

**Original Research Article**

**Formulation Development and Characterization of Darunavir and  
Ritonavir Sustained Release Tablets using Quality by Design  
Approach**

UNDER PEER REVIEW

**Abstract:** *Darunavir is a nonpeptidic inhibitor of protease and is primarily metabolized by cytochrome P450 3A (CYP3A) isoenzymes. It is usually coadministered with low-dose ritonavir (Darunavir/r). Ritonavir is an inhibitor of CYP3A isoenzymes and pharmacologically enhances Darunavir which leads to increased plasma concentrations of darunavir and allows for daily lower dose. Here, we have developed combination SR formulation of Darunavir and Ritonavir and evaluated. In vitro drug release of all formulations was carried out in dissolution medium 900ml of pH 3.0, 0.05 M Sodium Phosphate Buffer + 2% Tween 20 for 75 RPM USP II apparatus (paddle). The results shown that, all the formulations of matrix tablets shown the good release of drug from trialed formulations however all formulations were not releasing the drug in enough amount. In matrix tablets F6, the release of drug shows NLT 80%. So, the formulation F6 have been considered as suitable for the SR tablet of Darunavir and Ritonavir. Tablets were also evaluated though Quality by Design (QbD) method.*

**Keywords:** **Darunavir, Ritonavir, Sustained release, Tablet, Dissolution, Quality by design**

## 1. INTRODUCTION

For each disease condition or the disorder of the patient, appropriate treatment is very important to maintain good health of the patient. For the same, the drug is administered conventionally by one or more of several well defined and popular routes of drug administration which include but not limited to oral, parenteral, rectal, alveolar, ocular and topical etc.<sup>01,15</sup> Nowadays, oral drug delivery system is the preferred way for the administration of drugs because of easy administration, better patient compliance and flexible design of the dosage forms.<sup>02</sup> In recently era, much technical advancement have been done resulting in the development of new techniques for drug delivery. These techniques are capable of controlling the rate of drug delivery, sustaining the duration of therapeutic activity and/or targeting the delivery of drug to a tissue i.e. targeted drug delivery system. These advancements have led to the development of several “Novel Drug Delivery System”.<sup>03</sup> There are several terms used interchangeably viz. controlled release, programmed release, sustained release, prolonged release, timed release, extended release etc. The most important objective for the development of these systems is to furnish an extended duration of action and thus assure greater patient compliance.<sup>06-14</sup>

Sustained release system is a type of modified drug delivery system that can be used as an alternative to conventional drug delivery system. These system sustain the release of drug and maintain the plasma drug concentration in therapeutic window except any fluctuation and increase the therapeutic efficacy of drug.<sup>04</sup> Darunavir is a medicine used to decrease the amount of HIV virus in your body and make your immune system stronger. Darunavir is always used with other HIV medicines.<sup>15</sup> Darunavir inhibits and is primarily metabolized by cytochrome P450 3A (CYP3A) isoenzymes and is coadministered with low-dose ritonavir (Darunavir/r); ritonavir is an inhibitor of CYP3A isoenzymes and pharmacologically enhances Darunavir, resulting in increased plasma concentrations and allowing for a lower daily dose. The t<sub>1/2</sub> (terminal elimination half-life) of Darunavir is 15 h in the presence of ritonavir. An extensive Darunavir/r drug-drug interaction programme has been undertaken, covering a wide range of therapeutic areas.<sup>16,17,18,19,20,21</sup>

It is known that Darunavir is rapidly absorbed from the intestine after oral administration, reaching peak plasma concentrations after 2.5–4.0 h. It is also known that P-glycoprotein expressed in intestinal epithelial cells is able to decrease the absorption of orally administered and low levels of intestinal absorption together with CYP450 activity are major factors in the reduced bioavailability of these drugs. Darunavir, co-administered with ritonavir (both medications are HIV-1 protease inhibitors), is indicated for use in the treatment of HIV-1 infection in combination with other antiretroviral medications.<sup>17,18,21,22,23,24,25,26,27</sup>

The objective of developing oral sustained drug delivery systems of Darunavir in current research study is to avoid other combination treatment to reduce frequency of drug administration, to improve patient compliance, to reduce blood level oscillation characteristic of multiple dosing of conventional dosage forms, to reduce amount of drug administered. The recommended maximum dose for the Darunavir is 800 mg and for ritonavir is 200 mg. Same dose has been selected and formulated for the combination drug. The objective of developing oral combination immediate drug delivery systems of Darunavir and Ritonavir is to facilitate patients with ease of combination treatment

and to evaluate the properties of both drugs which might lead to provide information about other new formulations of these drug.

## 2. DOSE CALCULATION

Ke = Elimination rate constant (Zero-order kinetic), T1/2 = Half-life, DI = Initial Dose,

DM = Maintenance Dose, DL = Loading Dose, Tmax = Time to achieve maximum plasma concentration

$$K_e = 0.693/T_{1/2}$$

$$= 0.693/15$$

$$= 0.0462 \quad R = K_e * DI$$

$$= 0.0462 * 400$$

$$= 18.48/H$$

$$DM = R * T_{max}$$

$$= 18.48 * 20$$

$$= 369.6$$

$$\approx 375 \text{ mg} \quad DL = DI - R * T_{max}$$

$$= 400 - 18.48 * 4$$

$$= 326.08$$

$$\approx 325 \text{ mg}$$

$$\text{Total Dose} = DM + DL$$

$$= 375 + 325 = 700 \text{ mg}$$

Summary: 325 mg of drug should be released in first 4 hrs. Of total 24 hrs. Remaining 375 mg drug should be released in remaining 20 hrs. Of total 24 hrs.

The SR tablet with different methods will be prepared in different batches. Outcome from the data analysis after dissolution test will be analyzed by statistical method comparing.

## 3. MATERIAL AND METHODS

### Materials:

Darunavir, Ritonavir, Microcrystalline Cellulose (Avicel PH-101), Lactose monohydrate (Granulac 200), Hypromellose (METHOCEL™ K100 Premium LVCR), Hypromellose (METHOCEL™ K4M Premium CR), FD & C Green No.40, Magnesium stearate, Opadry White, Purified water.

### Preparation of sustained release tablets:

The powder blends were prepared by taking required quantities of drug and polymer. They were mixed thoroughly. After that microcrystalline cellulose (MCC) was added as directly compressible filler, binder. Finally magnesium stearate was added as a lubricant. These powder blends were then passed through sieve to break any lumps or aggregates. The formulas are indicated in below Table.

### Preparation of sustained release matrix tablets

The powder blends were compressed into tablets by direct compression technique on rotary tableting machine. The compression force was optimized by proper adjustment of upper and lower punches. The tablets formed did not show any defects like capping or chipping. These tablets of each formulation type (F-1 to F-6) were evaluated for various properties such as thickness, diameter, weight variation, uniformity of drug content, hardness, and friability.

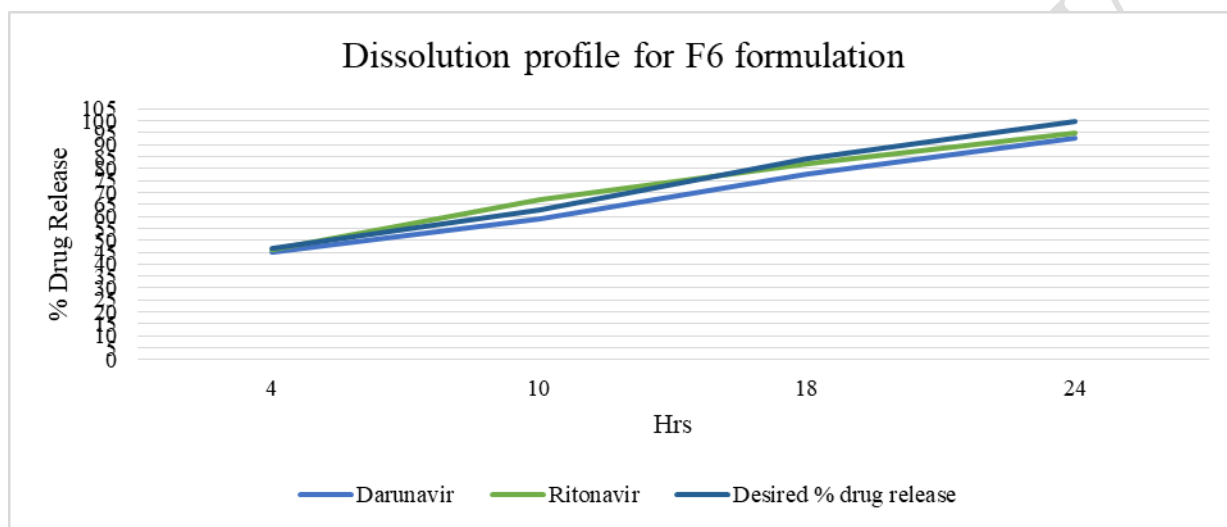
The result of excipients compatibility study is presented in table below.

**Table 1: Formulations and dissolution profile for sustained release Darunavir and Ritonavir tablet**

	F1		F2		F3		F4		F5		F6	
Ingredients	Quantity Required (mg/Tablet)		Quantity Required (mg/Tablet)		Quantity Required (mg/Tablet)		Quantity Required (mg/Tablet)		Quantity Required (mg/Tablet)		Quantity Required (mg/Tablet)	
Darunavir	375.00		375.00		375.00		375.00		375.00		375.00	
Ritonavir	110.00		110.00		110.00		110.00		110.00		110.00	
Microcrystalline Cellulose (Avicel PH-112)	52.00		42.00		36.00		31.00		27.50		22.00	
Lactose monohydrate (Granulac 200)	52.00		42.00		36.00		31.00		27.50		22.00	
Hypromellose (METHOCEL™ K100 Premium LVCR)	100.00		120.00		130.00		138.00		142.00		150.00	
Hypromellose (METHOCEL™ K4M Premium CR)	25.00		25.00		27.00		29.00		32.00		35.00	
FD & C Green No.40	1.00		1.00		1.00		1.00		1.00		1.00	
Magnesium Stearate	5.00		5.00		5.00		5.00		5.00		5.00	
Opadry White (03B28796)	20.00		20.00		20.00		20.00		20.00		20.00	
Purified Water	q. s.		q. s.		q. s.		q. s.		q. s.		q. s.	
<b>Total Weight</b>	740.00		740.00		740.00		740.00		740.00		740.00	
<b>Dissolution profile</b>												
<b>%Drug Release</b>												
Time Points	F1		F2		F3		F4		F5		F6	
	D	R	D	R	D	R	D	R	D	R	D	R
<b>4 hr (NMT 50 %)</b>	75	78	68	75	64	69	60	63	50	56	45	46
<b>10 hr (55-75)</b>	85	88	79	83	74	79	70	74	68	72	59	67

	F1		F2		F3		F4		F5		F6	
Ingredients	Quantity Required (mg/Tablet)		Quantity Required (mg/Tablet)		Quantity Required (mg/Tablet)		Quantity Required (mg/Tablet)		Quantity Required (mg/Tablet)		Quantity Required (mg/Tablet)	
18 hr (70-90)	92	97	89	90	86	87	84	85	82	86	78	82
24 hr (NLT 90)	99	99	97	98	95	97	94	96	94	96	93	95

D = Darunavir; R = Ritonavir



**Graph 1: Dissolution profile for F6 formulation**

#### Quality Target Product Profile for the Antiretroviral Sustained Release Tablets

The Quality Target Product Profile (QTPP) is “a prospective summary of the quality characteristics of a drug product that ideally will be achieved to ensure the desired quality, taking into account safety and efficacy of the drug product.” The QTPP is an essential element of a QbD approach and forms the basis of design for the development of the product. For SR Tablets, the target should be defined early in development based on the properties of the drug substance (DS), characterization of the product. By beginning with the end in mind, the result of development is a robust formulation and manufacturing process with an acceptable control strategy that ensures the performance of the drug product.

A critical quality attribute (CQA) is “a physical, chemical, biological, or microbiological property or characteristic that should be within an appropriate limit, range, or distribution to ensure the desired product quality.”<sup>1</sup> The identification of a CQA from the QTPP is based on the severity of harm to a patient should the product fall outside the acceptable range for that attribute.

All quality attributes are target elements of the drug product and should be achieved through a good quality management system, appropriate formulation/process design and development. From the perspective of pharmaceutical development, we only investigate the subset of CQAs of the drug product that also have a high potential to be impacted by the formulation or process variables. Our investigation culminates in an appropriate control strategy.

#### 4. QUALITY BY DESIGN

**Study Plan:** A 2<sup>3</sup> full Factorial design was used and three center points was included to evaluate any curvature effects exist. Batch size of 700 units was executed at lab scale as per different combination of factor as per DOE plan. All Processing parameter like granulation, wet milling, drying, milling, blending and compression are kept constant to reduce additional noise. All factors are numeric factor and the drug product CQA to be evaluated is **% Drug released at 24 hrs. For both API**. Study design is given in **Table**. Formulation composition and experimental results for drug release profile for all designed experiments are given.

**Power Calculation for design:** For selected design power also found to be above 80% for all three factors. Data of signal/Noise ratio and power obtained for the selected design at. 5% alpha level to detect specified signal to noise ratio are presented in below Table.

**Table 2: Design of the Regular 2<sup>3</sup> Factorial DOE with 1 Centre Points to Study Impact of Critical Excipients**

Factors : Formulation Variables		Levels		
		-1	+1	Center point level
A	Hypromellose (METHOCELTM K100 Premium LVCR)	142.50 mg/tab	157.50 mg/tab	150 mg/tab
B	Hypromellose (METHOCELTM K4M Premium CR)	33.25 mg/ tab	36.55 mg/ tab	35 mg/ tab
C	Magnesium Stearate	4.0 mg/tab	5.13 mg/tab	5 mg/tab
Response		Goal	Acceptance Range	
Y1	Dissolution of Darunavir at 4 Hrs. (%)	Minimum	NMT 50%	
Y2	Dissolution of Darunavir at 18Hrs. (%)	Minimum	70- 90%	
Y3	Dissolution of Ritonavir at 4 Hrs. (%)	Minimum	NMT 50%	
Y4	Dissolution of Ritonavir at 18Hrs. (%)	Minimum	70- 90%	

**Power Calculation for design:** For selected design power also found to be above 90 % for all three factors. Data of signal/Noise ratio and power obtained for the selected design at. 5% alpha level to detect specified signal to noise ratio are presented in below Table.

**Table 3: Design Power Evaluation**

Name	Unit	Difference to detect delta (Signal)	Est. Std. Deviation Sigma (Noise)	Delta/Sigma (Signal/Noise ratio)	Power
Dissolution of Darunavir at 4 Hrs. (%)	%	2	0.5	4	99.8
Dissolution of Darunavir at 18Hrs. (%)	%	2	0.5	4	99.8
Dissolution of Ritonavir at 4 Hrs. (%)	%	2	0.5	4	99.8

Dissolution of Ritonavir at 18Hrs. (%)	%	2	0.5	4	99.8
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**Table 5: Experimental Results of the DOE to study Drug release**

Batch No.	Factors : Formulation Variables			Response			
	Factor 1 A: Controlled Release Polymer Hypromellose (METHOCEL™ K100 Premium LVCR)	Factor 2 B: Controlled Release Polymer Hypromellose (METHOCEL™ K4M Premium CR)	Factor 3 C: Level of Lubrication Magnesium Stearate	Dissolution (%)			
				4 Hrs. NMT 50%		18Hrs. 70- 90%	
				Darunavir R1	Ritonavir R2	Darunavir R1	Ritonavir R2
DOE-1	157.50	36.57	5.13	42	46	72	78
DOE-2	150.00	35.00	5.00	43	48	79	82
DOE-3	157.50	36.57	4.88	43	46	73	78
DOE-4	142.50	36.57	4.88	46	48	81	87
DOE-5	142.50	33.25	4.88	46	49	82	88
DOE-6	150.00	35.00	5.00	44	48	77	81
DOE-7	142.50	33.25	5.13	45	48	81	86
DOE-8	142.50	36.75	5.13	44	48	79	84
DOE-9	157.50	33.25	4.88	42	47	77	80
DOE-10	150.00	35.00	5.00	42	47	78	83
DOE-11	157.50	33.25	5.13	42	46	78	80

**ANOVA for selected factorial model**

**Response 1: Darunavir-Dissolution 4 hr (NMT 50%)**

Source	Sum of Squares	df	Mean Square	F-value	p-value	
<b>Model</b>	18.88	7	2.70	2.20	0.2771	not significant
A-Hypromellose(Methocel K 100)	3.12	1	3.12	2.55	0.2083	
B-Hypromellose(Methocel K4M)	0.1250	1	0.1250	0.1022	0.7702	
C-Magnesium Stearate	1.12	1	1.12	0.9195	0.4083	
AB	1.13	1	1.13	0.9195	0.4083	
AC	10.12	1	10.12	8.28	0.0637	
BC	0.1250	1	0.1250	0.1022	0.7702	
ABC	3.12	1	3.12	2.55	0.2083	
<b>Residual</b>	3.67	3	1.22			
Lack of Fit	1.67	1	1.67	1.67	0.3254	not significant
Pure Error	2.00	2	1.0000			
<b>Cor Total</b>	22.55	10				

Factor coding is **Coded**.

Sum of squares is **Type III - Partial**

The **Model F-value** of 2.20 implies the model is not significant relative to the noise. There is a 27.71% chance that an F-value this large could occur due to noise.

**P-values** less than 0.0500 indicate model terms are significant. In this case there are no significant model terms. Values greater than 0.1000 indicate the model terms are not significant. If there are many insignificant model terms (not counting those required to support hierarchy), model reduction may improve your model.

The **Lack of Fit F-value** of 1.67 implies the Lack of Fit is not significant relative to the pure error. There is a 32.54% chance that a Lack of Fit F-value this large could occur due to noise. Non-significant lack of fit is good -- we want the model to fit.

**Fit Statistics**

**Std. Dev.** 1.11 **R<sup>2</sup>** 0.8372

**Mean** 43.64 **Adjusted R<sup>2</sup>** 0.4573

**C.V. %** 2.53 **Predicted R<sup>2</sup>** -16.5596

**Adeq Precision** 4.2405

A negative **Predicted R<sup>2</sup>** implies that the overall mean may be a better predictor of your response than the current model. In some cases, a higher order model may also predict better.

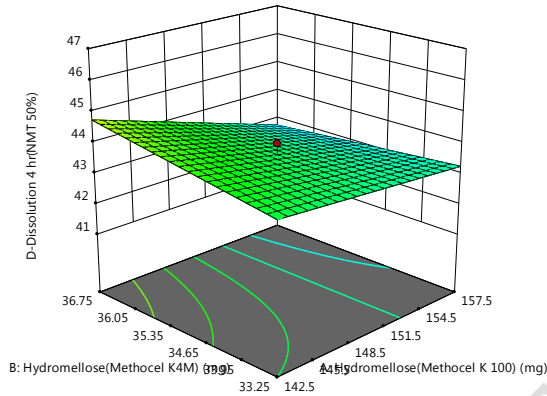
**Adeq Precision** measures the signal to noise ratio. A ratio greater than 4 is desirable. Your ratio of 4.240 indicates an adequate signal. This model can be used to navigate the design space.

Design-Expert® Software  
Factor Coding: Actual

D-Dissolution 4 hr(NMT 50%)  
● Design points above predicted value  
○ Design points below predicted value  
42 46

X1 = A: Hydromellose(Methocel K 100)  
X2 = B: Hydromellose(Methocel K4M)

Actual Factor  
C: Magnesium Stearate = 5



Graph 2: Dissolution 4 hrs (factorial design)

Response 2: Darunavir-Dissolution 18 hr (70-90 %)

Source	Sum of Squares	df	Mean Square	F-value	p-value	
<b>Model</b>	57.88	7	8.27	8.70	0.0515	not significant
A-Hypromellose(Methocel K 100)	15.12	1	15.12	15.91	0.0282	
B-Hypromellose(Methocel K4M)	6.12	1	6.12	6.44	0.0848	
C-Magnesium Stearate	1.13	1	1.13	1.18	0.3563	
AB	6.13	1	6.13	6.44	0.0848	
AC	28.12	1	28.12	29.58	0.0122	
BC	1.12	1	1.12	1.18	0.3563	
ABC	0.1250	1	0.1250	0.1315	0.7409	
<b>Residual</b>	2.85	3	0.9508			
Lack of Fit	0.8523	1	0.8523	0.8523	0.4534	not significant
Pure Error	2.00	2	1.0000			
<b>Cor Total</b>	60.73	10				

Factor coding is **Coded**.

Sum of squares is **Type III - Partial**

The **Model F-value** of 8.70 implies there is a 5.15% chance that an F-value this large could occur due to noise.

**P-values** less than 0.0500 indicate model terms are significant. In this case A, AC are significant model terms. Values greater than 0.1000 indicate the model terms are not significant. If there are many insignificant model terms (not counting those required to support hierarchy), model reduction may improve your model.

The **Lack of Fit F-value** of 0.85 implies the Lack of Fit is not significant relative to the pure error. There is a 45.34% chance that a Lack of Fit F-value this large could occur due to noise. Non-significant lack of fit is good -- we want the model to fit.

**Fit Statistics**

**Std. Dev.** 0.9751 **R<sup>2</sup>** 0.9530  
**Mean** 78.45 **Adjusted R<sup>2</sup>** 0.8434  
**C.V. %** 1.24 **Predicted R<sup>2</sup>** -2.3456

**Adeq Precision** 10.8233

A negative **Predicted R<sup>2</sup>** implies that the overall mean may be a better predictor of your response than the current model. In some cases, a higher order model may also predict better.

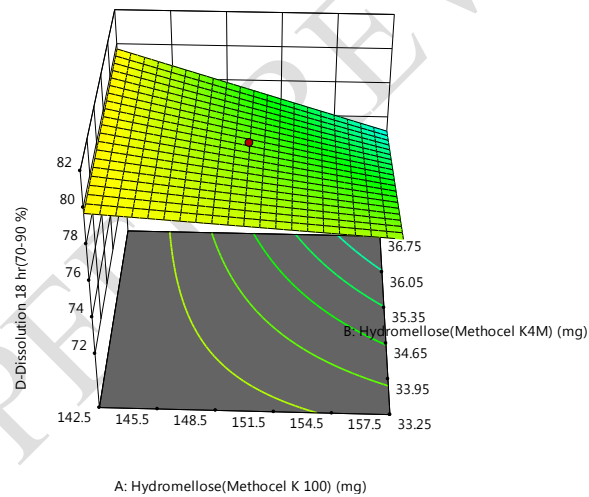
**Adeq Precision** measures the signal to noise ratio. A ratio greater than 4 is desirable. Your ratio of 10.823 indicates an adequate signal. This model can be used to navigate the design space.

Design-Expert® Software  
 Factor Coding: Actual

**D-Dissolution 18 hr(70-90 %)**  
 ● Design points above predicted value  
 ○ Design points below predicted value  
 73 82

X1 = A: Hydromellose(Methocel K 100)  
 X2 = B: Hydromellose(Methocel K4M)

**Actual Factor**  
 C: Magnesium Stearate = 5



**Graph 3: Dissolution 18 hrs (factorial design)**

**ANOVA for selected factorial model**

**Response 3: Ritonavir -Dissolution 4 hr(NMT 50%)**

Source	Sum of Squares	df	Mean Square	F-value	p-value	
<b>Model</b>	7.88	7	1.13	3.96	0.1428	not significant
A-Hypromellose(Methocel K 100)	1.13	1	1.13	3.96	0.1407	
B-Hypromellose(Methocel K4M)	0.1250	1	0.1250	0.4400	0.5545	
C-Magnesium Stearate	0.1250	1	0.1250	0.4400	0.5545	
AB	1.12	1	1.12	3.96	0.1407	
AC	3.13	1	3.13	11.00	0.0452	
BC	1.13	1	1.13	3.96	0.1407	
ABC	1.13	1	1.13	3.96	0.1407	
<b>Residual</b>	0.8523	3	0.2841			
Lack of Fit	0.1856	1	0.1856	0.5568	0.5333	not

significant

Pure Error 0.6667 2 0.3333  
**Cor Total** 8.73 10

Factor coding is **Coded**.

Sum of squares is **Type III - Partial**

The **Model F-value** of 3.96 implies the model is not significant relative to the noise. There is a 14.28% chance that an F-value this large could occur due to noise.

**P-values** less than 0.0500 indicate model terms are significant. In this case AC is a significant model term. Values greater than 0.1000 indicate the model terms are not significant. If there are many insignificant model terms (not counting those required to support hierarchy), model reduction may improve your model.

The **Lack of Fit F-value** of 0.56 implies the Lack of Fit is not significant relative to the pure error. There is a 53.33% chance that a Lack of Fit F-value this large could occur due to noise. Non-significant lack of fit is good -- we want the model to fit.

**Fit Statistics**

**Std. Dev.** 0.5330 **R<sup>2</sup>** 0.9023  
**Mean** 47.45 **Adjusted R<sup>2</sup>** 0.6745  
**C.V. %** 1.12 **Predicted R<sup>2</sup>** -4.1019  
**Adeq Precision** 6.6000

A negative **Predicted R<sup>2</sup>** implies that the overall mean may be a better predictor of your response than the current model. In some cases, a higher order model may also predict better.

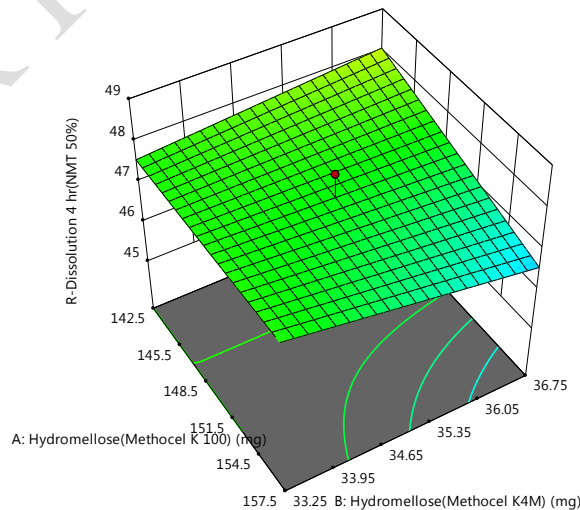
**Adeq Precision** measures the signal to noise ratio. A ratio greater than 4 is desirable. Your ratio of 6.600 indicates an adequate signal. This model can be used to navigate the design space.

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 Factor Coding: Actual

**R-Dissolution 4 hr(NMT 50%)**  
 ● Design points above predicted value  
 ○ Design points below predicted value  
 46 49

X1 = A: Hydromellose(Methocel K 100)  
 X2 = B: Hydromellose(Methocel K4M)

**Actual Factor**  
 C: Magnesium Stearate = 5



**Graph 4: Dissolution 4 hrs (factorial design)**

**ANOVA for selected factorial model**

\*\*Response 4: Ritonavir -Dissolution 18 hr (70-90 %) \*\*

Source	Sum of Squares	df	Mean Square	F-value	p-value
<b>Model</b>	94.87	7	13.55	8.54	0.0528 not

A-Hypromellose(Methocel K 100)	21.13	1	21.13	13.31	0.0355		significant
B-Hypromellose(Methocel K4M)	1.13	1	1.13	0.7088	0.4617		
C-Magnesium Stearate	0.1250	1	0.1250	0.0788	0.7972		
AB	10.12	1	10.12	6.38	0.0857		
AC	55.13	1	55.13	34.73	0.0098		
BC	1.13	1	1.13	0.7088	0.4617		
ABC	6.13	1	6.13	3.86	0.1442		
<b>Residual</b>	4.76	3	1.59				
Lack of Fit	2.76	1	2.76	2.76	0.2385		not significant
Pure Error	2.00	2	1.0000				
<b>Cor Total</b>	99.64	10					

Factor coding is **coded**.

Sum of squares is **Type III - Partial**

The **Model F-value** of 8.54 implies there is a 5.28% chance that an F-value this large could occur due to noise.

**P-values** less than 0.0500 indicate model terms are significant. In this case A, AC are significant model terms. Values greater than 0.1000 indicate the model terms are not significant. If there are many insignificant model terms (not counting those required to support hierarchy), model reduction may improve your model.

The **Lack of Fit F-value** of 2.76 implies the Lack of Fit is not significant relative to the pure error. There is a 23.85% chance that a Lack of Fit F-value this large could occur due to noise. Non-significant lack of fit is good -- we want the model to fit.

#### Fit Statistics

<b>Std. Dev.</b>	1.26	<b>R<sup>2</sup></b>	0.9522
<b>Mean</b>	82.82	<b>Adjusted R<sup>2</sup></b>	0.8407
<b>C.V. %</b>	1.52	<b>Predicted R<sup>2</sup></b>	-5.5523
		<b>Adeq Precision</b>	9.3078

A negative **Predicted R<sup>2</sup>** implies that the overall mean may be a better predictor of your response than the current model. In some cases, a higher order model may also predict better.

**Adeq Precision** measures the signal to noise ratio. A ratio greater than 4 is desirable. Your ratio of 9.308 indicates an adequate signal. This model can be used to navigate the design space.

Design-Expert® Software  
Factor Coding: Actual

R-Dissolution 18 hr(70-90 %)

● Design points above predicted value

○ Design points below predicted value

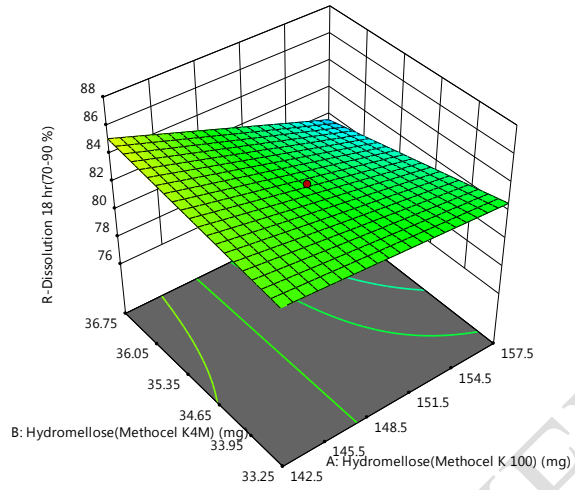
78  88

X1 = A: Hydromellose(Methocel K 100)

X2 = B: Hydromellose(Methocel K4M)

Actual Factor

C: Magnesium Stearate = 5



Graph 5: Dissolution 18 hrs (factorial design)

UNDER PEER REVIEW

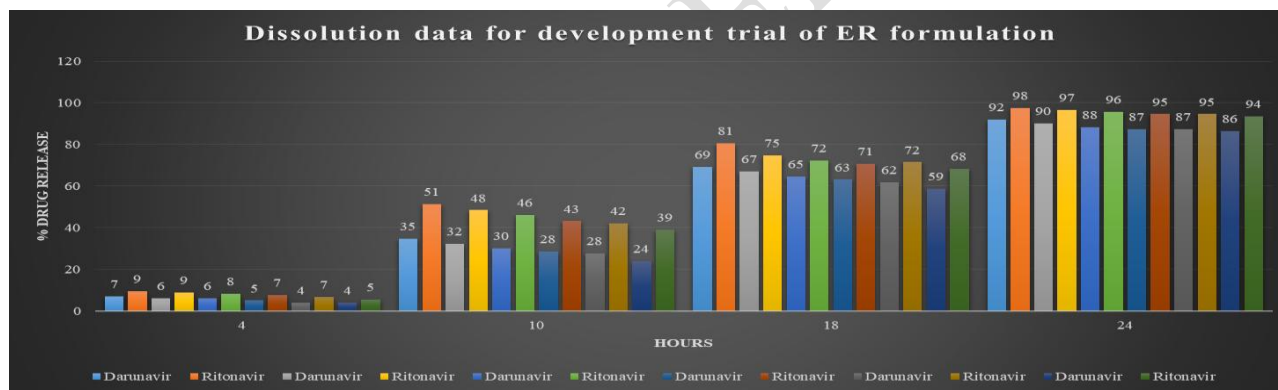
**Table 6: Development Trial of Darunavir and Ritonavir Extended release tablet**

Ingredients		D1		D2		D3		D4		D5		D6	
DRY MIXING		Qty mg/Tab	%W/W	Qty mg/Tab	%W/W	Qty mg/Tab	%W/W	Qty mg/Tab	%W/W	Qty mg/Tab	%W/W	Qty mg/Tab	%W/W
1	Darunavir	375.00	50.68	375.00	50.68	375.00	50.68	375.00	50.68	375.00	50.68	375.00	50.68
2	Ritonavir	110.00	14.86	110.00	14.86	110.00	14.86	110.00	14.86	110.00	14.86	110.00	14.86
3	Microcrystalline Cellulose (Avicel PH 101)	52.00	7.03	42.00	5.68	36.00	4.86	31.00	4.19	27.50	3.72	22.00	2.97
4	Lactose monohydrate (Granulac 200)	52.00	7.03	42.00	5.68	36.00	4.86	31.00	4.19	27.50	3.72	22.00	2.97
5	Hypromellose (METHOCEL™ K100 Premium LVCR)	100.00	13.51	120.00	16.22	130.00	17.57	138.00	18.65	142.00	19.19	150.00	20.27
6	Hypromellose (METHOCEL™ K4M Premium CR)	25.00	3.38	25.00	3.38	27.00	3.65	29.00	3.92	32.00	4.32	35.00	4.73
7	FD & C Green No.40	1.00	0.14	1.00	0.14	1.00	0.14	1.00	0.14	1.00	0.14	1.00	0.14
<b>GRANULATION:</b>													
8	Purified Water	q.s	---	q.s	---	q.s	---	q.s	---	q.s	---	q.s	---
<b>LUBRICATION:</b>													
10	Magnesium Stearate	5.00	0.68	5.00	0.68	5.00	0.68	5.00	0.68	5.00	0.68	5.00	0.68
<b>Total</b>		<b>720.00</b>		<b>720.00</b>	-	<b>720.00</b>		-	<b>720.00</b>	-	<b>720.00</b>	-	<b>720.00</b>
<b>FILM COATING:</b>													
11	Opadry White (03B28796)	20.00	2.70	20.00	2.70	20.00	2.70	20.00	2.70	20.00	2.70	20.00	2.70
12	Purified Water <sup>#</sup>	q.s		q.s	--	q.s	--	q.s	--	q.s	--	q.s	--
<b>Total</b>		<b>740.00</b>		<b>740.00</b>	100.00	<b>740.00</b>	100.00	<b>740.00</b>	100.00	<b>740.00</b>	100.00	<b>740.00</b>	100.00
D = Development trial													

**Table 7: Extended Release Dissolution profile in matrix formulation**

900mL pH 3.0 0.05 M Sodium Phosphate Buffer + 2% Tween 20 , 75 RPM												
Batch No.	D1		D2		D3		D4		D5		D6	
	% Release		% Release		% Release		% Release		% Release		% Release	
Time (hr)	Darunavir	Ritonavir	Darunavir	Ritonavir	Darunavir	Darunavir	Darunavir	Ritonavir	Darunavir	Ritonavir	Darunavir	Ritonavir
4 (NMT 10%)	7	9	6	9	6	8	5	7	4	7	4	5
10 (20 -45 %)	35	51	32	48	30	46	28	43	28	42	24	39
18 (55-75 %)	69	81	67	75	65	72	63	71	62	72	59	68
24 (NLT 80 %)	92	98	90	97	88	96	87	95	87	95	86	94

D = Development trial

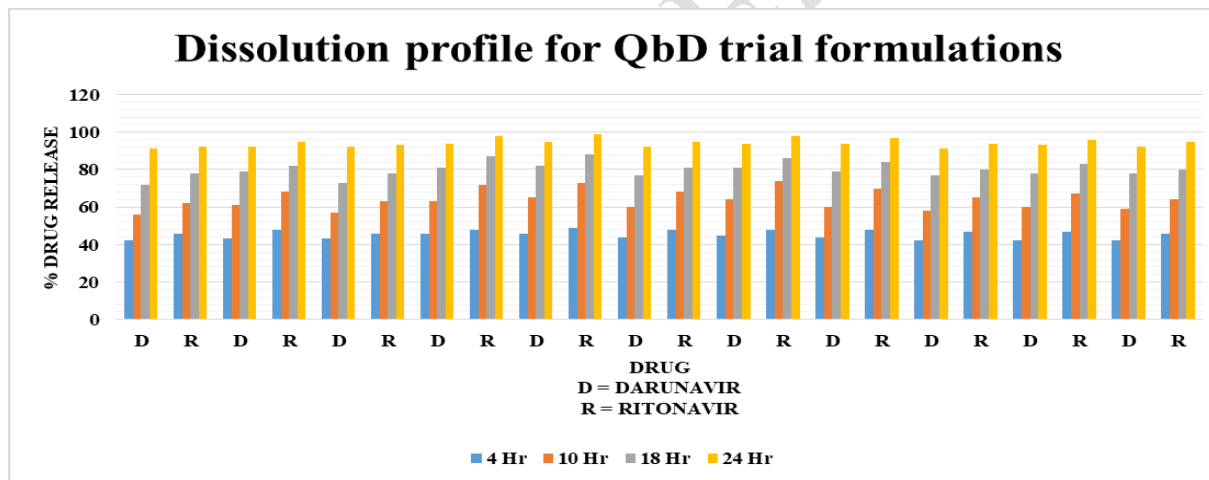


**Graph 6: Dissolution data for development trial of ER formulation**

Table 8: QbD data for Extended Release profile in matrix formulation

Time point	Acceptable range	F1		F2		F3		F4		F5		F6		F7		F8		F9		F10		F11	
		% Drug Release		% Drug Release		% Drug Release		% Drug Release		% Drug Release		% Drug Release		% Drug Release		% Drug Release		% Drug Release		% Drug Release		% Drug Release	
		D	R	D	R	D	R	D	R	D	R	D	R	D	R	D	R	D	R	D	R	D	R
4 hrs	NMT 50 %	42	46	43	48	43	46	46	48	46	49	44	48	45	48	44	48	42	47	42	47	42	46
10 hrs	55 – 75 %	56	62	61	68	57	63	63	72	65	73	60	68	64	74	60	70	58	65	60	67	59	64
18 hrs	70 – 90 %	72	78	79	82	73	78	81	87	82	88	77	81	81	86	79	84	77	80	78	83	78	80
24 hrs	NLT 90 %	91	92	92	95	92	93	94	98	95	99	92	95	94	98	94	97	91	94	93	96	92	95

D = Darunavir, R = Ritonavir



Graph 7: Dissolution profile for QbD trial formulation

## ANOVA for selected factorial model

### Response 1: D-Dissolution 4 hr(NMT 50%)

Source	Sum of Squares	df	Mean Square	F-value	p-value	
<b>Model</b>	18.88	7	2.70	2.20	0.2771	not significant
A-Hydromellose(Methocel K 100)	3.12	1	3.12	2.55	0.2083	
B-Hydromellose(Methocel K4M)	0.1250	1	0.1250	0.1022	0.7702	
C-Magnesium Stearate	1.12	1	1.12	0.9195	0.4083	
AB	1.13	1	1.13	0.9195	0.4083	
AC	10.12	1	10.12	8.28	0.0637	
BC	0.1250	1	0.1250	0.1022	0.7702	
ABC	3.12	1	3.12	2.55	0.2083	
<b>Residual</b>	3.67	3	1.22			
Lack of Fit	1.67	1	1.67	1.67	0.3254	not significant
Pure Error	2.00	2	1.0000			
<b>Cor Total</b>	22.55	10				

Factor coding is **Coded**.

Sum of squares is **Type III - Partial**

The **Model F-value** of 2.20 implies the model is not significant relative to the noise. There is a 27.71% chance that an F-value this large could occur due to noise.

**P-values** less than 0.0500 indicate model terms are significant. In this case there are no significant model terms. Values greater than 0.1000 indicate the model terms are not significant. If there are many insignificant model terms (not counting those required to support hierarchy), model reduction may improve your model.

The **Lack of Fit F-value** of 1.67 implies the Lack of Fit is not significant relative to the pure error. There is a 32.54% chance that a Lack of Fit F-value this large could occur due to noise. Non-significant lack of fit is good -- we want the model to fit.

### Fit Statistics

<b>Std. Dev.</b>	1.11	<b>R<sup>2</sup></b>	0.8372
<b>Mean</b>	43.64	<b>Adjusted R<sup>2</sup></b>	0.4573
<b>C.V. %</b>	2.53	<b>Predicted R<sup>2</sup></b>	-16.5596
		<b>Adeq Precision</b>	4.2405

A negative **Predicted R<sup>2</sup>** implies that the overall mean may be a better predictor of your response than the current model. In some cases, a higher order model may also predict better.

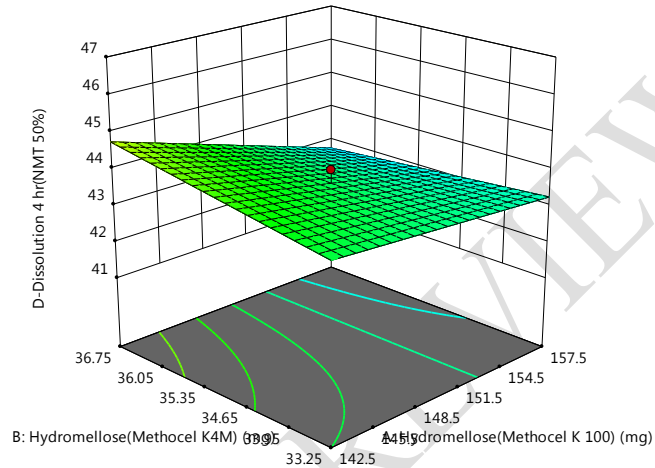
**Adeq Precision** measures the signal to noise ratio. A ratio greater than 4 is desirable. Your ratio of 4.240 indicates an adequate signal. This model can be used to navigate the design space.

Design-Expert® Software  
Factor Coding: Actual

**D-Dissolution 4 hr(NMT 50%)**  
● Design points above predicted value  
○ Design points below predicted value  
42 46

X1 = A: Hydromellose(Methocel K 100)  
X2 = B: Hydromellose(Methocel K4M)

**Actual Factor**  
C: Magnesium Stearate = 5



**Graph 8: Dissolution 4 hrs (factorial design)**

**ANOVA for selected factorial model**

**Response 2: D-Dissolution 18 hr(70-90 %)**

Source	Sum of Squares	df	Mean Square	F-value	p-value
<b>Model</b>	57.88	7	8.27	8.70	0.0515 not significant
A-Hydromellose(Methocel K 100)	15.12	1	15.12	15.91	0.0282
B-Hydromellose(Methocel K4M)	6.12	1	6.12	6.44	0.0848
C-Magnesium Stearate	1.13	1	1.13	1.18	0.3563
AB	6.13	1	6.13	6.44	0.0848
AC	28.12	1	28.12	29.58	0.0122
BC	1.12	1	1.12	1.18	0.3563
ABC	0.1250	1	0.1250	0.1315	0.7409
<b>Residual</b>	2.85	3	0.9508		
Lack of Fit	0.8523	1	0.8523	0.8523	0.4534 not significant
Pure Error	2.00	2	1.0000		
<b>Cor Total</b>	60.73	10			

Factor coding is **Coded**.

Sum of squares is **Type III - Partial**

The **Model F-value** of 8.70 implies there is a 5.15% chance that an F-value this large could occur due to noise.

**P-values** less than 0.0500 indicate model terms are significant. In this case A, AC are significant model terms. Values greater than 0.1000 indicate the model terms are not significant. If there are many insignificant model terms (not counting those required to support hierarchy), model reduction may improve your model.

The **Lack of Fit F-value** of 0.85 implies the Lack of Fit is not significant relative to the pure error. There is a 45.34% chance that a Lack of Fit F-value this large could occur due to noise. Non-significant lack of fit is good -- we want the model to fit.

### Fit Statistics

<b>Std. Dev.</b>	0.9751	<b>R<sup>2</sup></b>	0.9530
<b>Mean</b>	78.45	<b>Adjusted R<sup>2</sup></b>	0.8434
<b>C.V. %</b>	1.24	<b>Predicted R<sup>2</sup></b>	-2.3456
		<b>Adeq Precision</b>	10.8233

A negative **Predicted R<sup>2</sup>** implies that the overall mean may be a better predictor of your response than the current model. In some cases, a higher order model may also predict better.

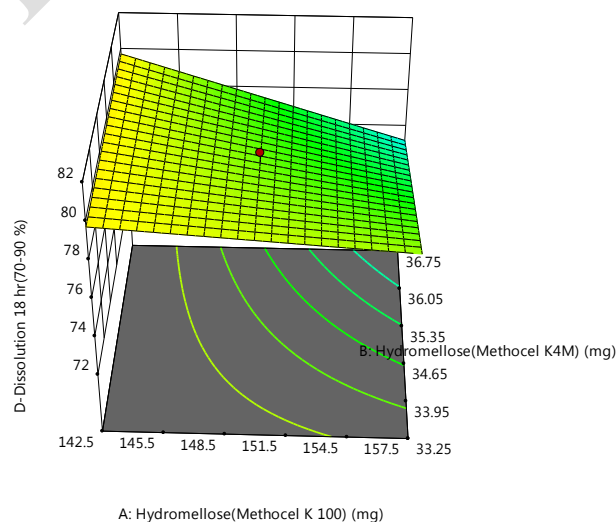
**Adeq Precision** measures the signal to noise ratio. A ratio greater than 4 is desirable. Your ratio of 10.823 indicates an adequate signal. This model can be used to navigate the design space.

Design-Expert® Software  
Factor Coding: Actual

**D-Dissolution 18 hr(70-90 %)**  
 ● Design points above predicted value  
 ○ Design points below predicted value  
 73 82

X1 = A: Hydromellose(Methocel K 100)  
 X2 = B: Hydromellose(Methocel K4M)

**Actual Factor**  
 C: Magnesium Stearate = 5



**Graph 9: Dissolution 18 hrs (factorial design)**

**ANOVA for selected factorial model**

**Response 3: R-Dissolution 4 hr(NMT 50%)**

Source	Sum of Squares	df	Mean Square	F-value	p-value
<b>Model</b>	7.88	7	1.13	3.96	0.1428 <sup>not significant</sup>
A-Hydromellose(Methocel K 100)	1.13	1	1.13	3.96	0.1407
B-Hydromellose(Methocel K4M)	0.1250	1	0.1250	0.4400	0.5545
C-Magnesium Stearate	0.1250	1	0.1250	0.4400	0.5545
AB	1.12	1	1.12	3.96	0.1407
AC	3.13	1	3.13	11.00	0.0452
BC	1.13	1	1.13	3.96	0.1407
ABC	1.13	1	1.13	3.96	0.1407
<b>Residual</b>	0.8523	3	0.2841		
Lack of Fit	0.1856	1	0.1856	0.5568	0.5333 <sup>not significant</sup>
Pure Error	0.6667	2	0.3333		
<b>Cor Total</b>	8.73	10			

Factor coding is **Coded**.

Sum of squares is **Type III - Partial**

The **Model F-value** of 3.96 implies the model is not significant relative to the noise. There is a 14.28% chance that an F-value this large could occur due to noise.

**P-values** less than 0.0500 indicate model terms are significant. In this case AC is a significant model term. Values greater than 0.1000 indicate the model terms are not significant. If there are many insignificant model terms (not counting those required to support hierarchy), model reduction may improve your model.

The **Lack of Fit F-value** of 0.56 implies the Lack of Fit is not significant relative to the pure error. There is a 53.33% chance that a Lack of Fit F-value this large could occur due to noise. Non-significant lack of fit is good -- we want the model to fit.

**Fit Statistics**

<b>Std. Dev.</b>	0.5330	<b>R<sup>2</sup></b>	0.9023
<b>Mean</b>	47.45	<b>Adjusted R<sup>2</sup></b>	0.6745
<b>C.V. %</b>	1.12	<b>Predicted R<sup>2</sup></b>	-4.1019
		<b>Adeq Precision</b>	6.6000

A negative **Predicted R<sup>2</sup>** implies that the overall mean may be a better predictor of your response than the current model. In some cases, a higher order model may also predict better.

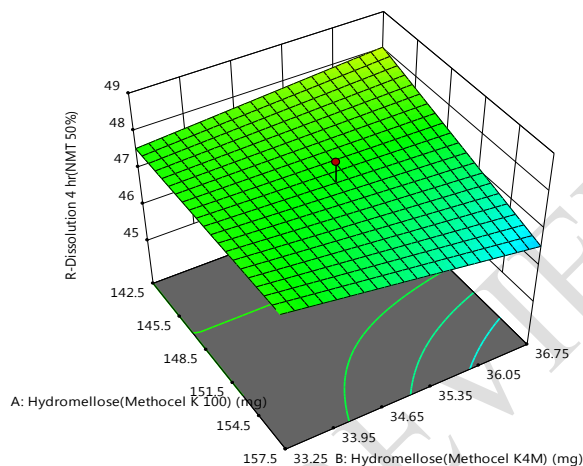
**Adeq Precision** measures the signal to noise ratio. A ratio greater than 4 is desirable. Your ratio of 6.600 indicates an adequate signal. This model can be used to navigate the design space.

Design-Expert® Software  
Factor Coding: Actual

R-Dissolution 4 hr(NMT 50%)  
● Design points above predicted value  
○ Design points below predicted value  
46 49

X1 = A: Hydromellose(Methocel K 100)  
X2 = B: Hydromellose(Methocel K4M)

Actual Factor  
C: Magnesium Stearate = 5



Graph 10: Dissolution 4 hrs (factorial design)

#### ANOVA for selected factorial model

\*\*Response 4: R-Dissolution 18 hr(70-90 %) \*\*

Source	Sum of Squares	df	Mean Square	F-value	p-value	
<b>Model</b>	94.87	7	13.55	8.54	0.0528	not significant
A-Hydromellose(Methocel K 100)	21.13	1	21.13	13.31	0.0355	
B-Hydromellose(Methocel K4M)	1.13	1	1.13	0.7088	0.4617	
C-Magnesium Stearate	0.1250	1	0.1250	0.0788	0.7972	
AB	10.12	1	10.12	6.38	0.0857	
AC	55.13	1	55.13	34.73	0.0098	
BC	1.13	1	1.13	0.7088	0.4617	
ABC	6.13	1	6.13	3.86	0.1442	
<b>Residual</b>	4.76	3	1.59			
Lack of Fit	2.76	1	2.76	2.76	0.2385	not significant
Pure Error	2.00	2	1.0000			
<b>Cor Total</b>	99.64	10				

Factor coding is **Coded**.

Sum of squares is **Type III - Partial**

The **Model F-value** of 8.54 implies there is a 5.28% chance that an F-value this large could occur due to noise.

**P-values** less than 0.0500 indicate model terms are significant. In this case A, AC are significant model terms. Values greater than 0.1000 indicate the model terms are not significant. If there are many insignificant model terms (not counting those required to support hierarchy), model reduction may improve your model.

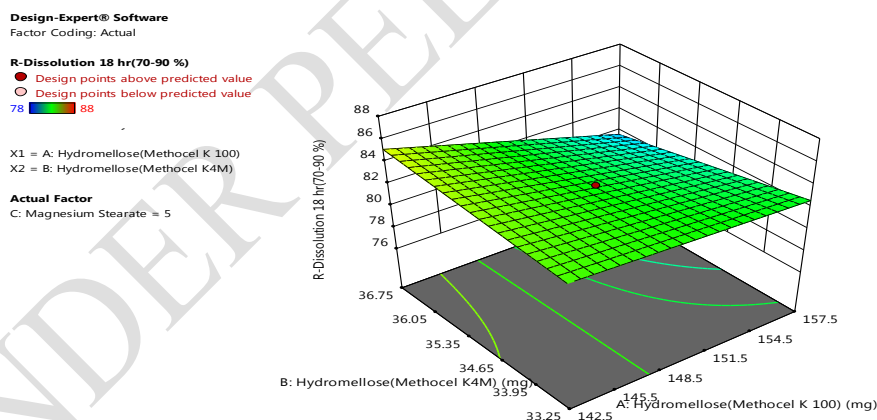
The **Lack of Fit F-value** of 2.76 implies the Lack of Fit is not significant relative to the pure error. There is a 23.85% chance that a Lack of Fit F-value this large could occur due to noise. Non-significant lack of fit is good -- we want the model to fit.

### Fit Statistics

<b>Std. Dev.</b>	1.26	<b>R<sup>2</sup></b>	0.9522
<b>Mean</b>	82.82	<b>Adjusted R<sup>2</sup></b>	0.8407
<b>C.V. %</b>	1.52	<b>Predicted R<sup>2</sup></b>	-5.5523
		<b>Adeq Precision</b>	9.3078

A negative **Predicted R<sup>2</sup>** implies that the overall mean may be a better predictor of your response than the current model. In some cases, a higher order model may also predict better.

**Adeq Precision** measures the signal to noise ratio. A ratio greater than 4 is desirable. Your ratio of 9.308 indicates an adequate signal. This model can be used to navigate the design space.



**Graph 11: Dissolution 18 hrs (factorial design)**

## 5. CONCLUSION

The present work was carried out to formulate and evaluate Darunavir and Ritonavir SR tablets. The drug excipient compatibility studies were carried out. Based on the results, it was confirmed that there is no interaction between drug and excipient at different conditions. Six formulation trials of Darunavir and Ritonavir tablets were conducted using different polymers at different concentrations.

In vitro drug release of all formulations was carried out in dissolution medium 900ml of pH 3.0, 0.05 M Sodium Phosphate Buffer + 2% Tween 20 for 75 RPM USP II apparatus (paddle). The

results shown that the all the formulations matrix tablets shown the good release of the drug from the formulations. In F6, the release of drug shows NLT 80% i.e. near to the desired target release. So, the formulation F6 is suited for the SR table of Darunavir and Ritonavir. Tablets were also evaluated for dissolution studies by QbD method.

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