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# Multivariate optimization of liquid chromatographic conditions for determination of dapagliflozin and saxagliptin, application to an in vitro dissolution and stability studies



Sunitha Gurrala<sup>1,2</sup>, Shiva Rai<sup>3\*</sup>, C. V. S. Subrahmanyam<sup>2</sup> and Panikumar D. Anumolu<sup>2</sup>

### Abstract

**Background:** Analytical quality by design driven HPLC method has been optimized for simultaneous estimation of dapagliflozin and saxagliptin in pharmaceutical dosage form. Response surface methodology was employed for optimization of experimental conditions using three factors such as organic phase (%), pH of aqueous phase, and flow rate of mobile phase.

**Results:** Virtuous separation of analytes was achieved with mobile phase consisted of acetonitrile: phosphate buffer, pH 5.8 (26:74% v/v) with flow rate 0.96 mL/min using SPOLAR C18 column (250  $\times$  4.6 mm, 5  $\mu$ ) with run time 6 min and UV detection at 236 nm. Retention time for dapagliflozin and saxagliptin were found to be 3.5 and 5.0 min, respectively. Method was validated as per ICH guidelines. The plot between peak area vs concentration for dapagliflozin and saxagliptin were rectilinear in the range of 0.2-300  $\mu$ g/mL and 0.1-150  $\mu$ g/mL respectively with detection and quantification limits were 0.061 and 0.18  $\mu$ g/mL for dapagliflozin and 0.014 and 0.043  $\mu$ g/mL for saxagliptin, respectively.

**Conclusion:** The proposed method was exploited for assay, in vitro dissolution, and stability studies of target drugs in marketed dosage form.

Keywords: Central composite design, Dapagliflozin, Saxagliptin, Dissolution, Stability

### **Background**

Dapagliflozin and saxagliptin are hypoglycemic agents used in the treatment of type II diabetes mellitus as a fixed-dose combination, and their tablets are available on the market under the FDA-approved brand name QTERN° (2017). Dapagliflozin is an inhibitor of sodium glucose co-transporter-2, which is responsible for glucose reabsorption from the glomerular filtrate in the kidney. Saxagliptin is a competitive and reversible

dipeptidyl peptidase-4 enzyme inhibitor that decreases glucagon-like peptide insulinotropic hormone-1 breakdown for improved glycemic regulation in patients with diabetes [1].

Analytical methods with HPLC technique are developed to support drug testing during production and quality release operations as well as during long-term stability studies against requirements. Due to numerous factors such as restricted availability of chromatographic columns, solvents and chemicals and critical physicochemical properties of analytes such as solubility, pKa, and stability, one factor at a time (OFAT) based analytical methods (trial and error based on one variable)

Full list of author information is available at the end of the article



<sup>\*</sup> Correspondence: srajkavadi@gmail.com

<sup>&</sup>lt;sup>3</sup>Department of Chemistry, University College of Science, Osmania University, Hyderabad 500007, India

experience several difficulties in optimizing robust chromatographic conditions. The model of analytical quality by design (AQbD) is a preferred and recommended approach to be adopted in the development of analytical methods to achieve regulatory versatility [2–7]. AQbD approach in chromatographic method development explores greater understanding of the parameters that must be regulated and tracked during the method's life cycle.

Literature survey revealed that there were few reports published on the simultaneous quantification of dapagliflozin and saxagliptin using UV [8], HPTLC [9], HPLC [10-18], and LC-MS [19, 20] techniques in bulk drug and pharmaceutical dosage forms. Few methods were reported for analysis of dapagliflozin and saxagliptin separately [21-24] and in combination with other drugs [25-29]. Literature methods concerted largely on the development and validation of a RP-HPLC methods using different organic phases at different wavelengths and sensitive spectroscopic methods. The literature HPLC methods were based on OFAT variation conditions, which were time consuming and may not have scientific evidence during development stage. Hence, present project has undertaken with the quality-by-design approach in robust HPLC method development for simultaneous analysis of dapagliflozin and saxagliptin in API and marketed dosage form.

### **Methods**

### Instrumentation

The HPLC system (Shimadzu Corporation, Japan) was connected to the LC solution software, consisting of binary pump (LC-20AD), Rheodyne syringe sample injector (20  $\mu\text{L}$ ), and an UV detector (SPD-20A). Type-II (paddle) dissolution apparatus (Electro lab TDT-08L), ultra-sonicator (RK 106, Spincotech), millipore (0.45  $\mu\text{m}$ ) filters, and digital pH meter (LI-120, Elico) were utilized for this work. Design-Expert version 11.0.5.0 software (Stat-Ease Inc. Minneapolis) was employed for designing of LC experiments and response-modeling to generate design space for optimized robust analytical method.

### Chemicals and reagents

The reference standards of saxagliptin and dapagliflozin were obtained from Hetero Laboratories, Hyderabad, India. Marketed formulation (QTERN) with label claim 10 mg of dapagliflozin and 5 mg of saxagliptin per tablet was procured from local pharmacy. All solvents used for mobile phase were of HPLC grade and obtained from Merck, Mumbai, India.

### Chromatographic conditions

Chromatographic separation of analytes was achieved on SPOLAR C18 (250  $\times$  4.6 mm, 5 $\mu$ ) column at ambient temperature with isocratic elution. The mobile phase consisting of acetonitrile: phosphate buffer, pH 5.8 (26: 74% v/v) at flow rate of 0.96 mL/min. Injection volume was 20  $\mu$ L and UV detection at 236 nm.

### Preparation of standard solution

In order to prepare binary standard solution, dapagliflozin (20 mg), and saxagliptin (10 mg) were weighed, decamped into volumetric flask of 10 mL and dissolved in diluent by sonication. Volume was contrived up to the mark with diluent solution and the flask was shaken well (solution A). Further 0.1 mL of this solution was diluted to 10 mL with diluent so as to get binary working standard solution concentration of 20  $\mu$ g/mL and 10  $\mu$ g/mL for dapagliflozin and saxagliptin, respectively.

### Method development using AQbD approach

Method optimization was premeditated with 20 experimental runs under (8 factorial points, 6 center points, and 6 axial points with  $\alpha$ = 0.6073) central composite design (CCD) to study the interactions of the three critical method parameters (CMPs), namely [S1], % acetonitrile  $(X_1)$ , aqueous phase pH  $(X_2)$ , and flow rate  $(X_3)$ . The chromatographic responses (resolution, capacity factor) were obtained through experimentation on HPLC instrument using working standard solution of dapagliflozin (20 μg/mL) and saxagliptin (10 μg/mL). Statistical analysis was performed together with CCD in the Design-Expert software. The significance of factors was calculated using analysis of variance (ANOVA). Design space was generated by numerical optimization where Derringer's desirability function was used to attain high method performance criteria.

### Method validation

The validation parameters like precision, linearity, accuracy, system suitability, sensitivity, and robustness were premeditated in accordance to ICH Q2(R1) guidelines.

### Linearity

Aliquots of 0.002, 0.5, 1.0, 1.5, 2.0, 2.5, and 3.0 mL were withdrawn from mixed standard stock and diluted to 10 mL with mobile phase such that the final concentration of dapagliflozin and saxagliptin and in the range of 0.2-300  $\mu$ g/mL and 0.1-150  $\mu$ g/mL was obtained respectively. Calibration curve was generated by plotting the peak area against the concentration of drug.

### System suitability

The efficiency of optimized method was monitored by system suitability test. It was carried out via injection of freshly prepared working standard solution into HPLC under optimized conditions for six times. The chromatographic responses (analytical attributes) studied were retention time, resolution, capacity factor, theoretical plate number, peak area, selectivity factor, and tailing factor of analyte peaks.

### Precision

Repeatability of proposed method was verified by analyzing 6 replicate injections of freshly prepared working standard solution of dapagliflozin (100  $\mu g/mL$ ) and saxagliptin (50  $\mu g/mL$ ) in mobile phase on the same day. Intermediate precision was performed by analyzing replicates of same concentration solution prepared in three consecutive days. The peak area of the analytes was determined and %RSD was calculated.

### Accuracy

Recovery of standard that spiked to target concentration of sample at three levels (80, 100, 120%) was studied for method accuracy. The % recovery of drug was calculated by measurement of its peak area in the chromatogram.

### Specificity

Specificity of the method was established by comparing the chromatogram of blank (mobile phase), placebo solution, and matrix (degradants) with test solution (analytes in mobile phase). The placebo solution comprises all the commonly used excipients for manufacturing of tablet dosage form. Stress degraded samples were prepared as mentioned in the forced degradation studies where the drugs were exposed to various stress conditions such as acid, base, peroxide, heat, and light to generate their degradation products.

### Sensitivity

Limit of detection (LOD) and limit of quantification (LOQ) were premeditated from the regression analysis data of linearity studies (LOD =  $3.3~\sigma/s$ , LOQ =  $10~\sigma/s$ ; where  $\sigma$ , standard deviation of response; s, slope of calibration curve).

### Robustness

In order to assess the robustness of the method, deliberate changes in method, critical parameters were made within the design space. The variations in the parameters include pH ( $\pm$  0.1), organic phase ( $\pm$  1 %), and flow rate ( $\pm$  0.04 mL/min) of mobile phase. The % RSD of theoretical plate number and retention time of chromatogram obtained for every variation was calculated.

### Assay of marketed formulation

Twenty tablets of marketed formulation (QTERN), each containing 10 mg of dapagliflozin and 5 mg of

saxagliptin were taken, average weight was measured and the fine powder was crushed. An accurately weighed amount of powder equal to 10 mg of dapagliflozin and 5 mg of saxagliptin was transferred to volumetric flask of 10 mL capacity containing methanol and sonicated for 15 min. The flask was shaken and volume was produced with methanol till the mark and filtered via Whatman filter paper (no: 41). From the filtrate, 0.2 mL was transferred into 10 mL volumetric flask and the volume was leveled to the mark with mobile phase. The amount of saxagliptin and dapagliflozin present in sample solution was determined.

### **Dissolution studies**

Dissolution testing of marketed formulation was carried out in FDA-recommended dissolution media, i.e., acetate buffer pH 4.5 (1000 mL) for dapagliflozin and 0.1 N HCl with pH 1.2 (900 mL) for saxagliptin using paddle apparatus (USP apparatus 2) at 50 rpm and 37 ± 0.5 °C for 45 min. Sampling aliquots of 5 mL were withdrawn at 10, 15, 20, 25, 30, and 45 min interval and replacing the fresh medium with an equal amount. After the end of each test time, sample aliquots were filtered. Filtrate of 0.1 mL was diluted 10 mL with mobile phase (phosphate buffer pH 5.8: acetonitrile, 74:26) and analyzed by the contemplated RP-HPLC method. Amount of drug dissolved (saxagliptin/dapagliflozin) was calculated using their respective calibration curves. The cumulative percentage of drug dissolved was plotted against the time.

### Forced degradation studies

Stress studies were executed on saxagliptin and dapagliflozin standards under acid, base, oxidative, thermal, and UV light conditions. Acid degradation was carried out with 5 mL of mixed standard stock solution, to this 5 mL of 1 N hydrochloric acid was added and kept at 60 °C for 60 min, then neutralized with 1 N sodium hydroxide. From this, 0.2 mL was diluted to 10 mL with mobile phase and injected into the HPLC system. Similarly, 1 N sodium hydroxide for base degradation and hydrogen peroxide (6%) for oxidative conditions were used. Dry heat degradation (105 °C, 6 h) and light degradation (UV Chamber, 200-Watt h/m<sup>2</sup>, 48 h) were performed on selected drugs in solid state. After degradation, the solid samples (20 mg dapagliflozin/saxagliptin 10 mg) were dissolved in 10 mL methanol and 0.1 mL of this solution was diluted to 10 mL with mobile phase. These solutions were injected into the HPLC system and sample stability assessment chromatograms were documented.

# **Results**Analytical method development Selection of detection wavelength

The individual standard solutions of dapagliflozin (20  $\mu$ g/mL) and saxagliptin (10  $\mu$ g/mL) were prepared in mobile phase and scanned in UV spectrophotometer [S3]. The overlaid UV spectra showed an iso-absorptive point at 236 nm, which was selected as detection wavelength in PDA detector.

### Design of experiments

Design of experiments (DoE) tool of RSM was employed to optimize the critical method parameters (CMP). Central composite design runs [S4] were executed on HPLC system with respect to three variables such as % acetonitrile  $(X_1)$ , aqueous phase pH  $(X_2)$ , and flow rate  $(X_3)$  to acquire resolution (Rs), capacity factor  $(K_1', K_2')$  as method control responses. CCD plan and experimental results were illustrated in Table 1. The data generated was analyzed using the statistical software.

### Statistical analysis

Statistical analysis of experimental observations was performed to evaluate significant factors that affect the chromatographic response, given in Table 2. Based on ANOVA results, it was found that the process model

**Table 2** ANOVA and regression analysis of selected models

	,		
Response	Rs	K <sub>1</sub> ′	K <sub>2</sub> '
Sum of squares	106.99	101.41	127.96
Df	9	9	9
Mean square	11.89	11.27	14.22
F value	130.4	151.5	44.5
p value	< 0.0001	< 0.0001	< 0.0001
Std. dev.	0.30	0.27	0.56
Mean	3.92	4.76	6.06
% CV	7.70	5.73	9.33
r <sup>2</sup>	0.996	0.993	0.978
Adjusted R <sup>2</sup>	0.984	0.986	0.954
Predicted R <sup>2</sup>	0.937	0.947	0.811
Adequate precision	41.8	46.5	25.9
PRESS	6.85	5.39	24.76

Rs resolution,  $K_1$ ' capacity factor of dapagliflozin,  $K_2$ 'capacity factor of saxagliptin

with  $X_1$ ,  $X_2$ , and  $X_3$  along with interactions is highly significant (p<0.05) at 95% confidence level with curvature effect. The following second-order polynomial (quadratic) equations were generated consistently with models for each critical quality attribute.

**Table 1** Central composite design and experimental results

Run Type		<i>X</i> <sub>1</sub>	<i>X</i> <sub>2</sub>	Х <sub>3</sub>	Y <sub>1</sub>	Υ <sub>2</sub>	Υ <sub>3</sub>
		Org. phase (%)	рН	Flow rate (mL/min)	Rs	<i>K</i> <sub>1</sub> ′	K <sub>2</sub> '
1	Factor	35	6	1	1.8	1.57	2.36
2	Center	30	5	0.8	3.64	5.11	6.3
3	Factor	25	6	1	10.5	1.59	3.07
4	Factor	25	4	1	8.6	7.8	8.2
5	Factor	35	6	0.6	6.2	4.53	4.9
6	Center	30	5	0.8	3.61	5.12	6.35
7	Center	30	5	0.8	3.54	5.18	6.31
8	Axial	38.5	5	0.8	1.4	1.2	2.3
9	Factor	35	4	1	1.4	2.33	2.59
10	Center	30	5	0.8	3.57	5.06	6.21
11	Factor	25	4	0.6	11.1	10.1	11.3
12	Center	30	5	0.8	3.42	5.3	6.38
13	Factor	35	4	0.6	1.8	3.2	4.6
14	Axial	30	3.31	0.8	6.4	6.6	9.3
15	Center	30	5	0.8	3.57	5	6.29
16	Axial	30	5	1.14	1.8	2.74	3.17
17	Axial	30	6.68	0.8	8.7	3.2	5
18	Axial	30	5	0.46	10.8	7.9	10.2
19	Factor	25	6	0.6	11.1	5.4	8.5
20	Axial	21.6	5	0.8	1.0	6.2	7.9

 $X_1$ ,  $X_2$ ,  $X_3$  factors;  $Y_1$ ,  $Y_2$ ,  $Y_3$  responses; Rs resolution;  $K_1$  capacity factor of dapagliflozin;  $K_2$  capacity factor of saxagliptin

$$Y_{1} (Rs) = 3.61 - 1.57X_{1}$$

$$+ 1.54X_{2} - 0.72X_{3} - 1.65X_{1}X_{2} - 0.072X_{1}X_{3}$$

$$+ 0.22X_{2}X_{3} - 0.232X_{1}^{2}$$

$$+ 0.829X_{2}^{2} - 0.143X_{3}^{2}$$

$$Y_{2} (K_{1}') = 5.13 - 1.59X_{1} - 1.18X_{2} - 1.36X_{3}$$

$$+ 1.44X_{1}X_{2} + 0.285X_{1}X_{3} - 0.45X_{2}X_{3}$$

$$- 0.515X_{1}^{2} - 0.09X_{2}^{2} + 0.06X_{3}^{2}$$

$$Y_{3} (K_{2}') = 6.33 - 1.91X_{1} - 1.11X_{2} - 1.82X_{3}$$

$$+ 1.0X_{1}X_{2}$$

$$+ 0.498X_{1}X_{3} - 0.36X_{2}X_{3} - 0.56X_{1}^{2}$$

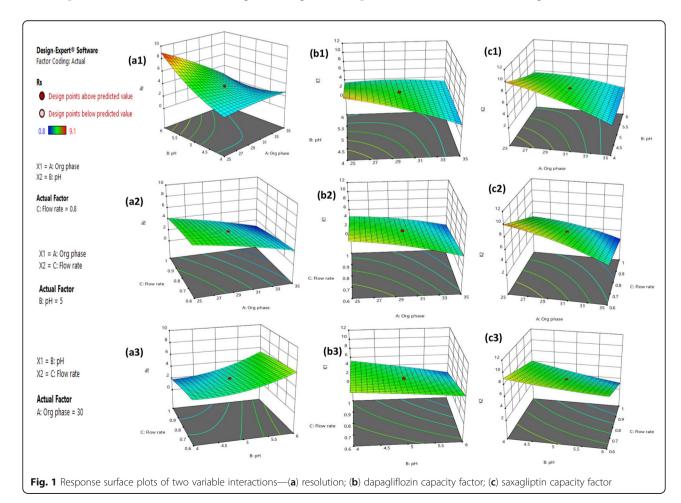
$$+ 0.167X_{2}^{2} + 0.003X_{3}^{2}$$

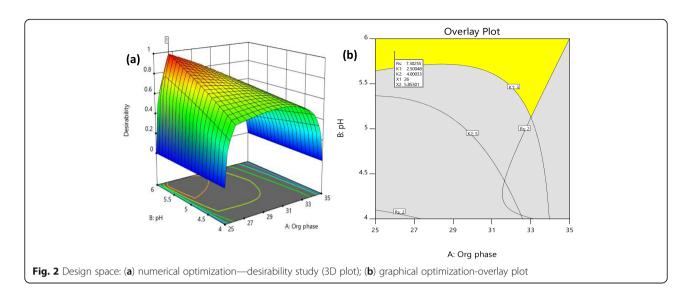
Factor-response relationship and probable interaction effects were studied by response surface analysis. Contour (2D) plots and response surface (3D) plots (Fig. 1) were produced as a function of significant variables while the third variable was kept constant at a definite level. Colored regions (blue-green-yellow-red) in Fig. 1 denoted the obtained response values from low to high with respect to variable interactions. Figure 1a signifies

the influence of variable interaction on resolution, high resolution values observed at high pH, low % acetonitrile, and minimum flow values. Figure 1b, c signifies the factor interactions on capacity factor of both drugs. Optimum K' values are accomplished with high pH and high flow rate at optimum % acetonitrile concentration.

### Design space

Multiple response optimization of variables for chromatographic separation of target analytes is carried using numerical optimization by setting up responses at desired goals. The one with desirability 1 was selected as an optimized solution out of three different solutions provided by the software. The design space generated through Derringer's desirability function was portrayed in Fig. 2a, indicated high method performance owing to maximum desirability value (equal to 1). The perusal to Fig. 2a indicated the different regions for desirability values, the region (marked red) showed the maximum value which meets the desired goals, i.e., maximum resolution and capacity factor at target rage (1-5). The overlay plot of responses obtained from graphical optimization as illustrated in Fig. 2b, which exhibited





method operable design region (MODR) and location of the optimized solution for the studied design. The perusal to Fig. 2b, indicated a region (yellow), which met the all-desired goals such as high resolution and optimum capacity factor values. The gray region with borders indicated the boundaries of responses against variables. MODR for the proposed method can be defined as organic phase (26-30%); pH (5.2-6.0); and flow rate (0.9-1.0 mL/min).

Optimum chromatographic conditions were derived from the design space as mobile phase consists of acetonitrile: phosphate buffer, pH 5.8 (26:74% v/v) at flow rate of 0.96 mL/min gave efficient chromatographic separation of analytes on SPOLAR C18 (250  $\times$  4.6 mm, 5 $\mu$ ) column with isocratic elution. Injection volume was 20  $\mu$ L and UV detection at 236 nm. The chromatogram of standard solution of saxagliptin and dapagliflozin under optimized method conditions was illustrated in Fig. 3. Retention time of dapagliflozin was 3.5 min, saxagliptin was 5.0 min and symmetrical peaks were observed. Tailing factor of both the drugs were less than 2.0, resolution was greater than 2, and plate count of both the drugs were greater than 2000.

### Method validation

The method was validated according to ICH Q2(R1) guidelines and obeyed all validation parameters. A linear calibration plot [S5] for the method was obeyed over the calibration range [S2, S6] of 0.2-300  $\mu g/mL$  for dapagliflozin (Y=17890X+111969,  $r^2=0.9991$ ) and 0.1-150 µg/mL for saxagliptin (Y=73126X-75712,  $r^2$ =0.999). Results of system suitability test and validation were represented in Table 3. System suitability test results indicated that analytical attributes produced with optimized method parameters were analogous within the predicted space. The % RSD values for intra-day and inter-day study were less than 2.0, endorsed the good repeatability of proposed method. Recovery tests confirmed the accuracy of the method and found it to be important within specification limits and afforded recovery 100.1-100.3% and 100.3-100.9% of dapagliflozin and saxagliptin, respectively.

Chromatograms of blank and placebo had lack of peak at the retention time of target analytes. The chromatograms of stressed samples showed well-separated peaks of the analytes and degradation products (resolution>2), without alteration of analyte peak retention times,

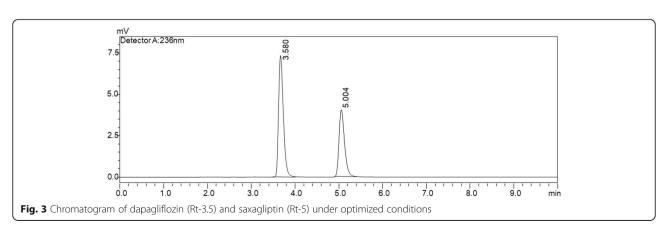


Table 3 Results of method validation

Parameter	Dapagliflozin			Saxagliptin		
Precision						
Repeatability	Peak area	SD	% RSD	Peak area	SD	% RSD
Intra-day	1989124	21017	1.05	3584561	21476	0.599
Inter-day	2018092	23326	1.15	3551248	26363	0.742
Accuracy						
Recovery level	Amount Spiked (µg/mL)	% Recovery	% RSD	Amount Spiked (µg/mL)	% Recovery	% RSD
80%	80	100.3	0.16	40	100.6	0.39
100%	100	100.2	0.10	50	100.9	0.40
120%	120	100.1	0.10	60	100.3	0.57
System suitability						
Retention time	3.5			5.0		
Peak area	515956			602550		
Tailing factor	1.3			1.2		
Theoretical plates	6822			9773		
Capacity factor (K')	2.5			4.0		
Selectivity factor (a)	1.6					
Resolution	7.3					

SD standard deviation, RSD relative standard deviation (n=6)

indicates method specificity. The LOD and LOQ values were found to be 0.061, 0.014  $\mu g/mL$ , and 0.18, 0.043  $\mu g/mL$  respectively for dapagliflozin and saxagliptin. Deliberate changes in method critical variables such as pH, flow rate, and organic phase (%) were made within MODR in order to facilitate the robustness of the method and results were represented in Table 4. The % RSD was below 2 for all variables, which point toward the robust method.

### Application of contemplated method

The results obtained for assay of saxagliptin and dapagliflozin were compared in the marketed formulation (QTERN) with the corresponding labeled quantities and

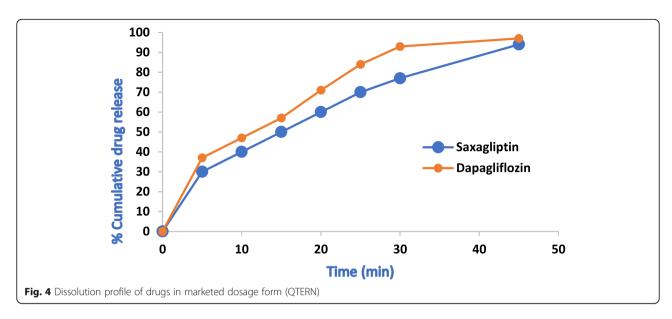
the chromatogram obtained for sample solution had dearth of additional peaks, indicated no interference of the formulation excipients used in the tablets. The % assay of dapagliflozin and saxagliptin and found to be 99.3% and 99.6%, respectively.

Dissolution testing of saxagliptin and dapagliflozin combined marketed tablet formulation was carried out in FDA-described dissolution media. The amount of drug dissolved was analyzed by the proposed RP-HPLC method without interference due to excipients. The cumulative drug release (%) was plotted against the time, depicted in Fig. 4. The perusal to Fig. 4 indicated that 97 and 94% of dapagliflozin and saxagliptin were released respectively from the marketed formulation (QTERN).

Table 4 Robustness data of optimized method

Factor	Change condition	Dapagliflozin	Dapagliflozin (%RSD)		6RSD)
		Rt	Тр	Rt	Тр
Flow rate (mL/min)	0.92	1.02	0.78	1.12	0.57
	0.96	0.31	0.72	0.47	0.41
	1.0	0.56	1.04	0.88	0.86
Organic phase (%)	25	1.11	0.85	1.11	0.66
	26	0.52	0.71	0.24	0.52
	27	0.57	0.83	0.84	0.69
Buffer pH	5.7	0.84	0.68	0.13	0.64
	5.8	0.21	0.71	0.46	0.58
	5.9	0.53	0.76	0.82	0.87

RSD relative standard deviation (n=3)



The maximum in vitro release of drugs was found within 45 min. Hence, the proposed method can be used for dissolution testing of targeted drugs.

Forced degradation studies have been carried out to establish the stability-indicating property of the method and the peaks of degraded products are well estranged from the peaks of title analytes at different retention times, recorded in Fig. 5, proven method's specificity. Summary of degradation studies was reported in Table 5. It was observed that saxagliptin was prone oxidative degradation. Significant degradation of dapagliflozin was observed with acidic and alkali hydrolysis condition. Stress degradation study results of both drugs revealed their stability under neutral hydrolysis condition. However, these degradation products do not be (or less) observed in normal course of degradation.

### Discussion

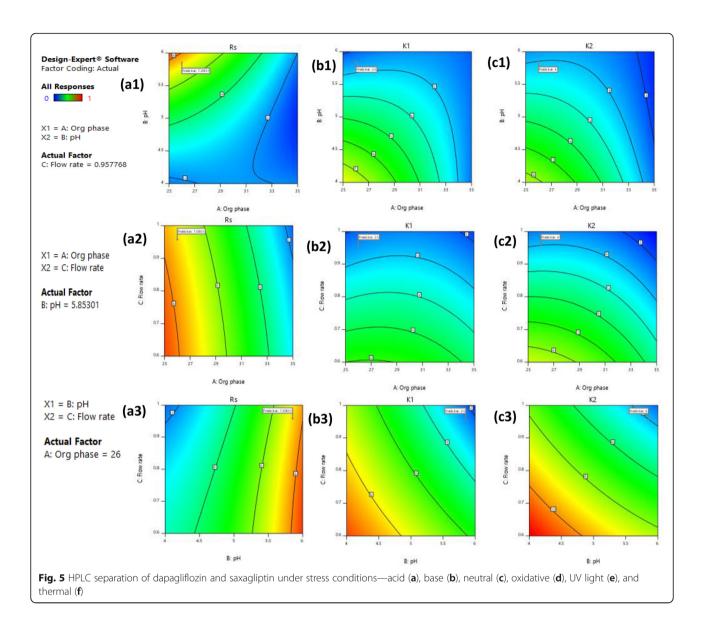
Central composite design (CCD) with response surface methodology (RSM) was exploited to optimize experimental conditions for HPLC separation of saxagliptin and dapagliflozin, since RSM in HPLC method development provides an effective, competent

approach for studying all critical parameters at the same instance, and at two levels (low, high) with lowest number of runs. Statistical analysis revealed that factor effects and interactions are highly significant (p<0.05) at 95% confidence levels. The highest least squares regression values ( $r^2>0.993$ ) of responses are obtained for selected models (quadratic). Coefficient of variation (CV) found to be less than 10% indicates reproducibility of model. A strong relationship between the experimental results and those of the models fitted was suggested. The model aptness is endorsed by lowest PRESS value, high adjusted  $R^2$  value (~1), high adequate precision (>4), non-significant lack of fit (p>0.05), and agreement of predicted  $R^2$ with adjusted  $R^2$  (difference<0.2). High degree of interaction among the studied CMPs on the method attributes is revealed by curvatures in response surface plots.

Figure 6 showed the contour plots of all responses and design space for optimized method are characterized with respect to critical method variables. It can be used to find out the response for a given set of

**Table 5** Results of stress degradation studies

Stress condition	Saxagliptin		Dapagliflozin		
	% Drug recovered	% Drug decomposed	% Drug recovered	% Drug decomposed	
Controlled sample	99.6		99.3		
Acidic (1N HCl, 60 °C, 1 h)	91.2	8.4	84.1	15.2	
Alkali (1N NaOH, 60 °C, 1 h)	92.3	7.3	85.6	13.7	
Neutral ( $H_2O$ , 60 °C, 1 h)	97.3	2.3	96.6	2.7	
Oxidative (6% v/v H <sub>2</sub> O <sub>2</sub> )	80.2	19.4	90.2	9.1	
UV light (48 h)	93.7	5.9	94.1	5.2	
Thermal (105 °C, 6 h)	89.1	10.5	90.6	8.7	

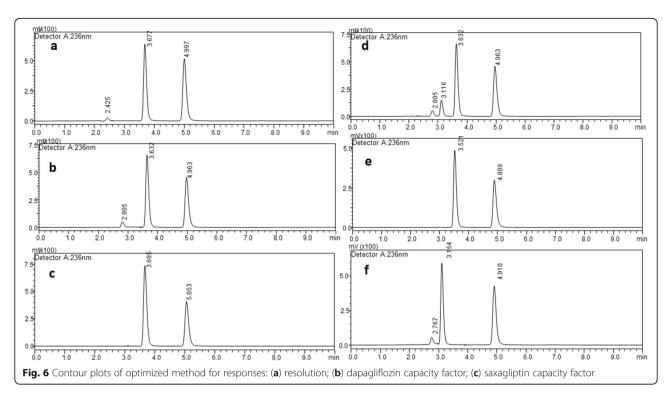


input variables. The perusal to Fig. 6a showed the two variable interaction effect on resolution. The resolution is evidenced to vary non-linearly descending manner with increase in % acetonitrile. The maximum resolution observed with extremely high levels of buffer pH values. High resolution between analyte peaks is recommended to get good separation, accomplished with high flow rate. Figure 6b represents the influence of variables on capacity factor of dapagliflozin. The % organic phase revealed a curvilinear trend with aqueous buffer pH of mobile phase. The influence of variables on the saxagliptin capacity factor is markedly different, portrayed in Fig. 6c. It showed significant interaction between  $X_1$  and  $X_2$ . Low-capacity factor values were observed with mobile phase containing low % organic component and buffer with high pH value.

The accuracy of predicted conditions within design space is verified by actual experimentation, results obtained are compared with predicted space and observed that the % prediction error (= [(observed-predicted)/predicted] $\times$ 100) associated with predicted set of conditions is less than  $\pm$  10%. This assured the consistency of method performance as per intended use.

### Method comparison with literature methods

Present method was found to be cost-effective owing to utilization of low % (26%) organic mobile phase than latest literature [10]. The analytical run duration for the proposed method is short (6 min); hence, this method was rapid and superior than literature methods [8, 11–13]. When compared with previous HPLC reports [14–18] for this drug combination, the proposed method has advantages such as high resolution (7.3) and wide linearity range (0.1-150  $\mu$ g/mL, 0.2-300  $\mu$ g/mL). Low LOD and



LOQ values furnished in the method signify the high sensitivity over literature methods. Moreover, present method was applied to in vitro dissolution as well as stability studies of target analytes. The proposed AQbD method was found to be superior to literature methods due to its ability to simulate the interactive effects of variables on performance of the method. The present strategy of response surface methodology can overcome the limitations of reported methods such as high number of experiments, no variable-interaction study, less robust, and less feasibility for method transfer.

### Conclusion

New RP-HPLC stability portentous method was developed for simultaneous analysis of dapagliflozin and saxagliptin in bulk and pharmaceutical dosage form using analytical quality by design approach. In this strategy, various constraints related to efficient separation of both analytes were considered and critical method parameters were optimized using response surface methodology. The developed method was validated for specificity, accuracy, linearity, precision, robustness, and it conformed all validation parameters of ICH Q2(R1) guidelines. The proposed HPLC method is highly robust for method transfer, regulatory flexibility within design space and allows continuous improvement.

### **Abbreviations**

ANOVA: Analysis of variance; API: Active pharmaceutical ingredient; AQbD: Analytical quality by design; CCD: Central composite design; CMP: Critical method parameters; C: Coefficient of variation; DoE: Design of experiments;

FDA: Food and drug administration; HPTLC: High performance thin-layer liquid chromatography; ICH: International conference on hormonization; K<sub>1</sub>: Capacity factor of dapagliflozin; K<sub>2</sub>: Capacity factor of saxagliptin; LC-MS: Liquid chromatography-tandem mass spectrometry; LOD: Limit of detection; LOQ: Limit of quantification; MODR: Method operable design region; OFAT: One factor at a time; PDA: Photo diode array; RP-HPLC: Reverse phase high performance liquid chromatography; Rs: Resolution; RSD: Relative standard deviation; RSM: Response surface methodology; USP: United States Pharmacopeia; UV: Ultra violet

### **Supplementary Information**

The online version contains supplementary material available at https://doi.org/10.1186/s43094-021-00229-z.

**Additional file 1: S1: Table.** Critical method parameters for optimization. **S2: Table.** Data of linearity studies. **S3: Figure.** Overlaid UV-absorption spectra of Dapagliflozin (20 μg/ml) Saxagliptin (10 μg/ml). **S4: Figure.** Chromatograms obtained under CCD runs. **S5: Figure.** Calibration plot of dapagliflozin (a) and saxagliptin (b). **S6: Figure.** Overlaid chromatogram - linearity studies.

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### Authors' contributions

We assure that "all authors have read and approved the manuscript." SG had analyzed the samples and completed this work under the supervision of SR and SCVS. PDA helped in experimental work and data analysis. All authors together contributed for this research work.

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### Availability of data and materials

The data generated during this study is included in the supplementary file of this article.

### **Declarations**

## **Ethics approval and consent to participate** Not applicable.

### Consent for publication

Not applicable.

### Competing interests

No competing interests to declare.

### Author details

<sup>1</sup>Department of Pharmacy, University College of Technology, Osmania University, Hyderabad 500007, India. <sup>2</sup>Gokaraju Rangaraju College of Pharmacy, Osmania University, Hyderabad 500007, India. <sup>3</sup>Department of Chemistry, University College of Science, Osmania University, Hyderabad 500007, India.

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