

# Supplementary Materials: DoE-Based Design of a Simple but Efficient Preparation Method for a Non-Effervescent Gastro-Retentive Floating Tablet Containing Metformin HCl

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**Table S1.** The full factorial design for a screening step and dissolution profiles (mean  $\pm$  SD,  $n = 12$ ), and information on similarity factors, floating lag time and floating retention time of non-EFTs.

Run order	X variable (Factor)		Y variable (Response)			Sample mean ( $f_2$ )	FLT (s)	FRT (h)
	(A)Cetyl alcohol (mg)	(B)HPMC (%)	(R1)Mean diss. at 60min (%)	(R2)Mean diss. at 240min (%)	(R3)Mean diss. at 480min (%)			
F1	50	5	53.0 $\pm$ 3.5	94.4 $\pm$ 2.6	98.9 $\pm$ 1.6	29.35	-	Not
F2	50	5	51.1 $\pm$ 3.2	92.5 $\pm$ 3.0	98.7 $\pm$ 1.0	30.56	-	Not
F3	250	5	44.1 $\pm$ 0.7	78.5 $\pm$ 1.4	91.0 $\pm$ 1.3	41.90	< 3	> 24
F4	250	5	43.4 $\pm$ 0.6	77.7 $\pm$ 1.0	90.4 $\pm$ 1.8	42.92	< 3	> 24
F5	50	25	41.6 $\pm$ 2.9	74.0 $\pm$ 3.2	94.0 $\pm$ 2.0	44.67	< 3	> 24
F6	50	25	41.0 $\pm$ 1.2	73.4 $\pm$ 1.8	93.3 $\pm$ 2.9	45.77	< 3	> 24
F7	250	25	30.9 $\pm$ 0.9	58.1 $\pm$ 1.4	75.3 $\pm$ 1.7	69.38	< 3	> 24
F8	250	25	31.4 $\pm$ 0.6	58.6 $\pm$ 1.0	76.0 $\pm$ 1.5	71.46	< 3	> 24
F9	150	15	38.6 $\pm$ 1.0	74.1 $\pm$ 1.2	92.1 $\pm$ 1.0	47.26	< 3	> 24
F10	150	15	37.4 $\pm$ 0.4	73.4 $\pm$ 1.4	92.0 $\pm$ 1.6	48.50	< 3	> 24

**Table S2.** The response surface methodology for optimization step and dissolution profiles (mean  $\pm$ SD,  $n=12$ ), and information on similarity factors, floating lag time and floating retention time of non-EFTs.

Run order	X variable (Factor)		Y variable (Response)			Sample mean ( $f_2$ )	FLT (s)	FRT (h)
	(A) Cetyl alcohol (mg)	(B) HPMC (%)	(R1) Mean diss. at 60min (%)	(R2) Mean diss. at 240min (%)	(R3) Mean diss. at 480min (%)			
M1	150	15	39.6 $\pm$ 0.4	73.5 $\pm$ 0.5	91.8 $\pm$ 0.5	47.36	< 3	> 24
M2	250	15	37.1 $\pm$ 1.5	67.9 $\pm$ 2.2	86.4 $\pm$ 2.2	58.15	< 3	> 24
M3	150	25	34.1 $\pm$ 0.6	64.8 $\pm$ 1.5	84.3 $\pm$ 2.1	69.08	< 3	> 24
M4	250	25	30.2 $\pm$ 1.1	58.2 $\pm$ 2.3	77.1 $\pm$ 3.3	75.47	< 3	> 24
M5	150	20	37.0 $\pm$ 0.6	69.0 $\pm$ 1.5	88.5 $\pm$ 2.0	55.41	< 3	> 24
M6	250	20	33.1 $\pm$ 0.8	62.9 $\pm$ 1.2	83.2 $\pm$ 1.9	76.76	< 3	> 24
M7	200	15	38.1 $\pm$ 1.1	68.9 $\pm$ 1.0	86.0 $\pm$ 0.8	56.15	< 3	> 24
M8	200	25	32.8 $\pm$ 0.7	62.3 $\pm$ 1.1	79.4 $\pm$ 1.7	77.88	< 3	> 24
M9	200	20	34.5 $\pm$ 0.7	64.8 $\pm$ 1.0	80.1 $\pm$ 1.6	69.32	< 3	> 24
M10	200	20	34.1 $\pm$ 0.6	63.8 $\pm$ 1.1	79.6 $\pm$ 1.2	71.74	< 3	> 24

**Table S3.** Comparison of dissolution profiles (mean  $\pm$  SD,  $n = 12$ ) for the external validation set, model prediction accuracy and bootstrap analysis.

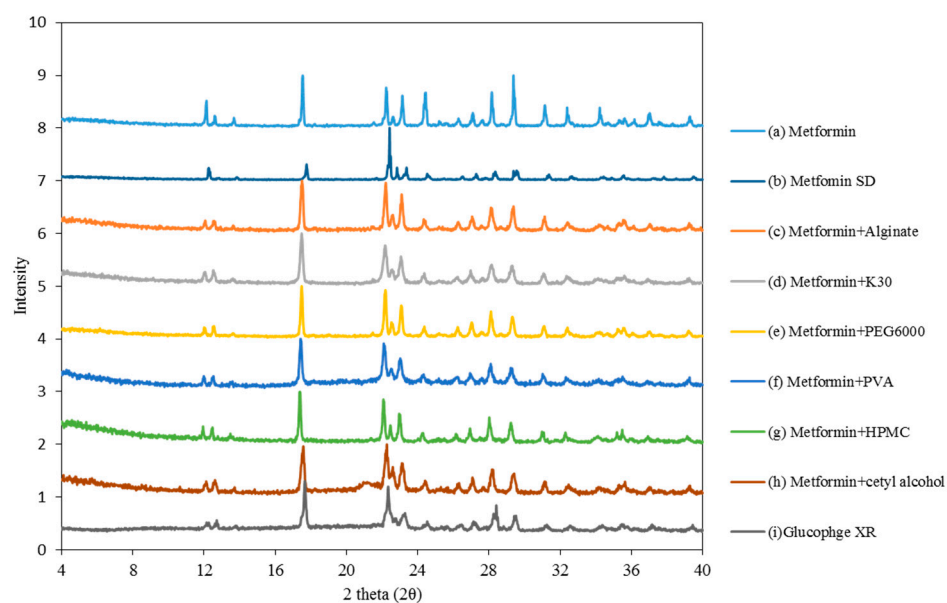
Code	Cetyl alcohol (mg)	HPMC (%)	Cont.	60 min (%)	240 min (%)	480 min (%)	Sample mean ( $f_2$ )	Bootstrap analysis (500)			
								$f_2$	E( $f_2$ )	PI	Bca
			Glucophage XR	29.4 $\pm$ 0.9	60.4 $\pm$ 1.2	81.6 $\pm$ 1.3	-	-	-	-	-
E1	150	5	True mean	43.9 $\pm$ 0.7	82.3 $\pm$ 2.6	97.1 $\pm$ 1.4	37.68	37.69	37.67	(36.88, 38.49)	(36.84, 38.44)
			Linear Pred. mean	45.6	81.2	98.3	37.18				
			Quad. Pred. mean	45.6	81.2	99.9	36.58				
E2	150	10	True mean	40.9 $\pm$ 0.6	77.1 $\pm$ 0.5	94.2 $\pm$ 0.3	42.98	42.98	42.95	(42.29, 43.62)	(42.35, 43.67)
			Linear Pred. mean	42.7	77.0	94.4	42.13				
			Quad. Pred. mean	42.7	77.0	96.0	41.39				
E3	200	5	True mean	43.3 $\pm$ 0.4	79.4 $\pm$ 0.8	95.4 $\pm$ 0.5	40.09	40.06	40.04	(39.47, 40.69)	(39.50, 40.74)
			Linear Pred. mean	43.9	78.1	95.3	40.58				
			Quad. Pred. mean	43.9	78.1	93.0	41.50				
E4	200	10	True mean	40.2 $\pm$ 0.2	73.4 $\pm$ 0.9	90.7 $\pm$ 1.3	47.68	47.73	47.64	(46.79, 48.53)	(46.91, 48.75)
			Linear Pred. mean	41.0	73.9	91.4	46.45				
			Quad. Pred. mean	41.0	73.9	89.1	47.55				
E5	250	5	True mean	44.1 $\pm$ 0.6	77.1 $\pm$ 1.1	91.9 $\pm$ 0.8	42.40	42.45	42.43	(41.69, 43.18)	(41.70, 43.19)

E6	250	10	Linear Pred. mean	42.2	75.1	92.4	44.46				
			Quad. Pred. mean	42.2	75.1	93.9	43.74				
			True mean	39.6 ± 0.9	72.0 ± 1.1	87.9 ± 2.0	50.70	50.81	50.64	(49.41, 51.73)	(49.43, 51.79)
			Linear Pred. mean	39.3	70.9	88.5	51.60				
			Quad. Pred. mean	39.3	70.9	90.0	50.68				
			Linear model	RMSEP	1.34	1.18	0.66				
	Quadratic model	RMSEP	1.34	1.18	2.17						

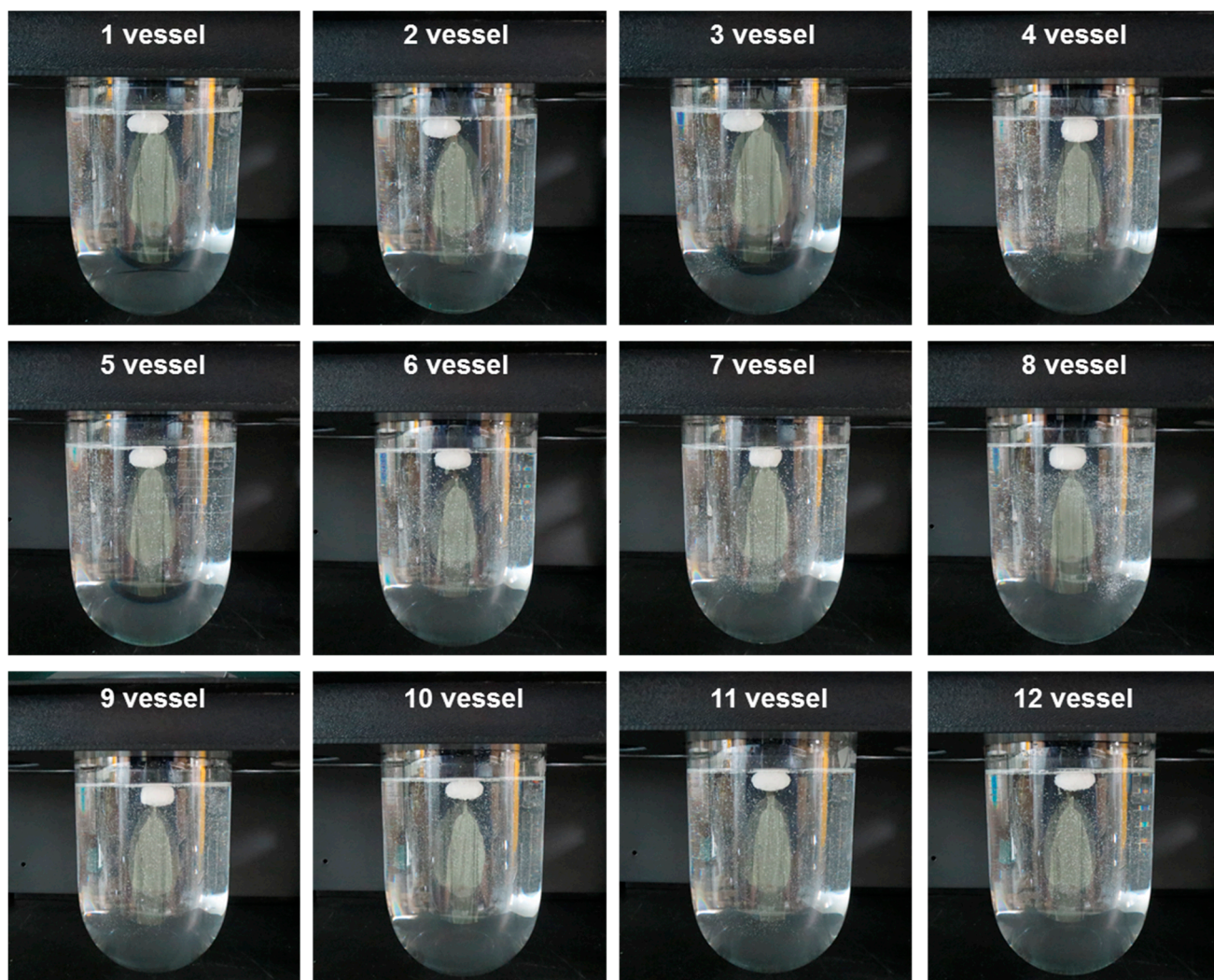
PI: The percentile confidence interval; Bca: The bias corrected and accelerated confidence interval, RMSEP: The root mean squared error of prediction.

**Table S4.** Comparison of the dissolution profiles for the optimized formulations predicted using the RSM with the experimentally obtained dissolution profiles.

Code	Cetyl alcohol (mg)	HPMC (%)	Mean dissolution profile	60min (%)	240min (%)	480min (%)	Sample mean (f <sub>2</sub> )
P1	150	16	Predicted	39.14	71.99	89.74	50.04
			Experimentally obtained	39.1 ± 0.7	72.2 ± 0.9	89.7 ± 1.5	49.85
P2		17	Predicted	38.55	71.16	88.96	51.68
			Experimentally obtained	38.1 ± 0.3	70.7 ± 0.3	89.0 ± 0.9	52.41
P3		18	Predicted	37.96	70.33	88.18	53.44
			Experimentally obtained	37.2 ± 0.4	70.3 ± 0.4	88.5 ± 0.4	53.80
P4		19	Predicted	37.37	69.49	87.40	55.36
			Experimentally obtained	37.3±0.5	69.6±0.9	88.4±1.1	54.63

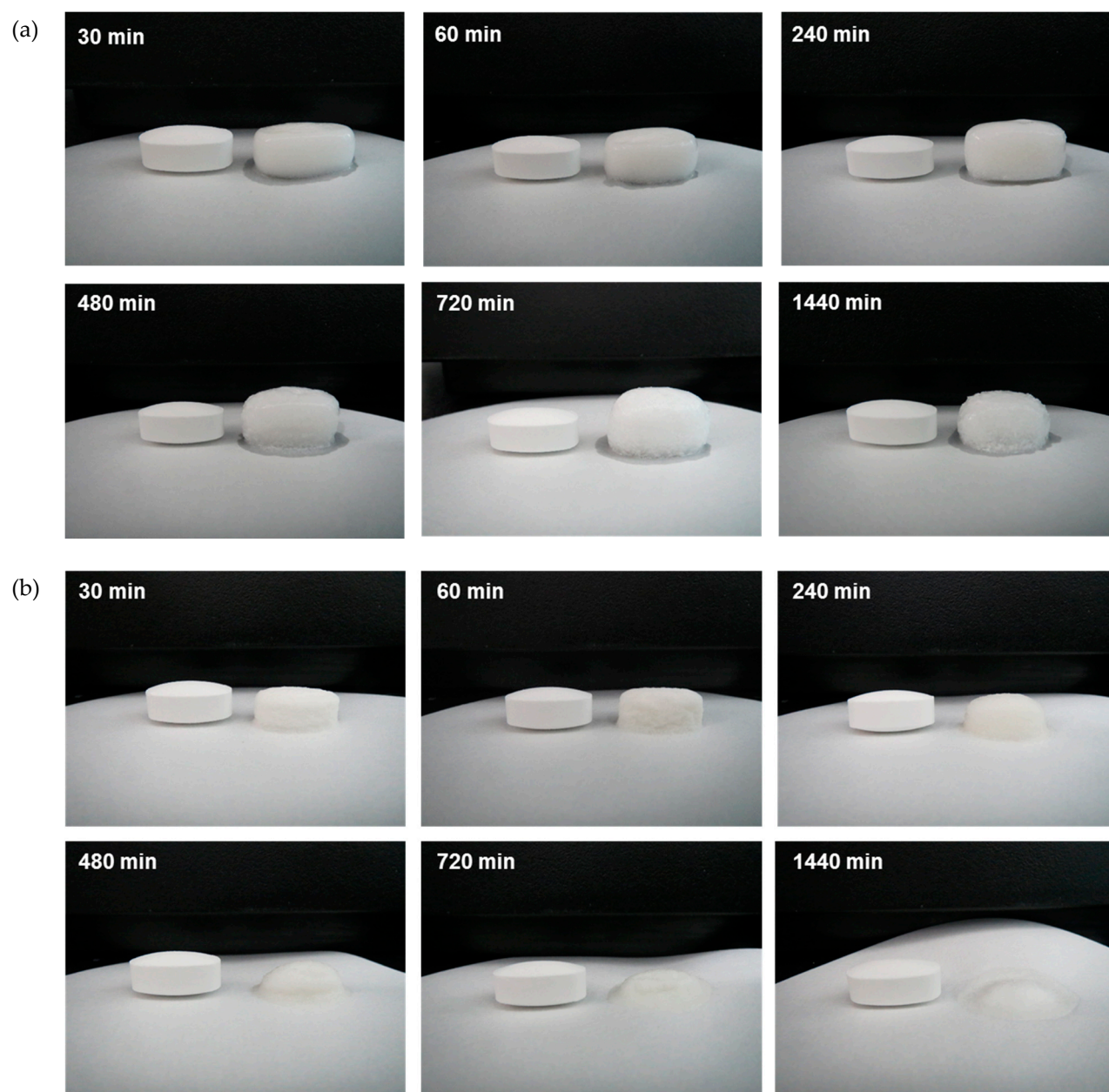


**Figure S1.** PXRD patterns for crystalline metformin HCl, co-spray dried solid dispersions, and Glucophage XR.



**Figure S2.** Images of non-EFTs in the vessels after the 24-h dissolution test ( $n = 12$ ).





**Figure S3.** Images of non-EFTs during (a) swelling and (b) erosion tests over time. Non-EFTs were taken out from the vessels at predetermined times during the dissolution test.