Research Article | Pharmaceutical Sciences | Open Access | MCI Approved

Online ISSN: 2230-7605, Print ISSN: 2321-3272

**UGC Approved Journal** 

# RP-HPLC Method Development and Validation of Chrysin in Bulk and Marketed Formulation

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Received: 7 Oct 2018/ Accepted: 9 Nov 2018/ Published online: 01Jan 2019 Corresponding Author Email: Shaheen.pharmchem@gmail.com

## **Abstract**

Chrysin has interesting activity profile. It exerts anti-inflammatory, anticancer, hypoglycemic, antioxidant and anti-ageing properties. In this study a novel, simple, precise and linear reveres phase high performance liquid chromatography method (RP-HPLC) is described for bulk and marketed formulation of chrysin. Methanolic solution of chrysin was injected (injection volume  $20\mu l$  at a flow rate 1ml/min) in reversed phase column using acetonitrile and methanol (65:35V/V) as mobile phase. Detection was set at 268 nm ( $\lambda$ max). The method was found to be linear with correlation coefficient of 0.99 (0.2-6.4 mg of chrysin). Using this method it was possible to detect chrysin at 3.39 min of retention time and method is validated by determining all the validation parameters.

#### Keywords

Chrysin, Flavonoid, Method validation, Marketed formulation, RP-HPLC.

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# INTRODUCTION

Chrysin (Fig.1: 5, 7-di hydroxyl-2-phenyl-4Hchromen-4-one) belongs to flavone class possessing potential antioxidant activity. The other noteworthy pharmacological activities of chrysin hepatoprotection, anti-inflammatory and antiaromatase activity. Passiflora caerulea. passiflora incarnate extracts contain noticeable amounts of chrysin. Literature indicated that upon oral administration it is rapidly excreted from the body and hence its bioavailability is very poor [1].

Ultraviolet-spectroscopy, High Performance Liquid Chromatography (HPLC), reversed-phase HPLC (RP-HPLC) based methods were developed for the analysis and validation of chrysin (**Table.1**) [2-14]. Recently Ying Jia et al developed and validated ultraperformance liquid chromatography tandem mass spectrometry (UPLC-MS/MS) for the simultaneous estimation of chrysin and tectochrysin (*Alpinia oxyphylla* fruits extract) [14]. With an aim to estimate



chrysin at less retention time it was planned to develop and validate a simple RP-HPLC method.

#### **MATERIALS AND METHOD**

#### Instruments used

Denver M-220D Analytical balance was used for weighing, Enerteck Sonicator was used for the sonication, Shimadzu UV-1800 Ultraviolet-Visible Spectrophotometer was used for spectrophotometric determination and Shimadzu L201447 HPLC was used for the chromatographic separation.

#### Chemicals

Pharmaceutical grade pure chrysin (Active pharmaceutical ingredient) sample was procured from Sigma Aldrich, Reference Standard chrysin was purchased from MRM manufacturer, Disodium hydrogen orthophosphate dehydrate, methanol, sodium chloride, potassium bromide, acetonitrile (HPLC Grade) and methanol (HPLC Grade) were purchased from Merck chemicals.

# UV-SPECTROPHOTOMETRIC ESTIMATION OF CHRYSIN

#### Solvent selection

Different solvents like water, methanol, 0.1N sodium hydroxide and phosphate buffer (PH 6.8) were used but as phosphate buffer gave a single distinct peak with good absorbance for chrysin, it was employed as the solvent.

# Preparation of buffer PH 6.8

3.5g of disodium hydroxide orthophosphate dehydrate was dissolved in 100ml of water. The P $^{\rm H}$  was adjusted to 6.8 $\pm$  0.05 with ortho phosphoric acid and filtered through 0.45 $\mu$  membrane filter.

# Determination of $\lambda_{\text{max}}$

 $10\mu g/ml$  chrysin solution was prepared by using phosphate buffer as a solvent. A single distinct peak with good absorbance was obtained at 268 nm hence it was considered as  $\lambda_{max}$ .

# METHOD DEVELOPMENT OF CHRYSIN USING RP-

Preparation of solutions used in the present work

## 1) Preparation of mobile phase solution

A degassed mixture of acetonitrile and methanol in ratio of 65:35 v/v was prepared, and the  $P^{H}$  6.8 adjusted by using phosphate buffer.

# 2) Preparation of standard solution and sample solution

100 mg of chrysin was accurately weighed and transferred into a 100 ml volumetric flask, dissolve with diluents. Take 1ml of this solution into 10 ml volumetric flask and then makeup mark with diluents. Similarly test samples were prepared using chrysin

## **OPTIMIZATION OF THE METHOD**

Reverse-Phase column (Hypersil BDS  $C_{18}$ :  $5\mu m$ , spherical,  $250 \times 4.6$  mm dimension) was selected for chromatographic separation. Chrysin was freely soluble in acetonitrile and DMSO (Dimethyl sulfoxide) where as it was slightly soluble in methanol. Four trials were performed by changing the concentrations of mobile phase (Table-2). At higher concentrations of acetonitrile broad peak was obtained where the ratio of acetonitrile and methanol was set at 65:35, symmetric peak was observed at room temperature. The diluent was used to prepare standard and sample solutions. The chromatogram of chrysin was observed by injecting  $20\mu l$  solution and the flow rate was adjusted to 1ml/min.

# **VALIDATION PARAMETERS**

Validation of the optimized method was performed according to the ICH Guidelines [15-17].

## **System Suitability**

System suitability was carried out with six replicate injections of chrysin solution into the chromatographic system. The results were obtained as SD (Standard Deviation) and %RSD (Relative Standard Deviation) (Table.3).

#### Linearity

For determination of linearity, appropriate aliquots were prepared with concentrations of 25%, 50%, 75%, 100%, 125% and 150%. To prepare these solutions, 0.2 – 6.4 ml standard solutions were pipetted out into a series of 10 ml volumetric flasks and volume was made with the solvent. Average peak area was determined by introducing standard solutions into the column (in duplicate) Calibration curve was plotted with observed peak area against concentration followed by the determination of regression equation and calculation of the correlation coefficient. Calibration curve for chrysin was shown in Fig.3 and their corresponding linearity parameters were given in **Table-4**.

# Specificity

The blank was injected to check the interference for the sample peak by injecting  $20\mu l$  of blank solution (**Table-5**).

## **Method Precision**

The repeatability of the method was verified by calculating the % RSD for the area of six replicate injections of chrysin (16  $\mu g/ml)$ . The results were given in **Table.6**. % RSD for peak area and for retention time were found to be 0.67 and 0.65 respectively. To perform the intermediate precision the above mentioned method was followed on different days by applying same chromatographic



conditions. The % RSD for peak area was found to be within the specified limits.

#### Robustness

To report robustness, variation in flow rate and wavelength were studied (**Table-7**). Injections were delivered by changing the flow rate (range: 0.8-1.2 ml/min; optimized flow rate: 1 ml/min).

To study the effect of variation in wavelength standard solutions were eluted and their presence was detected at different wavelengths (265 nm & 270 nm; optimized wavelength 268 nm).

#### Accuracy

To calculate % recovery, the standard addition method was followed and the results were given in Table-8. For this samples were spiked with  $20\mu l$  of solutions of 50, 100 and 150% standard solutions of chrysin.

# LOD (Limit of Detection)

The LOD and LOQ were calculated by using the formulas LOD= $3.3\sigma/s$  and LOQ= $10\sigma/s$ .

LOD=3.3xstandard deviation of intercept/average of slope

LOD=3.3x78571.24/102323 =2.533

The LOD of standard solution of chrysin was found to be 2.533

## LOQ (Limit of Quantification)

LOQ= 10x Standard deviation of intercept/average slope

**LOQ=**10x78571.24/102323 =7.678

The LOD of the standard solution of chrysin was found to be 7.678

## **RESULTS AND DISCUSSION**

To optimize the RP-HPLC method, a series of four trials were made with different mobile phase compositions and finally 65:35; v/v mixture of acetonitrile and methanol was selected as mobile

phase. Chrysin was found to have appreciable absorbance at 268 nm when determined spectrophotometrically and hence it was selected as detection wavelength. At optimized chromatographic conditions, symmetric peak and better resolution were observed for chrysin. An optimized chromatogram showing retention values was shown in figure-2.

%RSD value for peak area was found to be 1.4 and for retention time it was obtained as 0.19. As %RSD values are within specified limits (<2) it can be said that system suitability had met the requirements of the method validation. Within the concentration range of 0.2- 6.4 mg/ml of chrysin the method resulted in linear plot with correlation coefficient of 0.99 indicating that the method is linear. The proposed method was found to be precise and reproducible with %RSD was found to be <2 % (0.67 for retention time and 0.65 for peak area). The accuracy of the method was verified by performing recovery studies and values are in the range of 99.0-100.6 which were within the acceptance criteria (98 – 102%). The method was found to be specific after verifying the chromatograms showing that there is no interference of the blank. The limit of detection and quantification values for chrysin was found to be 2.533 and 7.678 respectively. The results of robustness of present method had shown that changes made in flow rate and wavelength did not produce significant changes in analytical results. As the changes are not significant it can be said that the method is robust.

This method is suitable to estimate chrysin using RP-HPLC at retention time of 3.39 min, fulfilling the objective of the present study. The study also concludes that the developed method is a simple and precise.

Table: 1 Details of various analytical methods used for the estimation of Chrysin (single or in combination)

Reference	Details of the	e method	Compounds analyzed
	Method employed:	Liquid chromatography/mass	
	spectrometry(LC/MS	) with turbo ion spray	
	Column	: C <sub>18</sub> narrow bore column	
Careri M et al.,	Mobile phase	: formic acid and acetonitrile	Chrysin
1999[2]	(20:80, v/v)		Chrysin
	Flow rate	: 200µL/min	
	Detection wavelengt	h: 280 nm- 370 nm and elution of	
	the solvent mixture f	or 7 min	
	Method employed	: HPLC with UV detection	
	Column	: chromsep RP 18	
Gremiao et al., 2003	Mobile phase	: water and acetonitrile	Chrysin (Propilis
[3]		(97.5:2.5 ,v/v)	extract)
	Detection wavelengt	h : 310 nm	
	Retention time	: 29 min	

Chrysin

Chrysin

Various alkyl and acyl

derivatives of chrysin

Chrysin (Oroxylum

Chrysin (Oroxylum

Chrysin and Galangin

indicum)

indicum)



Method employed: HPLC with Photodiode array

detector

Mobile phase : methanol (55%) and triflouro acetic

acid (0.3%)

Walle et al., 2001[4] Detection wavelength: 268 nm

> Retention time: Chrysin-19.8min Chrysin glucuronide - 3.7 min Chrysin sulphate -6.7 min

Method employed: HPLC with UV detector

Column: Nova pack C<sub>18</sub> column,

Kyoung Soon Kim et al., 2002 [5]

Mobile phase: methanol and phosphoric acid (79:21),

Column temperature: room temperature

Flow rate : 1ml/min Detection wavelength: 280nm Retention time : within 10 min

Method employed: HPLC with diode array detector

Column : Luna C<sub>18</sub> column,

Column temp : 30°C.

Maria C et al., 2003

[6]

[7]

Mobile phase : formic acid:acetonitrile:2-propanol

(70:22:8)

Flow rate : 0.2 ml/min

Detection wavelength: 306 nm- 370 nm.

: < 20 min Retention time

Method employed: RP-HPLC with UV detection

Column : Hypersil BPS-RP C<sub>18</sub>

Mobile phase : water: methanol: acetonitrile: ortho

Zaveri M et al., 2008

phosphoric acid(60:30:38:1 v/v/v/v)

: 1 ml/min Flow rate Detection wavelength: 262nm

Retention time : 12.31 min

Method employed: HPTLC(High pressure thin layer

chromatography)

Saraf Aparna et al.,

2011 [8]

Mobile phase : chloroform: methanol: formic acid

(8.8:0.7:0.5)

Detection wavelength: 254nm and 366 nm Scanner : Camag TLC scanner

Method employed: HPLC method : Diamonsil C<sub>18</sub> Column

: methanol : phosphoric acid Mobile phase

Nie P et al., 2013 [9] : 1 ml/min Flow rate

Detection wavelength: 268 nm

: 25 min(Chrysin) and 8 min(Galangin) Retention time

Method employed: RP-HPLC (High performance liquid chromatography) with MS (mass spectrometry) and

diode array detection

Cui-ping Z et al.,

: HP-C<sub>18</sub> column Column 2014 [10]

Mobile phase : 1% formic acid and methanol

Flow rate : 1.0 ml/min

Detection wavelength: 280 nm

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Chrysin (Chinese

propilis)



Retention time : around 65 min Method employed: LC-UV-ESI-Q/TOF/MS Column :C<sub>18</sub> (4.6 mm x 150 mm, 5μm) Wen-TingTang et Flow rate : 0.3 ml/min Chrysin (Scutellaria Detection wavelength: 278 nm al.,2014 [11] baicalensis) Retention time : 14.81 min Method employed: HPLC method with MS(Mass spectrometry) Keumhan Noh et al.. : Atlantis dc C<sub>18</sub> column Column Chrysin 2016 [12] Mobile phase : 0.1% formic acid and acetonitrile : 0.25 ml/min Flow rate Retention time: 7 min Method employed: Simultaneous RP-HPLC Column: Hypersil BDS C<sub>18</sub> column Vallabaneni Mobile phase: water and acetonitrile (55:45v/v) madhavarao et Column temp: 25°c Chrysin al.,2018 [13] Flow rate : 1.0ml/min Detector : photo diode array Detection wavelength: 263nm Method employed: UPLC-MS/MS(Mass spectrometry) Column : ACQUITY UPLC BEH C<sub>18</sub> Mobile phase : 0.1% Formic acid and methanol Chrysin and Ying Jia et al., 2018 Detection : Multiple monitoring with electrospray Tectochrysin [14] ionization in the positive ion mode (Alpinia oxyphylla) Retention time : 2.09 min (Chrysin) 3.30 min (Tectochrysin)

Table -2: Trial conditions in HPLC method development

Chromatographic Conditions	Trial Condition-1	Trial Condition-2	Trial Condition- 3	Trial Condition-4	
Column	Hypersil BDS C <sub>18</sub> , 250 x 4.6mm, 5μ	Hypersil BDS C <sub>18</sub> , 250 x 4.6mm, 5μ	Hypersil BDS C <sub>18</sub> , 250 x 4.6mm, 5μ	Hypersil BDS C <sub>18</sub> , 250 x 4.6mm, 5μ	
Flow rate  Mobile phase	1 mL/min Acetonitrile: methanol (75:25)%v/v	1 mL/min Acetonitrile: methanol (60:40) % v/v	1 mL/min Acetonitrile: methanol (50:50) % v/v	5mL/min Acetonitrile: methanol (65:35)% v/v	
Detection wavelength	UV, 268 nm	UV, 268 nm	UV, 268 nm	UV, 268 nm	
Injection volume	20μΙ	20μΙ	20μΙ	20μΙ	
Column temperature	Ambient	Ambient	Ambient	Ambient	
Run time and Run mode	Gradient, 10 min	Gradient, 10 min	Gradient, 10 min	Gradient, 10 min	
Conclusion	Peak shape was not good.	Broad peak was obtained.	Slight changes in peak symmetry were observed.	The peak was symmetric	



Table-3: Suitability studies for chrysin

S.NO	Chrysin peak area	Retention	time
1	<u>2202835</u>	3.140	
2	2238912	3.140	
3	2226511	3.409	
4	2225246	3.411	
5	2204540	3.394	
6	2288455	3.412	
AVG	2231083.16	3.392	
SD	31337.8	0.0067	
%RSD	1.4	0.19	

Table-4: Linearity of standard chrysin

Level solution for linearity	Concentration (mg/ml)	Peak area
25%	0.2	592923
50%	0.4	884188
75%	0.8	1195447
100%	1.6	2148712
125%	3.2	3960322
150%	6.4	6905918

Table-5: Specificity

Peak Name	Retention time	Retention ratio
Chrysin	3.3	0.76

Table-6: Method precision

Injection	Peak area	Retention time
1	3.374	636209
2	3.373	679711
3	3.375	671889
4	3.424	673617
5	3.393	684188
6	3.417	675641
AVG	3.3926	676875.8
SD	0.0228	4448.908
%RSD	0.67	0.65



**Table-7: Robustness parameters** 

S.no	Parameter Variations		Area of chrysin	Standard deviation	%RSD
			583538		
		0.8ml	582068	2886.63	0.49
			587638		0.43
1	Flow rate		598999		
1	1 Flow rate	1.2ml	595014	2446.05	0.41
			599464		
			677678		
	2 Wavelength	266nm	6739373	3270.58	0.49
			671157	3270.38	0.43
2		ength 270nm	671421		
۷	wavelength		677858	4175.54	0.62
			679247	41/3.34	0.02

Table-8: Accuracy data for chrysin

Level	Peak area	Found concentration(mg/ml)	%Recovery
	83692	1.012397	101.2
50%	83924	1.971074	98.5
	84176	3.012397	100.4
	2338912	1.028262	102
100%	2202835	1.985743	99
	2626511	2.957501	98
	6207572	1.018297	101
150%	7668229	3.012937	99
	7601714	1.985642	100



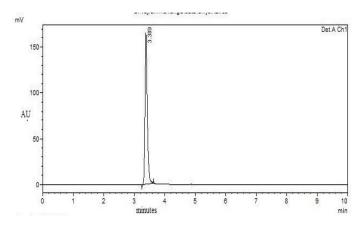


Fig-2: Optimized chromatogram of chrysin



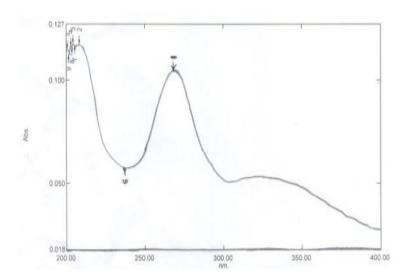


Fig-3: Maximum absorption wavelength ( $\lambda_{max}$ ) of Chrysin- UV Determination

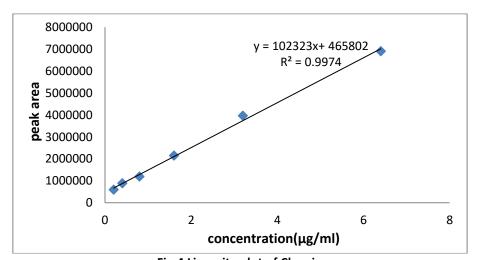


Fig-4 Linearity plot of Chrysin

Regression equation parameters: Slope-102323; Y-intercept-465802; Correlation coefficient: 0.9974

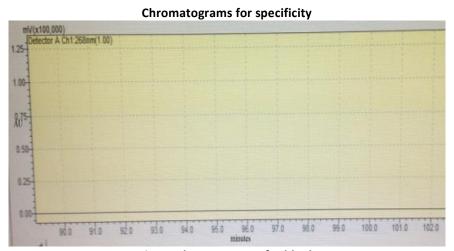


Fig-5: Chromatogram for blank



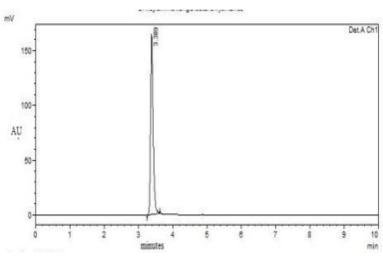


Fig-6: Chromatogram for chrysin (retention time 3.3 min)

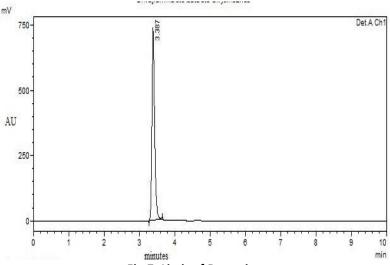
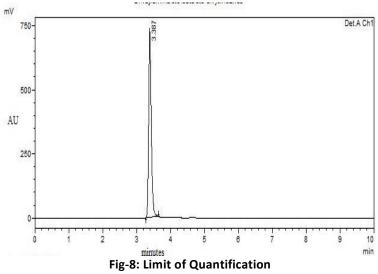


Fig-7: Limit of Detection





#### **ACKNOWLEDMENTS**

We are grateful to DST-CURIE for providing HPLC facility

#### **CONFLICTS**

None

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