

**Table S1.** Release curves comparison using Fit factors f1 and f2.

<b>Incubation of the matrices with PBS without any medium replacement</b>			
<b>Matrix type</b>	<b>5% PCL/SRL/10% HSA</b>	<b>5% PCL/SRL/10% HSA/3%DMSO</b>	<b>5% PCL/SRL</b>
5% PCL/SRL/10% HSA	-	f1 = 7.3 f2 = 70.2	f1 = 4.5 f2 = 76.8
5% PCL/SRL/10% HSA/3%DMSO	f1 = 7.3 f2 = 70.2	-	f1 = 8.0 f2 = 67.7
5% PCL/SRL	f1 = 4.5 f2 = 76.8	f1 = 8.0 f2 = 67.7	-
<b>Incubation of matrices with PBS with medium replacement</b>			
<b>Matrix type</b>	<b>5% PCL/SRL/10% HSA</b>	<b>5% PCL/SRL/10% HSA/3%DMSO</b>	<b>5% PCL/SRL</b>
5% PCL/SRL/10% HSA	-	f1 = 26.6 f2 = 35.6	f1 = 17.0 f2 = 42.4
5% PCL/SRL/10% HSA/3%DMSO	f1 = 26.6 f2 = 35.6	-	f1 = 16.4 f2 = 49.4
5% PCL/SRL	f1 = 17.0 f2 = 42.4	f1 = 16.4 f2 = 49.4	-
<b>Incubation of the matrices with BP without any medium replacement</b>			
<b>Matrix type</b>	<b>5% PCL/SRL/10% HSA</b>	<b>5% PCL/SRL/10% HSA/3%DMSO</b>	<b>5% PCL/SRL</b>
5% PCL/SRL/10% HSA	-	f1 = 27.3 f2 = 29.7	f1 = 18.8 f2 = 36.7
5% PCL/SRL/10% HSA/3%DMSO	f1 = 27.3 f2 = 29.7	-	f1 = 12.6 f2 = 50.9
5% PCL/SRL	f1 = 18.8 f2 = 36.7	f1 = 12.6 f2 = 50.9	-
<b>Incubation of the matrices with BP with medium replacement</b>			
<b>Matrix type</b>	<b>5% PCL/SRL/10% HSA</b>	<b>5% PCL/SRL/10% HSA/3%DMSO</b>	<b>5% PCL/SRL</b>
5% PCL/SRL/10% HSA	-	f1 = 24.9 f2 = 32.0	f1 = 1.8 f2 = 74.3

5% PCL/SRL/10%	f1 = 24.9	f1 = 31.0
HSA/3%DMSO	f2 = 32.0	f2 = 32.0
5% PCL/SRL	f1 = 1.8	f1 = 31.0
	f2 = 74.3	f2 = 32.0

**Table S2.** Release curves comparison using Fit factors f1 and f2. Comparison of kinetic curves of SRL release from the parent and the two-fold expanded matrices. Incubation of the matrices in BP with medium replacement.

Matrix type	5% PCL/SRL/10% HSA expanded	5% PCL/SRL/10% HSA/3%DMSO expanded	5% PCL/SRL expanded
5% PCL/SRL/10%	f1 = 2.4	-	-
HSA	f2 = 77.6	-	-
5% PCL/SRL/10% HSA/3%DMSO	-	f1 = 2.6 f2 = 75.0	-
5% PCL/SRL	-	-	f1 = 2.6 f2 = 76.9

Fit factors were calculated as:

$$f_1 = \{[\sum_{t=1}^n |R_t - T_t|] / [\sum_{t=1}^n R_t]\} \cdot 100$$

$$f_2 = 50 \cdot \log \{[1 + (1/n) \sum_{t=1}^n (R_t - T_t)^2]^{0.5} \cdot 100\}$$

R<sub>t</sub> and T<sub>t</sub> are the cumulative percentage released at each of the selected n time points of the reference and test product respectively. The difference factor f<sub>1</sub> is proportional to the average difference between the two profiles, where as similarity factor f<sub>2</sub> is inversely proportional to the average squared difference between the two profiles, with emphasis on the larger difference among all the time-points. The f<sub>1</sub> value is 0 when the test and the reference profiles are identical and increases proportionally with the dissimilarity between the two profiles. The f<sub>2</sub> value is between 0 and 100. f<sub>2</sub> value between 50-100 indicates similarity between two dissolution profiles.

## References

Shah, V.P.; Tsong, Y.; Sathe, P.; Williams, R.L. Dissolution Profile Comparison Using Similarity Factor, f<sub>2</sub>. *Dissolution Technologies*, 1999, 6, 15.