



Mixture Design-driven Statistical Optimization and Method Validation in HPTLC: Targeting Rutin, Quercetin, and Gallic Acid

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Abstract

Rutin (RT), Quercetin (QT), and Gallic Acid (GA) are recognized for their potent antioxidant and anticancer properties, prevalent across numerous plant species. The precise quantification of RT, QT, and GA is pivotal for evaluating the therapeutic potential of plant-based substances. In response, a new, straightforward, cost-effective, and reliable method using High-Performance Thin-Layer Chromatography (HPTLC) has been developed and validated for the quantification of these compounds. The mobile phase optimization employed a mixture design approach, achieving chromatographic separation with a mobile phase mixture of toluene, ethyl acetate, menthol, and formic acid at specific ratios (3.56:3.70:0.94:1.80 v/v/v/v/v/v/v/v). Silica gel 60 F254 HPTLC plates were utilized for the analysis. The retention factors (R_f) observed for RT, GA, and QT were 0.21, 0.58, and 0.74, respectively. This method demonstrated a robust linear relationship for concentrations ranging from 400 to 2000 ng per band, with correlation coefficients (R²) of 0.9921 for RT, 0.9936 for QT, and 0.9912 for GA. The Limits of Detection (LOD) and quantification (LOQ) were established at 100.84, 102.38, and 84.54 ng per band for LOD, and 305.58, 310.25, and 256.18 ng per band for LOQ, respectively, for RT, QT, and GA. This validated HPTLC method developed through a Design of Experiment (DoE) approach was successfully employed for the quantification of GA, QT, and RT from the fruits of *Adansonia digitata* and leaves of *Grewia asiatica* ethanolic extracts.

Keywords: *Adansonia digitata*, Gallic Acid, *Grewia asiatica*, HPTLC, Quercetin, Rutin

Abbreviations

GA: Gallic acid, ICH: International Council for Harmonization, LOD: Limit of Detection, LOQ: Limit of Quantification, R²: Correlation coefficient, RSD: Relative Standard Deviation, RT: Rutin, QT: Quercetin, SD: Standard Deviation.

1. Introduction

Following the outbreak of the COVID-19 pandemic, there has been a marked increase in the utilization of plant-based medicinal products, highlighting a worldwide trend toward natural healing solutions¹. The primary factor behind this trend is the increasing recognition of the health advantages linked to

plant-derived antioxidants, immunomodulatory substances, and anticancer agents. These natural chemicals have shown encouraging outcomes in augmenting immune function and offering defensive actions against many illnesses, including cancer. Nevertheless, the growing use of these plant-derived products has emphasized the pressing need for standardization to guarantee their safety, effectiveness, and quality^{2,3}.

The process of standardizing plant products is essential to provide consistent medicinal effects and guarantee the safety of consumers. HPTLC is the predominant approach employed by researchers for this objective, in contrast to alternative chromatographic methods^{4,5}. HPTLC is used for standardizing plant products because of its cost-effectiveness, simplicity,

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and ability to handle several samples at once. This makes it an ideal method for analyzing the qualitative and quantitative composition of plant elements⁶.

Crucial phytochemicals from the flavonoid and phenolic acid groups include Rutin (RT), Quercetin (QT), and Gallic acid (GA), respectively (Figure 1). RT, a flavonol glycoside, helps scavenge free radicals and is well-known for its powerful antioxidant⁷⁻⁹ and anticancer¹⁰⁻¹² capabilities. Another flavonoid, QT, has been the focus of many oncology study¹³⁻¹⁶ and antioxidant potential^{14,17-19}. Equally noteworthy is GA, a trihydroxybenzoic acid, which has a powerful antioxidant capacity²⁰⁻²² and, according to new research, might be used to prevent and treat cancer²³⁻²⁵. These chemicals have medicinal value, and the simultaneous HPTLC approach helps comprehend their combined effects, especially in antioxidant and anticancer applications. Improving their use in nutraceuticals and pharmacotherapy, this analytical breakthrough shows great potential for studying and efficiently controlling the quality of these beneficial chemicals.

Despite several pharmacological effects of these substances having been extensively studied, researchers investigating their simultaneous analytical testing approach are noticeably lacking. This lack of research emphasizes the importance of our technique (HPTLC), which provides a simplified and cost-effective way to evaluate these phytochemicals in different herbal formulations both qualitatively and quantitatively.

Not only does our work address a significant gap in phytochemical analysis, but it also lays the groundwork for future studies to investigate their synergistic effects, which might improve their use in the creation of nutraceuticals and pharmaceuticals.

2. Materials and Methodology

2.1 Materials

Rutin (RT), Quercetin (QT), and Gallic acid (GA) were procured from Sigma-Aldrich, USA. Pre-coated TLC plates (Silica gel 60 F254, 10×10 cm, 2mm) from Merck, Germany and other analytical reagents were procured from S.D. Fine Chemicals Limited, Mumbai, India. *Grewia asiatica* leaves and *Adansonia digitata* fruits were collected from the Surat district, Gujarat, India.

2.2 Sample Preparation

2.2.1 Standard Solution Preparation

For this study, we produced separate solutions for analyzing RT, QT, and GA. Each compound was precisely measured to 5 mg and carefully transferred into individual 10 mL volumetric flasks. At first, these compounds were dissolved in 5 mL of AR-grade methanol. To achieve thorough dissolution and uniformity, the solutions underwent sonication for a period of 5 to 10 minutes. Following this process, each flask was filled to the mark with methanol to achieve the desired volume.

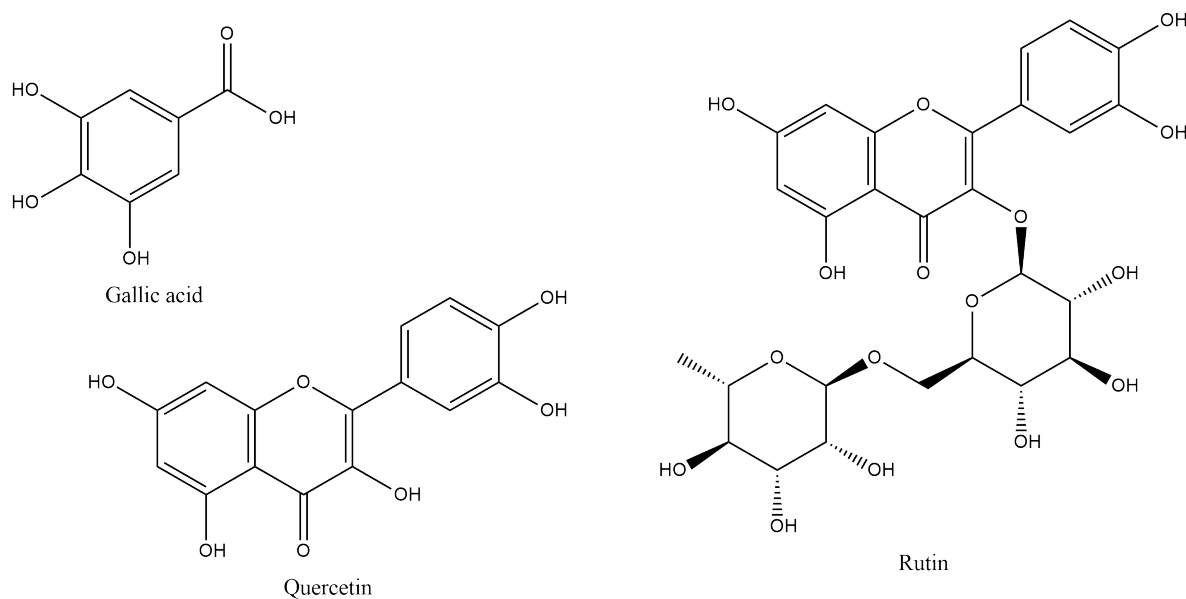


Figure 1. Chemical structure of Rutin, Quercetin, and Gallic acid.

2.2.2 Sample Preparation

The freshly harvested leaves of *G. asiatica* were meticulously collected and subsequently dried beneath a sunshade. At the same time, the fruits of *A. digitata* were delicately opened, and the pulp was carefully extracted and dried. Ethanolic (99.9%v/v) extract was prepared by immersing in the powder plant material for 72 hours at room temperature. The mixture was occasionally agitated to ensure complete extraction. After the maceration process, the extracts were filtrated using Whatman filter paper. The remaining mass was further treated with fresh ethanol to ensure full extraction. At last, the extracted substance was concentrated under reduced pressure using a rotary evaporator.

2.3 Instrumentation of HPTLC

The standard RT, QT, and GA solution were applied using a Linomat 5 device (CAMAG, Muttenz, Switzerland), with a Hamilton syringe (Bonaduz, Switzerland). TLC plates were developed into a twin-trough developing chamber. Following the successful development of Thin-Layer Chromatography (TLC), a UV cabinet was employed to observe the resulting TLC plates. The plates were scanned at 254 nm using the CAMAG TLC Scanner 3, which was equipped with a D2 lamp along with WinCATS 4 software (V 1.4.8).

2.4 Mobile Phase Selection

In the optimization of the mobile phase for HPTLC analysis, initial trials utilized a range of solvents including acetic acid, dioxane, formic acid, ethyl acetate, methanol, n-butanol, isopropanol, and toluene in various proportions. These first investigations uncovered several difficulties, including the low retention factor (R_f) values for RT, the issue of overlapping peaks between QT and GA, and the extremely high R_f values for QT. Modifying the methanol concentration in the mobile phase considerably enhanced the R_f values for RT and QT. However, R_f values were adversely impacted when the methanol percentage fell beyond the range of 2-3 %.

2.5 Statistical Optimization

After extensive testing with different combinations, a mixture of methanol, toluene, ethyl acetate,

and formic acid was used for optimization, as it provided better separation of RT, QT, and GA. The optimization was done by I-optimal Mixture design, considering the total volume of 10 mL, with the mixture components represented as pseudo-components X1 to X4, corresponding to toluene, ethyl acetate, methanol, and formic acid, respectively. The optimization focused on the proportions of these solvents, considering their impact on the peak area and R_f value for each analyte, with each mixture composition being tested once in random order and the centre point run replicated for thorough analysis. Table 1 describes the level of toluene, ethyl acetate, formic acid, and methanol used in statistical optimization. A total of 12 runs as suggested in the design were tested and results were analyzed in design expert software (V. 11.00.0) (Table 2).

2.6 Method Validation

The analytical method used to measure RT, QT, and GA underwent a thorough validation process. This process included tests to ensure accuracy, linearity, precision, repeatability, robustness, sensitivity, and specificity. The validation was performed using the most recent criteria Q2 (R1) established by the International Conference on Harmonization (ICH). The thorough validation process ensured that the method was reliable and met international standards²⁶.

2.6.1 Linearity

The linearity of the association was assessed by examining the correlation between the peak area and marker concentration. This was done by making six consecutive measurements around the concentration range between 400–2000 ng/band for GA, QT, and RT. The resulting correlation coefficient was determined.

Table 1. Mixture design set up for optimization of mobile phase

Factor Code	Factor name	Level		
		Low (-1)	Medium (0)	High (+1)
X1	Toluene	3.2	3.5	3.8
X2	Ethyl acetate	3.7	4	4.3
X3	Formic acid	0.7	1	1.3
X4	Methanol	1.2	1.5	1.8

Table 2. Design Metrix with responses

Run	Independent factors				Dependent Factors					
	X1	X2	X3	X4	Y1	Y2	Y3	Y4	Y5	Y6
	TOL	EA	FA	MeOH	Area-RTN	Area-QUC	Area-GA	RF-RTN	RF-QUC	RF-GA
1	3.2	4.3	1.3	1.2	6310.8	10698.7	9078.1	0.14	0.7	0.54
2	3.70	4.01	1.09	1.20	5524.9	10280.2	9341.8	0.08	0.64	0.46
3	3.80	4.30	0.70	1.20	6186.1	10949.5	9964	0.01	0.6	0.38
4	3.61	3.70	1.30	1.39	5956.4	9396	8831.3	0.14	0.68	0.5
5	3.41	4.26	0.94	1.40	6472	10551.9	4650.4	0.15	0.71	0.55
6	3.41	4.26	0.94	1.40	6472	10551.9	4650.4	0.15	0.71	0.55
7	3.54	3.93	1.02	1.52	7410.4	9261.5	9196	0.2	0.75	0.58
8	3.20	3.99	1.21	1.60	7374.2	10549.5	9941.4	0.23	0.76	0.62
9	3.80	3.71	0.88	1.61	5236.5	9766.6	8471	0.12	0.69	0.51
10	3.57	4.04	0.70	1.69	7941.8	10648.5	10534.6	0.1	0.77	0.6
11	3.20	4.30	0.70	1.80	5507.6	10663.5	9101.7	0.16	0.73	0.59
12	3.41	3.70	1.09	1.80	7737.5	10078.9	9327.4	0.24	0.71	0.57

2.6.2 Precision

The interday and intraday study was done to evaluate precision. The variability of the peak area was measured and shown as a percentage Relative Standard Deviation (% RSD). The experiment's intraday precision was evaluated by carrying out three repetitions of three distinct concentrations on the same day. Similarly, the precision study was undertaken on various days.

2.6.3 Accuracy

Recovery tests were done to evaluate the accuracy of the method. The experiment entailed introducing a pre-established quantity of a reference substance to samples at three distinct concentrations (50%, 100%, and 150%) and subsequently examining them using the proposed method. The analysis was conducted three times for optimal accuracy.

2.6.4 Limit of Detection (LOD) and Limit of Quantification (LOQ)

The sensitivity of the developed method was tested using LOD and LOQ which were calculated from the standard deviation of the response and the slope of the calibration curve of the marker using the following formula:

Where " σ " is the standard deviation of the response and " S " is the slope of the calibration curve.

2.6.5 Specificity

The specificity of the method was ascertained by standard markers and sample comparison. Through the comparison of their R_f values to the standard, the presence of biomarkers in the extract was established. It was found that RT, QT, and GA were detected.

2.6.6 Robustness

The proposed method's robustness was tested by making slight modifications to the methodology, such as altering the wavelength, saturation time, and mobile phase composition, all at the same concentration. The method's robustness was determined by calculating the %RSD, which indicated its ability to maintain accuracy and precision.

2.7 Analyzing Rutin, Quercetin, and Gallic Acid in Ethanol Extracts of *Grewia asiatica* and *Adansonia digitata*

To quantify the concentrations of RT, QT and GA in the ethanolic extracts of *G. asiatica* and *A. digitata*, a concentrated solution (10 mg/mL) was applied

on a precoated HPTLC plate. This was followed by chromatographic separation under an optimized solvent system. The quantification of RT, QT and GA was calculated using peak area. Results, expressed in grams per 100 grams of extract, were derived from triple replicate analyses for accuracy and reliability.

3. Results

3.1 Primary Study

In the development of an optimized HPTLC methodology for the detection of RT, QT, and GA, a comprehensive exploration of mobile phase combinations was undertaken to refine the chromatographic resolution of RT, QT, and GA peaks. Drawing on insights from prior research, which has established the HPTLC technique as a viable method for analyzing these compounds individually or in synergy with additional biomarkers, our experimental design included a diverse array of solvents - ranging from toluene, n-hexane, ethyl acetate, cyclohexane, methanol, formic acid, and glacial acetic acid. This initial phase involved varying solvent combinations and ratios to identify the most effective mobile phase. Ultimately, a mobile phase composition of toluene, ethyl acetate, menthol, and formic acid was identified as superior, yielding clear, well-defined peaks for each marker compound. The optimal conditions were further validated by the distinct R_f values observed for RT (0.21), QT (0.58), and GA (0.74), indicating successful separation. Further refinement of the chromatographic parameters was achieved through a mixture design approach, enhancing the method's efficacy and precision.

3.2 Advancing Chromatography: A Mixture Design Approach

3.2.1 Model Fitting and Regression Analysis

The incorporation of a mixture design approach in this study was predicated on its adaptability and its proficiency in refining HPTLC separations, permitting a detailed examination of both primary and interactive effects exerted by various factors. Initial experimental observations guided the selection of critical parameters for this optimization endeavour, focusing on the volumetric contributions of toluene, ethyl acetate, menthol, and formic acid. The optimization criteria

were centred around the chromatographic metrics of peak area (area under the curve) and retention factor. To mitigate the influence of extraneous variables and ensure the integrity of the outcomes, all experimental runs were randomized. Analytical assessment of the results employed diverse mixture models, enabling the determination of the most suitable model based on the minimization of residual errors and the overall quality of fit. A favourable outcome was indicated by a positive value, whereas a negative value suggested a relationship that went against optimization. The experimental design and outcomes are delineated in Table 2, with the model's validation conducted through Analysis of Variance (ANOVA). Selection criteria for the optimal model included a high coefficient of determination (R^2), minimal discrepancy between predicted and adjusted R^2 values, and a low Predicted Residual Sum of Squares (PRESS), as detailed in Table 3. The model's robustness was further assessed by examining changes in R^2 upon the exclusion of regression variables, with the refined model's validity confirmed via F-statistics analysis (Tables 4 and 5).

Design-Expert software was utilized to generate 3D surface and contour plots (Figure 2), providing a deeper insight into the impact of the variables (Table 6). An overlay plot was created to meet the specified targets for the optimal response variables, which are indicated by the yellow zone. (Figure 3) After statistical optimization, checkpoint analysis was done by selecting three different compositions and results were verified against the predicted R_f value and AUC (Table 7). Final optimized mobile phase composition X1:X2X3:X4 [(TOL:EA:FA:MeOH) (3.56:3.70:0.94:1.80)].

3.3 Method Validation

The validation of the method was rigorously conducted by the International Conference on Harmonization (ICH) Q2 (R1) guidelines, evaluating various parameters such as accuracy, linearity, Limit of Detection (LOD), Limit of Quantification (LOQ), precision, and robustness.

3.3.1 Linearity

The developed HPTLC method demonstrated a significant correlation for RT, QT, and GA within a concentration range of 400-2000 ng/band, illustrated in Figure 4. Linearity for each compound was established

Table 3. Model selection summary

Response	Source	SD	R ²	Adj. R ²	Pred. R ²	PRESS
Y1: Area-RTN	Linear	967.84	0.182	-0.124	-1.387	21872689
	Quadratic	236.84	0.988	0.933	-17.898	173193934
	Special Cubic	7.78	1.000	1.000		+
Y2: Area-QUC	Linear	383.680	0.6388	0.503	0.320	2218000
	2FI	567.200	0.8027	-0.085	-303.698	993500000
	Quadratic	14.140	0.9999	0.999		+
Y3: Area-GA	Linear	2135.577	0.1005	-0.237	-0.868	75769789
	2FI	1155.160	0.9342	0.638	-100.627	4122143337
	Quadratic	2.121	1.0000	1.000		+
Y4: RF-RTN	Linear	0.03	0.866	0.816	0.733	0.0
	2FI	0.02	0.972	0.847	-40.335	1.8
	Quadratic	0.01	0.999	0.988		+
Y5: RF-QUC	Linear	0.035	0.6260	0.486	0.033	0.025
	2FI	0.010	0.9930	0.961	-6.954	0.209
	Quadratic	0.007	0.9981	0.979		+
Y6: RF-GA	Linear	0.035	0.7981	0.722	0.452	0.027
	2FI	0.007	0.9980	0.989	-0.503	0.074
	Quadratic	0.007	0.9990	0.989		+

with regression equations of $y = 4.8221x + 2443$ ($r^2 = 0.9921$) for RT, $y = 5.366x + 3957$ ($r^2 = 0.9936$) for QT, and $y = 5.554x + 3594$ ($r^2 = 0.9912$) for GA, indicating precise analytical performance.

3.3.2 Precision

The inter-day precision for RT, QT, and GA was determined to have mean % RSD values of 1.02, 0.83, and 1.01 respectively where $n = 3$. The intraday precision for RT, QT, and GA was measured to be 1.6, 1.41, and 1.59 (mean % RSD, $n = 3$) respectively.

3.3.3 Limit of Detection (LOD) and Limit of Quantification (LOQ)

The Limits of Detection (LOD) for RT, QT, and GA were 100.84 ng/band, 102.38 ng/band, and 84.54 ng/band, respectively. Correspondingly, the Limits of Quantification (LOQ) for these compounds were 305.58

ng/band for RT, 310.25 ng/band for QT, and 256.18 ng/band for GA, showcasing the method's sensitivity.

3.3.4 % Recovery

The recovery studies aimed at evaluating the accuracy of the established HPTLC technique yielded average percentage recoveries for RT, QT, and GA, with values of 100.79%, 100.72%, and 105.09% across three trials, respectively.

3.4 Quantification of RT, QT and GA in Plant Extract

In the analysis of the ethanolic extracts from *G. asiatica* and *A. digitata*, chromatography identified three prominent peaks at R_f values of 0.74, 0.58, and 0.21, which corresponded to QT, GA, and RT which is aligned perfectly with the standards, indicating no detectable interference from other constituents within

Table 4. Regression analysis

Model term	Y1: Area-RTN			Y2: Area-QUC		
	Full model		Reduced Model	Full model		Reduced Model
	Coefficient	p-value	Coefficient	Coefficient	p-value	Coefficient
X1	-5282.98	0.0929	-5604.38	10642.6	0.3322	9623.38
X2	-148.37		336.18	10970.3		11793.88
X3	11789.4		12192.44	10752.48		9441.51
X4	7983.34		7806	12561.36		9994.77
X1X2	35423.5	0.0113	35200.44	1011.98	0.9216	-
X1X3	101.72	0.9804	-	-5605.38	0.5881	-
X1X4	20766.69	0.0335	21718	-5495.25	0.6157	-
X2X3	1754.8	0.6865	5598.34	-156.54	0.9877	-
X2X4	6165.58	0.2344	-	-3942.54	0.6971	-
X3X4	-6065.89	0.2602	-6374.2	-4878.33	0.6536	-
	Y3: Area-GA			Y4: RF-RTN		
X1	20688.55	0.5302	27921.22	-0.1056	0.0464	-0.06
X2	-13148.44		-8302.84	0.0714		0.08
X3	30775.56		26291	-0.0551		-0.09
X4	28132.8		26958.98	0.1578		0.24
X1X2	23879.97	0.3263	-	0.0894	0.8439	-
X1X3	-59637.44	0.0792	-62340.1	0.5788	0.2722	0.6
X1X4	-47356.85	0.1304	-58361.8	0.227	0.6358	-
X2X3	51.48	0.998	-	0.5061	0.3293	0.56
X2X4	5487.3	0.7878	-	0.1614	0.7163	-
X3X4	-66484.14	0.073	-55428.2	0.8914	0.1622	0.82
	Y5: RF-QUC			Y6: RF-GA		
X1	0.391	0.0167	0.355	0.16	0.0037	0.168
X2	0.411		0.396	0.19		0.166
X3	0.792		0.827	0.57		0.637
X4	0.524		0.586	0.49		0.518
X1X2	0.79	0.0361	0.889	0.82	0.0184	0.858
X1X3	-0.017	0.9195	-	0.12	0.3687	-
X1X4	1.047	0.0222	1.02	0.88	0.0167	0.84
X2X3	0.386	0.1273	0.36	0.63	0.0297	0.57
X2X4	1.042	0.0198	0.96	1	0.0114	0.99
X3X4	0.196	0.3411	-	0.17	0.2854	-

Table 5. Model summary statistics

Model	Model	Df	F-value	P-value (model)	R ²	SSE	MSE	No. of term omitted	p-value (Lack of Fit)	Fcalculated	Fcritical (α=0.05)
Y1: Area-RTN	1	9	17.93	0.0539	0.98775	112191	56095.4	2	0.058	0.117	3.677
	2	7	41.21	0.0014	0.9863	125300	31332.3				
Y2: Area-QUC	1	9	0.9	0.6281	0.8027	643441	321720	6	0.057	0.277	8.812
	2	3	4.72	0.0353	0.6388	1178000	147207				
Y3: Area-GA	1	9	3.16	0.2638	0.9342	2669000	1334000	3	0.501	0.576	4.099
	2	6	5.96	0.0345	0.8774	4973138	994628				
Y4: RF-RTN	1	9	7.75	0.1195	0.9721	0.0012	0.0006	3	0.272	0.132	4.099
	2	6	24.13	0.0015	0.9666	0.0015	0.0003				
Y5: RF-QUC	1	9	31.31	0.0313	0.993	0.00019	0.00009	2	0.478	0.818	3.677
	2	7	44.03	0.0013	0.9872	0.00034	0.00008				
Y6: RF-GA	1	9	111.9	0.0089	0.998	9.8E-05	4.9E-05	2	0.557	1.475	3.677
	2	7	115.91	0.0002	0.9951	0.00024	0.00006				

1=Full Model, 2=Reduced Model

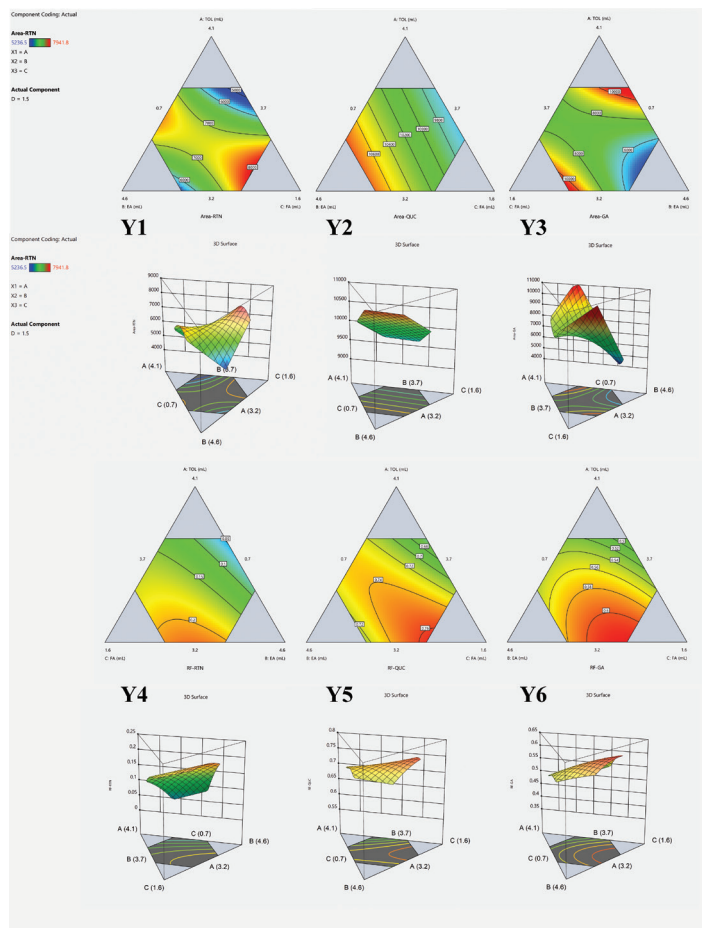


Figure 2. Contour and 3D surface plots for responses (Y1-Y6).

Table 6. Response target for optimization

Response	Response	Lower limit	Upper limit	Target set for optimization
Y1	Area-RTN	5236.5	7941.8	maximize
Y2	Area-QUC	9261.5	10949.5	maximize
Y3	Area-GA	4650.4	10534.6	maximize
Y4	RF-RTN	0.01	0.24	0.24
Y5	RF-QUC	0.6	0.77	0.7
Y6	RF-GA	0.38	0.62	0.5

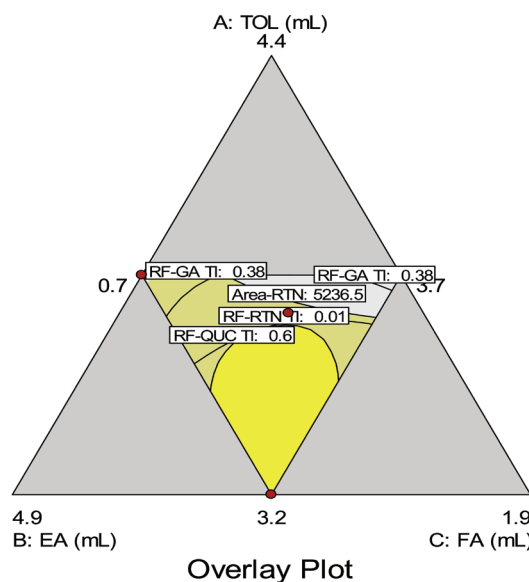


Figure 3. Design space (Yellow region) in overlay plot.

Table 7. Checkpoint analysis

Suggested solutions	TOL:EA:FA:MeOH (X1:X2X3:X4)	Y1: Area-RTN	Y2: Area-QUC	Y3: Area-GA	Y4: RF-RTN	Y5: RF-QUC	Y6: RF-GA
Predicted responses							
1*	3.56:3.70:0.94:1.80	7278	9774	9084	0.200	0.719	0.563
2	3.31:3.70:1.30:1.69	8275	9684	10317	0.240	0.723	0.573
3	3.34:3.70:1.26:1.69	8098	9690	9687	0.239	0.722	0.571
Observed responses							
1*	3.56:3.70:0.94:1.80	7337	10282	9699	0.21	0.74	0.58
2	3.31:3.70:1.30:1.69	7894	9626	9462	0.26	0.76	0.62
3	3.34:3.70:1.26:1.69	7648	10402	10724	0.21	0.67	0.53
% Prediction error							
1*	3.56:3.70:0.94:1.80	-0.81	-5.20	-6.77	-5.00	-2.94	-2.94
2	3.31:3.70:1.30:1.69	4.61	0.60	8.28	-8.33	-5.06	-8.22
3	3.34:3.70:1.26:1.69	5.56	-7.34	-10.70	12.23	7.16	7.26

*Final optimized mobile phase composition as there was least perdition error for all responses

the extracts. Results described in Table 8 showed that the ethanolic extracts of *G. asiatica* leaves contained higher levels of RT, QT, and GA compared to *A. digitata* fruits.

4. Discussion

Biomarker study is the most crucial part of herbal drugs. The development and validation of analytical

Table 8. Estimated content of RT, QT, and GA in ethanolic extracts of *Adansonia digitata* fruits and *Grewia asiatica* leaves

Biomarker	Ethanolic extract of AD	Ethanolic extract of GA
RT	832.56 ng/spot	1614.06 ng/spot
QT	2586.65 ng/spot	3411.85 ng/spot
GA	1743.4 ng/spot	1865.71 ng/spot

QT: Quercetin, RT: Rutin, GA: Gallic acid

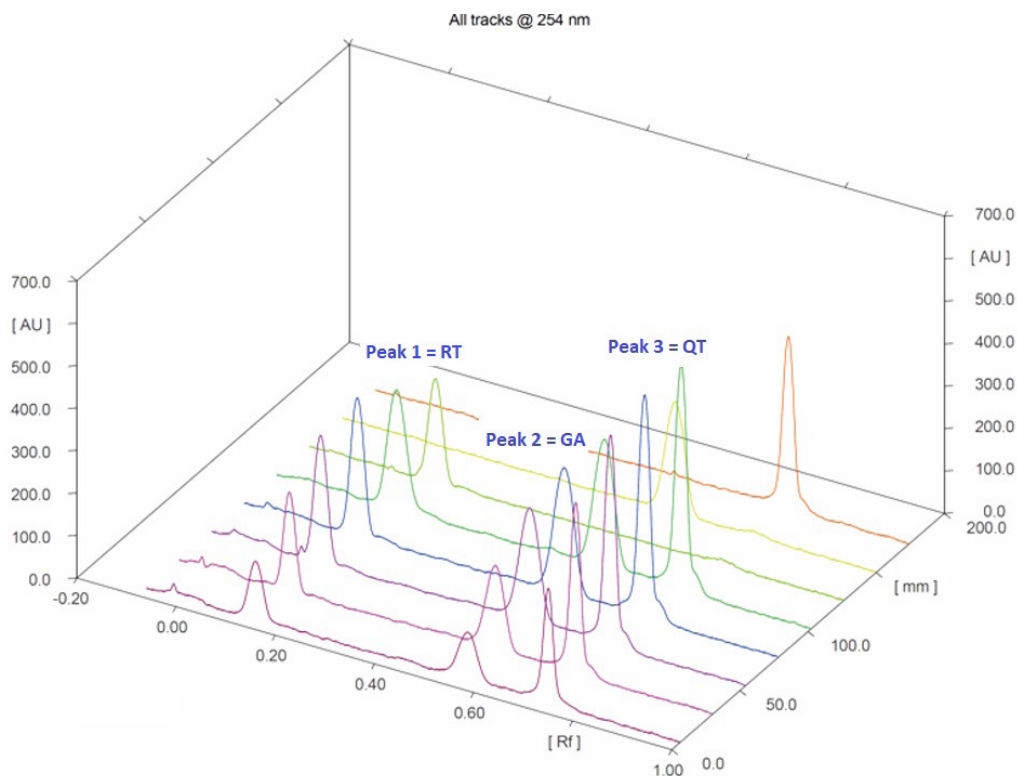


Figure 4. 3D chromatogram of Rutin, Quercetin, and Gallic acid.

methods are crucial for ensuring the quality of herbal medicines. Herbal industries utilize a range of sophisticated instrumental techniques, such as High-Performance Liquid Chromatography (HPLC), Liquid Chromatography-Tandem Mass Spectrometry (LC-MS/MS), Gas Chromatography (GC), Gas Chromatography-Tandem Mass Spectrometry (GC-MS/MS), and High-Performance Thin-Layer Chromatography (HPTLC), to quantify various phytochemicals in herbal formulations. RT, QT, and GA are significant phytochemicals whose simultaneous quantification is essential for the standardization of herbal drugs. Therefore, there is a need for an economical and user-friendly method that facilitates this process. Several HPTLC methods for simultaneous estimation of selected phytochemicals are available in the literature. In a study conducted by Kumar, and Lakshman²⁷, the concentration levels of RT, QT, and GA in *Terminalia chebula* were quantified using HPTLC. The author devised two separate analytical methods for the quantification of three compounds. For RT and QT, a mobile phase consisting of ethyl acetate, dichloromethane, formic acid, glacial acetic acid, and

water in the ratio of 10:2.5:1:1:0.1 (v/v/v/v/v/v) was employed. In contrast, a mixture of toluene, acetone, and glacial acetic acid in the ratio of 3:1:2 (v/v/v) was utilized specifically for the quantification of GA.

In another study, Alam *et al.*, also developed two distinct methods for analyzing RT, QT, and GA in an anti-HBV active extract of *Guiera senegalensis*. They used acetonitrile: water (4:6, v/v) and Toluene: Ethyl acetate: Formic acid (6:4:0.8, v/v/v) as solvents for the analysis of rutin, quercetin, and gallic acid, respectively. There are two distinct methods for measuring RT, QT and GA, which require more time and involve higher costs for analysis. Several other methods were also reported by various researchers²⁸⁻³⁰. Compared to reported studies, our method offers a fast easy to perform and cost-effective HPTLC method of analysis.

The mobile phase plays a crucial role in the method development. To develop a precise, sensitive, easy-to-operate and cost-effective HPTLC method, various mobile phases were experimented with to enhance the resolution between the compounds. Through our investigation, we discovered that the combination of toluene, ethyl acetate, formic acid, and methanol

provides superior resolution when compared to other solvents. Design mixture assists in identifying the optimal combination of solvents to achieve the desired R_f value within the design space.

An optimized mobile phase composition was achieved using a simple centroid design with axial points in a pseudo-component representation. The simple centroid design is a design that lies on the boundary. By conducting additional experiments within the simplex, we can accurately describe the properties of the mixture. Therefore, the typical simplex design was enhanced by the axial runs. In addition, the centroid was adjusted due to it not being a design point³¹.

The present study resolves the issue of resolution of all three phytochemicals in HPTLC analysis. The method successfully quantifies RT, QT, and GA levels in ethanolic extracts of *A. digitata* and *G. asiatica*. Further testing of the method in other plant extracts is needed to confirm the applicability of the designed method.

5. Conclusion

The study's results highlighted that the separation resolution of RT, QT, and GA is significantly affected by the mobile phase's composition. An effective optimization of the mobile phase was achieved through the application of a simplex centroid mixture design coupled with a desirability function approach. Furthermore, this process led to the development of a new analytical method characterized by its efficiency and user-friendliness, particularly for the quantification of RT, QT, and GA in ethanolic extracts of *A. digitata* and *G. asiatica*. This method offers time savings and presents a practical solution for routine analysis. Future directions include extensive testing and validation across various plant extracts to broaden its applicability for all extracts containing RT, QT and GA, thereby underscoring its potential value to the herbal pharmaceutical industry.

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