose.

**Stress degradation studies:** Stress testing of the drug product can help to identify the likely degradation products and specificity of the analytical procedures. The stress degradation of SMV tablets sample was conducted for hydrolytic degradation (acid and base treated) and oxidative degradation. For acid hydrolysis, 5 ml of 0.5 Molar (M) hydrochloric acid solution was added to the sample and flask was kept in water bath at 60° for 30 min, whereas for base hydrolysis, 5 ml of 0.1 M hydrochloric acid solution was added to the sample and it was kept in at room temperature for 10 min. After treatment period the samples were suitably neutralized with corresponding base or acid and further sample preparation was followed as given above. For oxidative degradation, 5 ml of 1 % Hydrogen peroxide (H<sub>2</sub>O<sub>2</sub>) solution was added to the sample and followed as above.

## **RESULTS AND DISCUSSION**

Achieving best separation with fastest possible runtime with highest possible sensitivity was our goal during the method development. To achieve best separation among the impurities and SMV, choosing the right column

with optimum selectivity is a key during development. After selection of right column, mobile phase pH and composition must be optimized to obtain the desired separation with shortest runtime without compromising selectivity and sensitivity of the method. The systematic approach of maximizing selectivity without compromising sensitivity is completely missing in the current literature for the development. The systematic approach of QbD uses more statistical concepts with experimental design plans (also referred as DoE) as an efficient and fast tool for method development which is achieved using Fusion AE™ software

In the first phase of method development, screening experiments were performed using the “Method screening” tool of Fusion AE™ software (S-Matrix Corporation, Eureka, CA). The screening of different chemistry for columns, selection of mobile phase pH,

selection of organic modifiers and gradient composition was studied using Method screening tool. The details of method variables are tabulated in Table 1.

Total 46 experiments were designed statistically using DoE principles. Few repeat trials were included in these designed experiments for statistical reason. Table 2 represents the experimental design used in the column and eluent combination scouting phase. The designed experiments were then exported to the chromatographic data software (Empower 2) as ready to run methods and sequence. This ready sequence was executed and data was generated in Empower 2 software on UPLC system. After completion of the sequence, the results were processed and selection of column, mobile phase pH, organic modifier and gradient time was done with predetermined goal using the “Numerical search tool” in Fusion AE™ software. Not less than 6 peaks and not

**TABLE 1: QbD METHOD VARIABLES**

Study variables	Range or levels
Column type	Acquity UPLC Bridged Ethyl Hybrid (BEH) Amide, 100×2.1 mm, 1.7 μm Acquity UPLC CSH C18, 100×2.1 mm, 1.7 μm Acquity UPLC High Strength Silica (HSS), 100×2.1 mm, 1.8 μm Acquity UPLC CSH Phenyl-Hexyl, 100×2.1 mm, 1.7 μm
pH of mobile phase	3.00, 4.50, 5.00, 7.40 (Buffers prepared with 5 ml OPA in 1000 ml water and pH adjusted with TEA)
Strong solvent/Organic modifier	Acetonitrile, Methanol
Gradient time (Min)	5.0 ≤ Gradient time ≤ 12.0

**TABLE 2: DESIGN OF EXPERIMENTS FOR SCREENING**

Run No.	Solvent	Gradient Time	pH	Column	Run No.	Solvent	Gradient Time	pH	Column
1	Acetonitrile	12	3.00	BEH Amide	24	Acetonitrile	12	7.40	BEH Amide
2	Acetonitrile	12	3.00	CSH C18	25	Acetonitrile	5	7.40	BEH Amide
3	Acetonitrile	5	3.00	CSH C18	26	Methanol	12	3.00	CSH Phenyl-Hexyl
4	Acetonitrile	5	3.00	CSH Phenyl-Hexyl	27	Methanol	12	3.00	HSS C18
5	Acetonitrile	5	3.00	CSH C18	28	Methanol	5	3.00	HSS C18
6	Acetonitrile	10.3	4.50	HSS C18	29	Methanol	5	3.00	BEH Amide
7	Acetonitrile	6.8	4.50	BEH Amide	30	Methanol	10.3	4.50	BEH Amide
8	Acetonitrile	8.5	4.50	CSH C18	31	Methanol	8.5	5.00	CSH Phenyl-Hexyl
9	Acetonitrile	8.5	4.50	CSH Phenyl-Hexyl	32	Methanol	8.5	5.00	CSH C18
10	Acetonitrile	8.5	5.00	CSH C18	33	Methanol	8.5	5.00	BEH Amide
11	Acetonitrile	5	5.00	HSS C18	34	Methanol	8.5	5.00	HSS C18
12	Acetonitrile	12	5.00	CSH Phenyl-Hexyl	35	Methanol	8.5	5.00	HSS C18
13	Acetonitrile	8.5	5.00	BEH Amide	36	Methanol	8.5	5.00	CSH Phenyl-Hexyl
14	Acetonitrile	8.5	5.00	CSH C18	37	Methanol	8.5	5.00	CSH C18
15	Acetonitrile	8.5	5.00	HSS C18	38	Methanol	10.3	5.00	CSH C18
16	Acetonitrile	8.5	5.00	CSH Phenyl-Hexyl	39	Methanol	5.0	5.00	CSH Phenyl-Hexyl
17	Acetonitrile	8.5	5.00	HSS C18	40	Methanol	8.5	5.00	BEH Amide
18	Acetonitrile	8.5	5.00	CSH Phenyl-Hexyl	41	Methanol	12	7.40	CSH C18
19	Acetonitrile	8.5	5.00	BEH Amide	42	Methanol	12	7.40	CSH Phenyl-Hexyl
20	Acetonitrile	12	7.40	HSS C18	43	Methanol	5	7.40	CSH C18
21	Acetonitrile	5	7.40	CSH Phenyl-Hexyl	44	Methanol	12	7.40	CSH Phenyl-Hexyl
22	Acetonitrile	8.5	7.40	CSH C18	45	Methanol	5	7.40	BEH Amide
23	Acetonitrile	12	7.40	HSS C18	46	Methanol	5	7.40	HSS C18

more than 10 min for the last peak in the sequence were set as goal parameters. The numerical search resulted in 5 best possible answers out of which first answer with 0.96 probabilities (Table 3) was selected as model run for optimization. The optimum run gave pH 4.5 (OPA-TEA) as buffer for mobile phase with acetonitrile as organic modifier on Acquity UPLC Charged Surface Hybrid (CSH) C18, 100×2.1 mm, 1.7 μm.

Few trials were taken for mobile phase composition optimization, which includes addition of 20 % acetonitrile in buffer to avoid precipitation and inclusion of 20 % methanol in acetonitrile for obtaining better polarity for separation.

The second phase optimization of method was done using “Method Optimization” tool of Fusion AE software. In the optimization experiments, 17 DoE experiments were generated (Table 4) and effects of three study parameters; Pump Flow rate (0.35 0.40 and 0.45 ml/min), Gradient time (6.0, 7.5 and 9.0 min) and column temperature (30, 35 and 40°); were simultaneously evaluated to assess the effects of these parameters on each of the five response variables. The five response variables are listed in Table 5. Column type, injection volume, mobile phase buffer (pH 4.5 TEA/OPA buffer:methanol 80:20) and organic modifier (acetonitrile:methanol 80:20) were kept constant. The designed optimization sequence was run on chromatographic data software that is Empower 2. After completion of sequence the chromatographic runs were processed and the results were imported in Fusion AE™ for further statistical evaluation.

This chromatographic data was processed with trend response in capabilities in Fusion AE™ and prediction equations (models) were generated automatically. These prediction equations quantitatively characterize the independent and interactive effects of the study parameters on each critical quality attributes (variables)

included in the study. These equations then used to predict the method performance region (Design space) within the experimental region. The performance region generated for estimation of SMV and its impurities is given in fig. 3. The unshaded region represents the design space where all the performance parameters of the method are successfully met. Based on these results, the final chromatographic conditions are given in Table 6. Fig. 4 provides three dimensional plots showing the combined effect of variables such as gradient flow and pump flow rate on no of peaks and USP resolution of  $\geq 1.5$  between all peaks. As we can see that when the gradient time and flow rate is increased there is negative relationship for the number of peaks obtained. The representative chromatogram of sample solution in the optimized method is given as fig. 5.

The developed method was then validated as per ICH Q2 (R2) guidelines. The validation results are discussed in details in following sections. The method specificity was proved by injecting the placebo solution, each individual impurity and standard solution and sample solution. All the impurities and the placebo peaks were very well separated from the main peak of SMV and all the impurities were well separated from each other (fig. 5). In addition, the peaks of impurities and SMV were found to be pure by peak purity analysis using Empower 2 software.

The calibration curves constructed for SMV and its impurities (A, B, C, D, E, F and G) were evaluated by its correlation coefficient. The peak area was found linear in the range of 5 % to 150 % of the specification levels as given in Table 7. The correlation coefficients for all the calibration plots of SMV and its impurities were more than 0.999.

The LOD and LOQ values were obtained from the linearity curve as per ICH guidelines and values are

**TABLE 3: NUMERICAL ANSWER SEARCH**

Variable Settings					
Variable	Level Setting				
Strong Solvent Type	Acetonitrile				
Gradient Type	12.00				
pH	4.50				
Column Type	CSH C18 10 Column				
Predicted results					
Response	Goal	Predicted result	Desirability	-2 Sigma Configuration limit	+2 Sigma Configuration limit
No. of Peaks	Maximize	8.3	1.0000	7.2	10.0
Last Peak-Retention Time	Minimize	9.09	1.0000	6.43	12.91

Cumulative desirability target=1.0000, Cumulative desirability result=1.0000

represented in Table 7. The obtained LOQ values were found to be less than 20 % of the specification level.

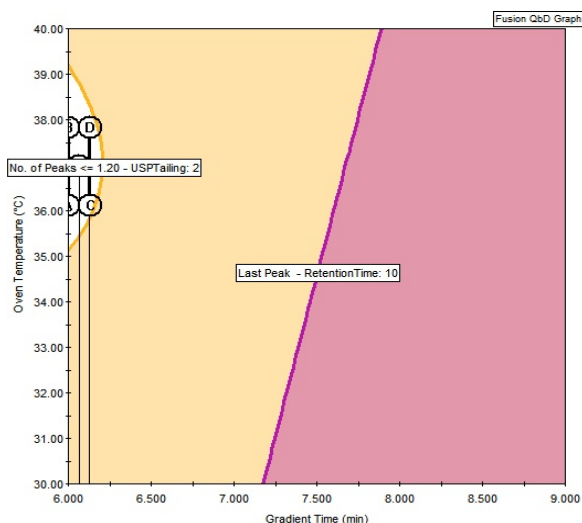
Accuracy for SMV and its impurities was determined by spiking known amount of analyte into formulation placebo at 50 %, 100 % and 150 % levels. Triplicate spikes were performed for all levels. The results are summarized in Table 8 and met the acceptance criteria.

**TABLE 4: DESIGN OF EXPERIMENTS FOR OPTIMIZATION**

Run No.	Pump flow rate (ml/min)	Gradient time (min)	Oven temp (°)
1	0.350	6.0	30.0
2	0.450	6.0	30.0
3	0.350	9.0	30.0
4	0.450	9.0	30.0
5	0.400	7.5	30.0
6	0.450	7.5	35.0
7	0.400	6.0	35.0
8	0.400	9.0	35.0
9	0.350	7.5	35.0
10	0.400	7.5	35.0
11	0.400	7.5	35.0
12	0.400	7.5	35.0
13	0.350	6.0	40.0
14	0.400	7.5	40.0
15	0.450	6.0	40.0
16	0.350	9.0	40.0
17	0.450	9.0	40.0

The coefficient of variation of six replicate injections of the standard solution was found to be 0.37 % which is well within the acceptance criteria. The coefficient of variation of peak area was within the range of 2 % limit signifying suitability of the system.

The six sample preparations showed the coefficient of variance within the acceptance criteria of less than 10 % for all the impurities indicating that the method



**Fig. 3: Method performance region.**

The unshaded region represents the design space where all the performance parameters of the method are successfully met

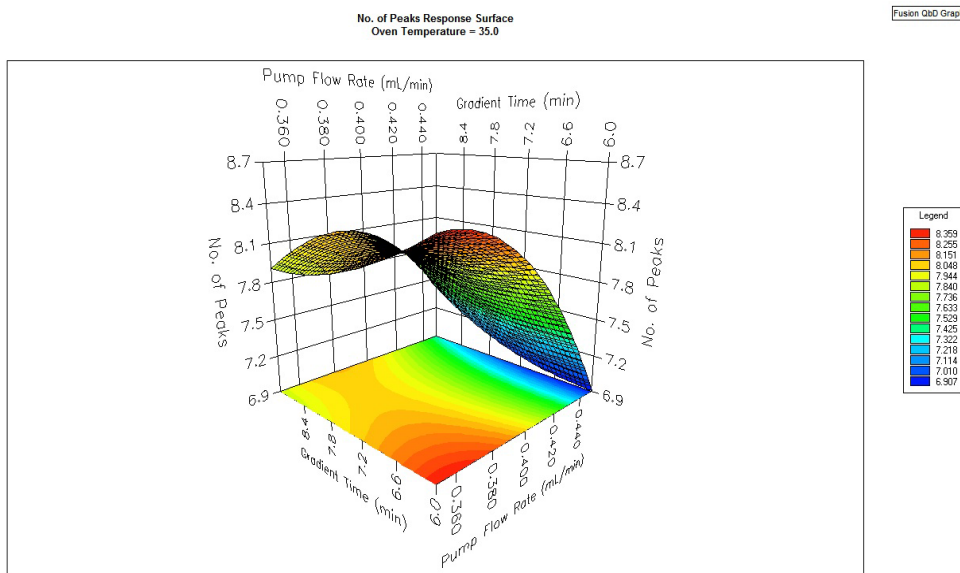
**TABLE 5: RESPONSE GOAL SETTINGS**

Response	Target	Lower bound	Upper bound	Relative rank
No. of peaks	Maximize	6.0	9.0	1
No. of peaks $\geq$ 1.50-USP resolution	Maximize	1.0	2.0	1
No. of peaks $\geq$ 2.00-USP resolution	Maximize	2.0	9.0	1
No. of Peaks $\leq$ 1.20-USP tailing	Minimize	1	9.0	1
Max Peak #1-USP resolution	Maximize	1.00	2.00	1
Last Peak-Retention time	Minimize	7.00	10.00	1

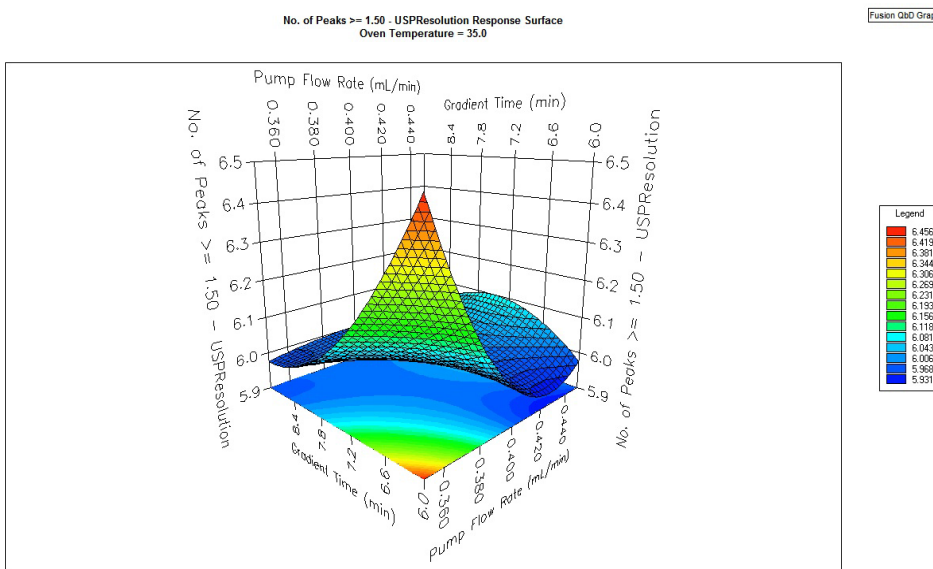
**TABLE 6: OPTIMIZED CHROMATOGRAPHIC CONDITIONS FOR ANALYSIS OF SMV AND ITS IMPURITIES**

Parameters	Condition		
Column	Waters Acquity CSH C18 1.7 $\mu$ m, 2.1 mm $\times$ 100 mm		
Flow rate	0.35 ml/min		
Column temperature	35°		
Sampler temperature	10°		
Injection volume	2 $\mu$ l		
Detector wavelength	UV, 238 nm		
Run time	13 min		
Diluent	Buffer:Acetonitrile (40:60 v/v)		
	Time	Mobile phase A	Mobile phase B
	0	50	50
Gradient Program	6.2	5	95
	8.7	5	95
	9.0	50	50
	13.0	50	50

Where, mobile phase A is buffer:acetonitrile (80:20) and mobile phase B is acetonitrile:methanol (80:20)

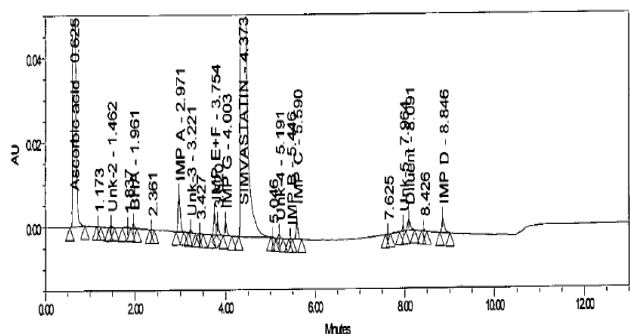


a)



b)

**Fig. 4: Three dimensional response surface plots.**  
 (a) Correlation of No. peaks, (b) Correlation of USP resolution  $\geq 1.5$



**Fig. 5: Representative sample chromatogram of the optimized method**  
 All impurities of SMV are well separated from main peak and from each other

is repeatable. The results are summarized in Table 8. The intermediate precision established by analyzing six sample on different day by different analyst, who has used different instrument and the chromatographic preparations, resulted in the similar results which are tabulated in Table 8.

The results of stress degradation studies given in Table 9, which indicates that sample is degraded in acidic condition and Impurity A is formed majorly at a concentration of 23 %. The peak purity threshold was found to be more than the purity angle for all the impurity peaks which indicates that that peaks of all



**TABLE 7: METHOD SPECIFICITY, LINEARITY, LOD AND LOQ**

Component	RT	RRT	Equation	Correlation Coefficient	Limit of Quantification (%)	Limit of Detection (%)
SMV	4.389	1.00	y=20166x-89.725	1.0000	0.003	0.001
Impurity A	2.887	0.66	y=7413x-143.39	0.9999	0.034	0.011
Impurity B	5.468	1.25	y=16235x-83.168	0.9999	0.012	0.004
Impurity C	5.611	1.28	y=17118x-234.68	0.9998	0.033	0.011
Impurity D	8.836	2.01	y=9386.1x-710.72	0.9991	0.079	0.028
Impurity E	3.754	0.85	y=20206x-365.72	0.9999	0.054	0.018
Impurity F	3.820	0.87	y=20377x-461.93	0.9999	0.067	0.022
Impurity G	4.020	0.92	y=18123x-83.198	0.9999	0.011	0.003

**TABLE 8: ACCURACY AND PRECISION DATA**

Component name	Mean recovery			Method precision		Intermediate precision	
	50 %	100 %	150 %	Analyst I	% RSD	Analyst II	% RSD
SMV	100.1	99.4	101.8	-	-	-	-
Impurity A	102.7	104.9	103	0.319	0.86	0.322	1.75
Impurity B	104.3	102	98.3	0.277	0.19	0.276	0.76
Impurity C	104.8	106.2	106.1	0.206	0.91	0.206	0.91
Impurity D	99.3	96.8	98.2	0.660	0.10	0.664	1.95
Impurity E	102.3	102.1	102.5	0.391	0.70	0.394	1.78
Impurity F	101.5	102.2	102.9	0.260	1.02	0.263	1.97
Impurity G	101.4	101.8	98.3	0.302	0.44	0.302	0.36
Unknown Max	-	-	-	0.048	1.56	0.048	1.07
Total Impurity	-	-	-	2.540	0.75	2.547	1.15

**TABLE 9: STRESS DEGRADATION STUDIES**

Condition	% Impurity			
	As such	Acid	Base	Peroxide
Impurity A	0.455	23.26	0.565	0.451
Impurity B	0.007	0.007	0.008	0.007
Impurity C	0.365	0.379	0.344	0.345
Impurity D	0.422	0.356	0.419	0.42
Impurity E	0.399	0.334	0.396	0.398
Impurity F	0.263	0.259	0.273	0.264
Impurity G	0.116	0.097	0.114	0.115
Unknown Max	0.038	0.042	0.038	0.038
Total	2.329	26.985	2.409	2.501

the known impurities were found to be pure. All the mass balance study showed results of more than 99 % indicate that a good correlation was observed between generation of impurities in the degradation sample and the assay value of the sample. Hence it can be confirmed that no other peaks were obtained during stress degradation studies except for Impurity A which is a known degradation product of SMV.

In conclusion, a simple and rapid reversed-phase UPLC method has been developed systematically for estimation of SMV impurities using Fusion AE™ method development software. Response surface plots illustrated the major effects of gradient time flow rate and oven temperature on the separation. This method

has been successfully validated for determination of SMV impurities from drug formulation. The QbD approach of method development is recommended by Food and Drug Administration (FDA) and provides a very clear picture of the robustness during the method development process. Unlike OFAT approach, it is a faster way of developing method, which helps choosing much better method condition during the development process through design space. The validation results confirmed the usefulness of the method. The method was found to be precise, accurate and linear, whereas the stress degradation studies identified the potential degradation products which can form during the shelf life of the product. This newly developed method is

having run time of about 13 min which is almost 5-8 times shorter than the Pharmacopoeial and literature methods.

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