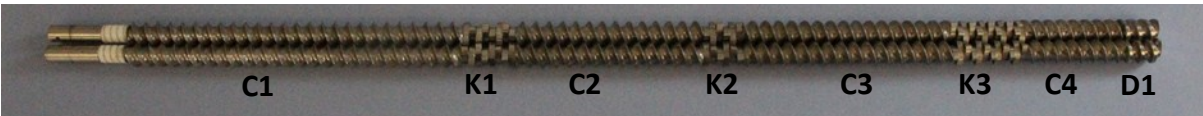


Electronic Supplementary information

Solid State Thermomechanical Engineering of High-Quality Pharmaceutical Salts via Solvent Free Continuous Processing

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1. Screw Configuration



	C1	K1	C2	K2	C3	K3	C4	D1
CFX-MA F1	100	160	160	160	200	200	200	200
CFX-MA F2	100	160	160	160	160	160	160	160
CFX-MA F3	80	120	120	120	140	140	140	140
CFX-MA F4	80	100	100	130	130	100	100	100
KTZ- OA	50	75	110	175	185	185	185	185

Fig. S1.1†: Complete screw profile showing the order of the Conveying (C), Kneading (K) and Discharge (D) zones



Fig. S1.2†: Screw configuration for Kneading zone 1



Fig. S1.3†: Screw configuration for Kneading zone 2



Fig. S1.4†: Screw configuration for Kneading zone 3



Fig. S1.5†: Screw configuration for the Conveying zones



Fig. S1.6†: Screw configuration for the Discharge zone

2. Particle Size and Morphology

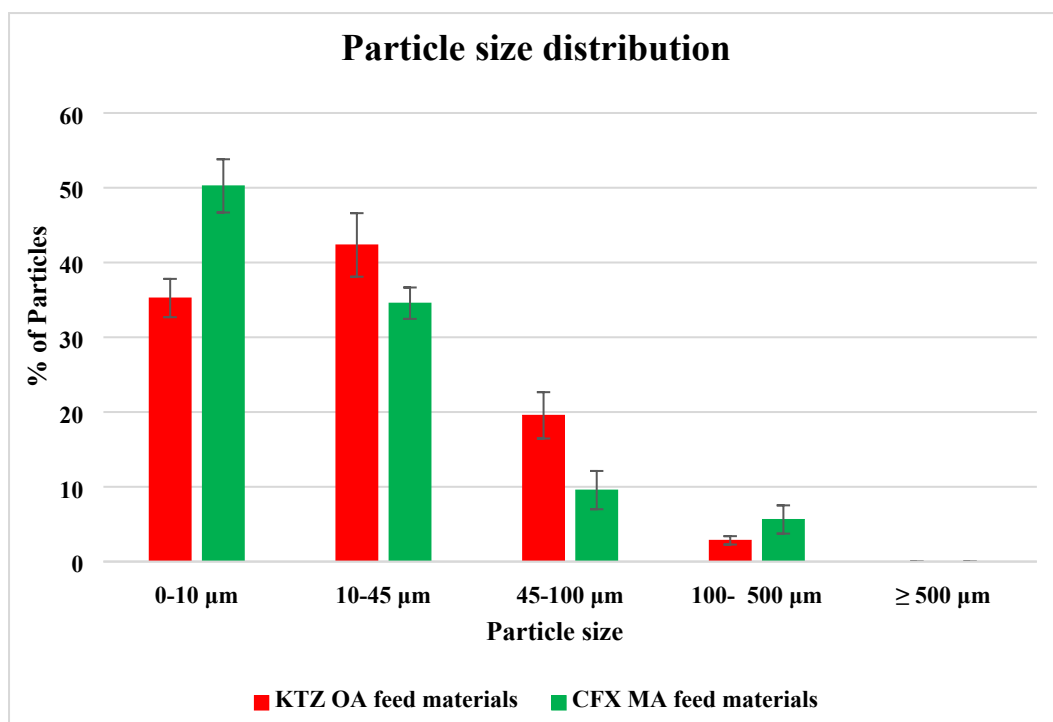


Fig. S2.1†. Particle size distribution of the feed materials for both mixtures used in ssTME and LAG processing.

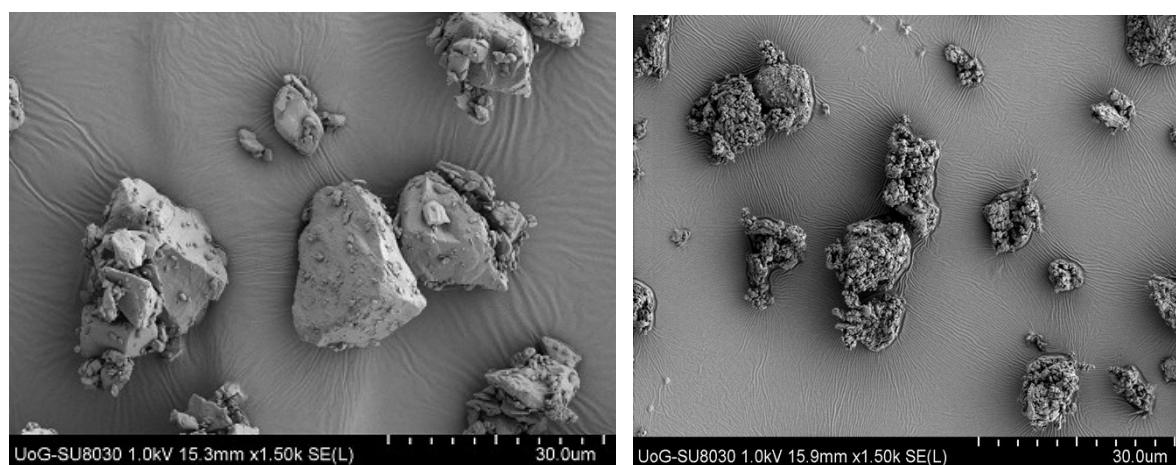


Fig. S2.2 †: SEM images of bulk KTZ (Left) and CFX (Right)

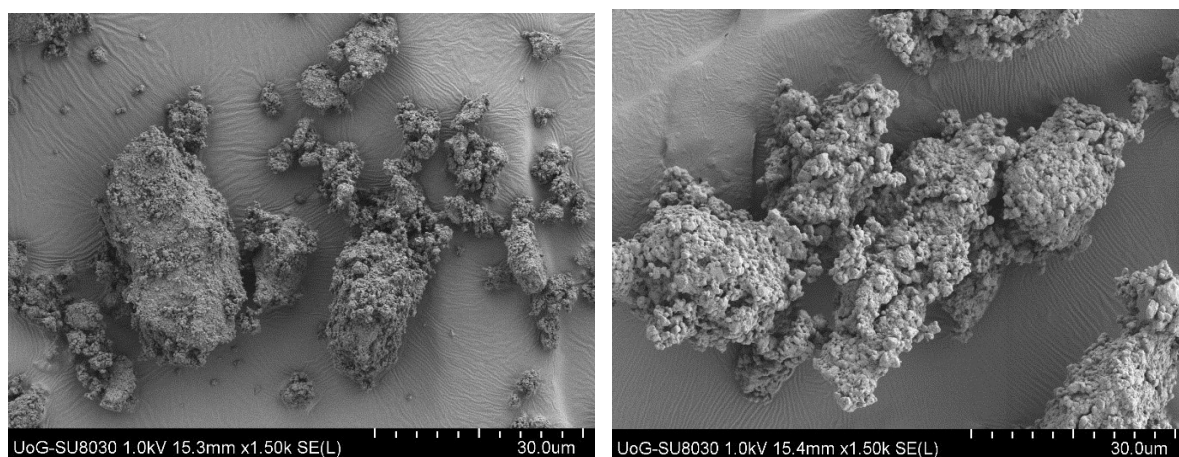


Fig. S2.3 †: SEM images of KTZ-OA salts and CFX-MA salts produced via LAG

3. Thermal Analysis

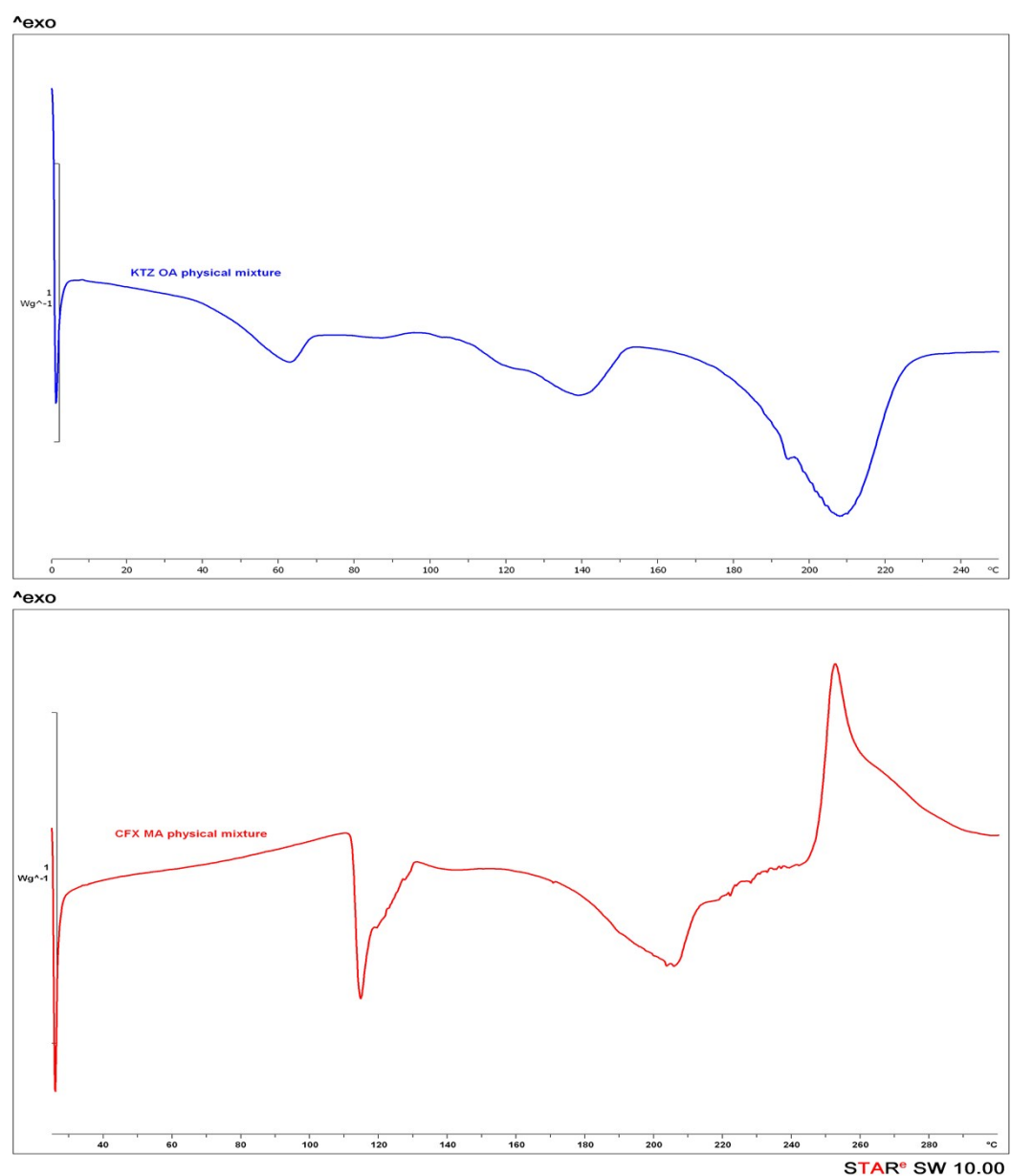


Fig. S3.1†. DSC thermogram of KTZ – OA and CFX – MA physical mixture.

4. PXRD analysis

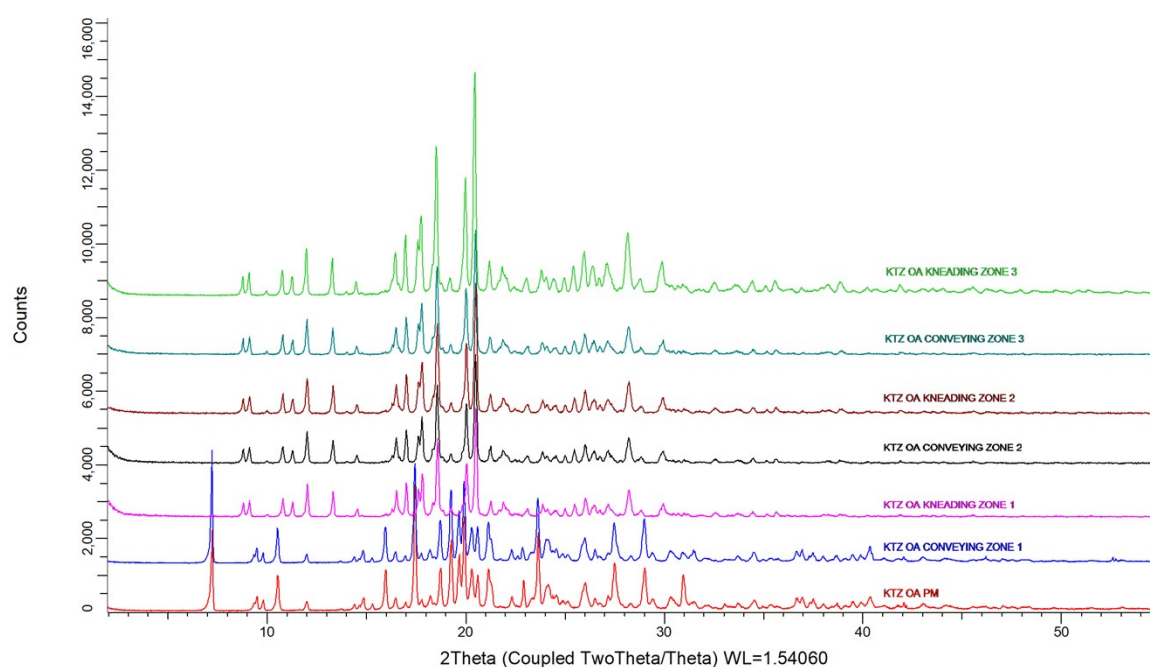


Fig. S4.1†. PXRD analysis of KTZ-OA salts taken from each individual conveying and kneading zone within the twin-screw extruder, showing the cocrystals transformation

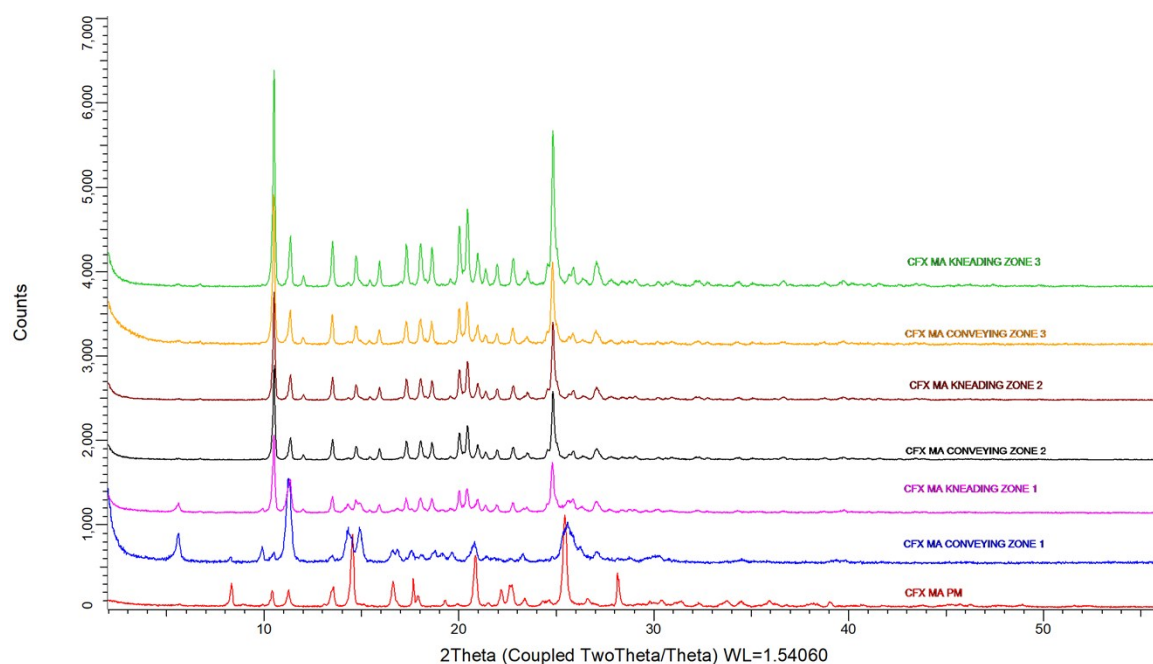
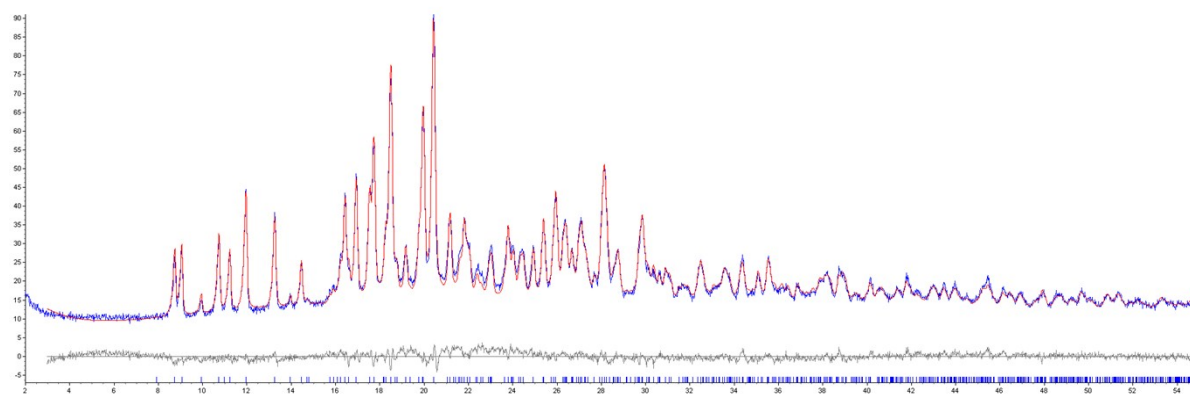


Fig. S4.2†. PXRD analysis of CFX-MA salts taken from each individual conveying and kneading zone within the twin-screw extruder, showing the cocrystals transformation pathway.

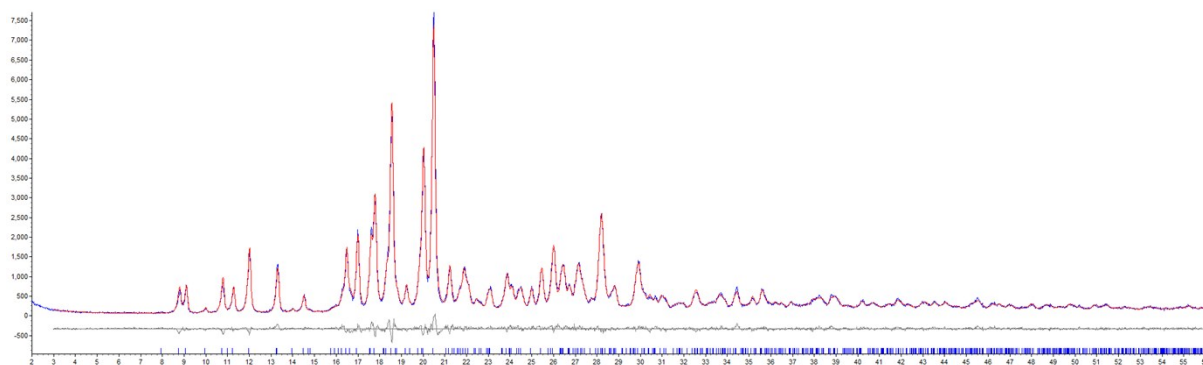


R-Values:

R_{exp} : 4.73 R_{wp} : 8.74 R_p : 6.69 GOF: 1.85

R_{exp} : 6.05 R_{wp} : 11.18 R_p : 9.14 DW: 0.64

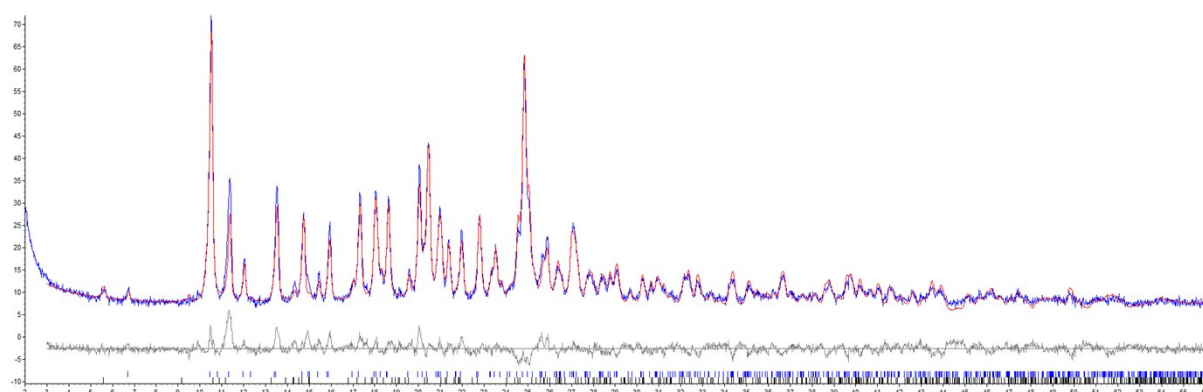
Fig. S4.3†: Rietveld refinement of the PXRD data for the KTZ OA ssTME, where the measured pattern is represented with blue line, the simulated pattern with the red line and the difference pattern in grey. Refinement values displayed.



R-Values:

Rexp : 4.79 Rwp : 6.78 Rp : 5.23 GOF : 1.42
 Rexp` : 5.74   Rwp` : 8.13 Rp` : 6.50 DW : 1.03

Fig. S4.4†: Rietveld refinement of the PXRD data for the KTZ OA LAG, where the measured pattern is represented with blue line, the simulated pattern with the red line and the difference pattern in grey. Refinement values displayed.



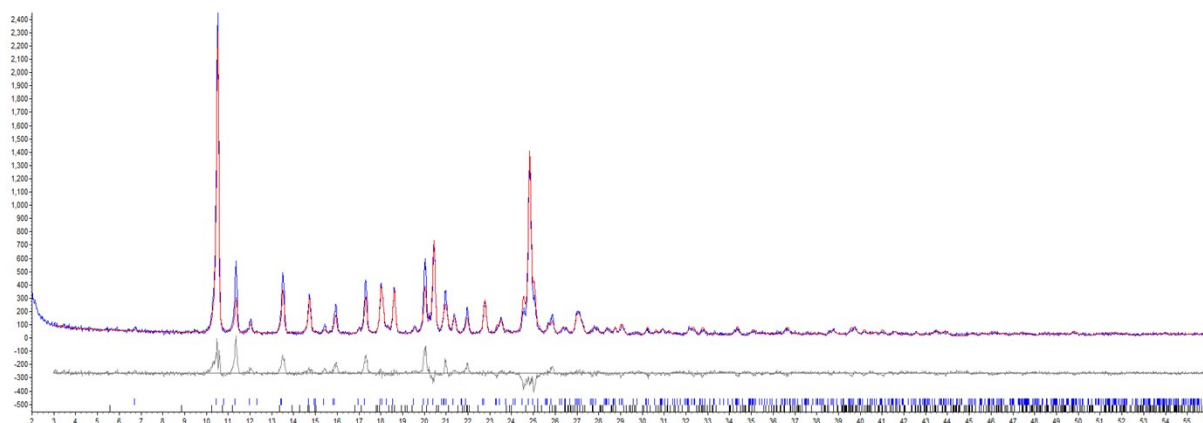
R-Values

Rexp : 7.70 Rwp : 16.17 Rp : 12.66 GOF : 2.10
 Rexp` : 8.31   Rwp` : 17.45 Rp` : 13.87 DW : 0.50

Quantitative Analysis - Rietveld

Phase 1 : "CFX MA Polymorph 2" 99.048 %
 Phase 2 : "CFX MA Polymorph 1" 0.952 %

Fig. S4.5†: Rietveld refinement of the PXRD data for the CFX MA ssTME, where the measured pattern is represented with blue line, the simulated pattern with the red line and the difference pattern in grey. Refinement values displayed.



R-Values

Rexp : 11.73 Rwp : 20.34 Rp : 15.96 GOF : 1.73
 Rexp` : 14.17 Rwp` : 24.57 Rp` : 19.65 DW : 0.70

Quantitative Analysis - Rietveld

Phase 1 : "CFX MA Polymorph 2"	99.001 %
Phase 2 : "CFX MA polymorph 1"	0.999 %

Fig. S4.6†: Rietveld refinement of the PXRD data for the CFX MA LAG, where the measured pattern is represented with blue line, the simulated pattern with the red line and the difference pattern in grey. Refinement values displayed.

5. Release Studies

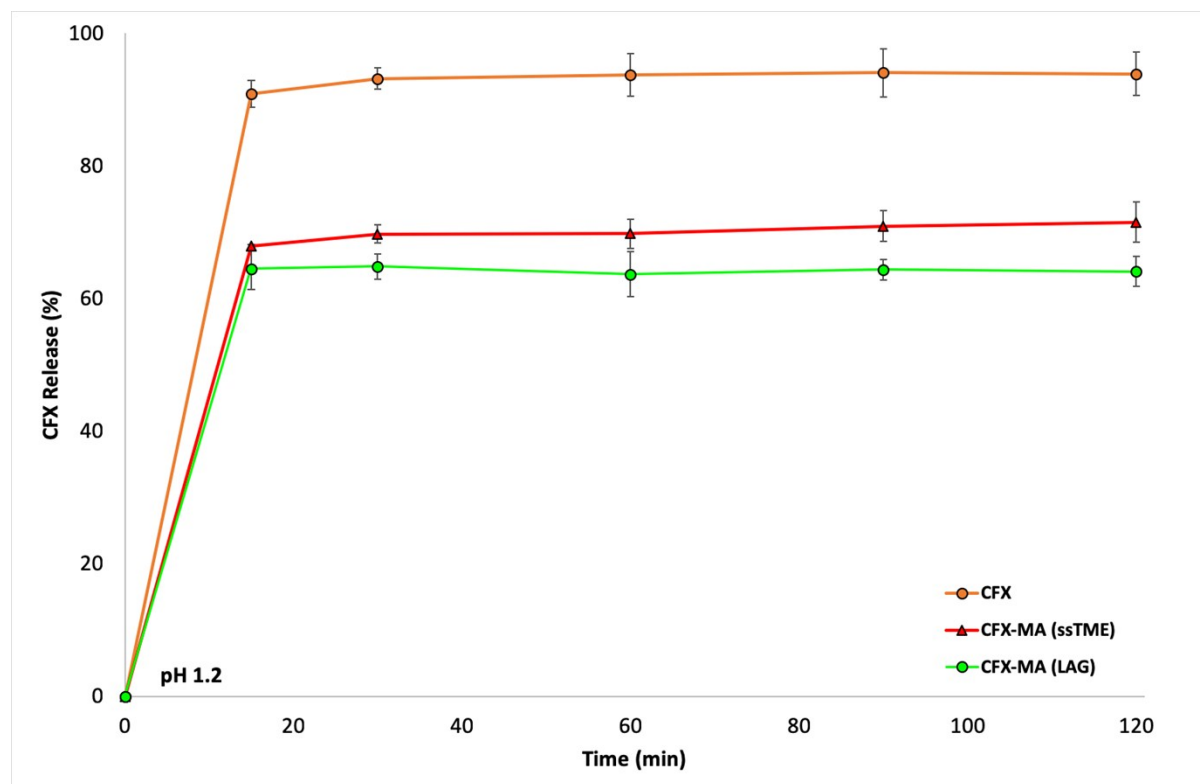


Fig. S5.1†: Dissolution profiles of bulk CFX and CFX-MA salt synthesised by ssTME and LAG.

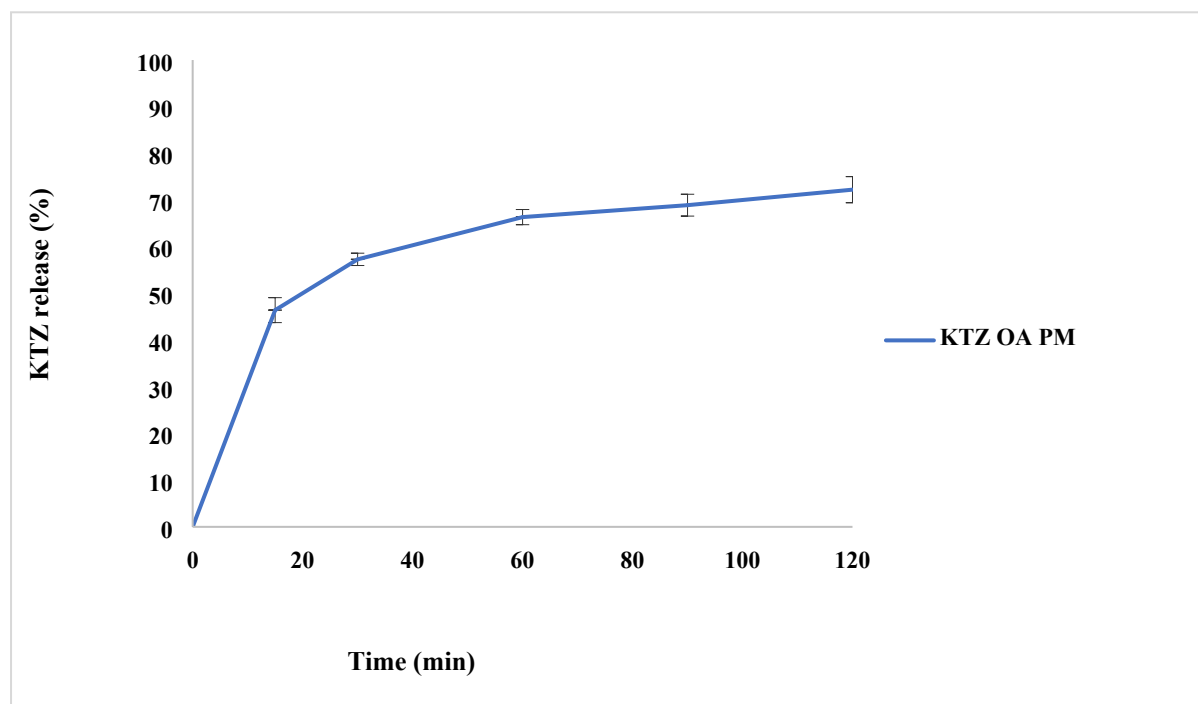


Fig. S5.2†: Dissolution profile for the KTZ-OA physical blend at pH 4.4.

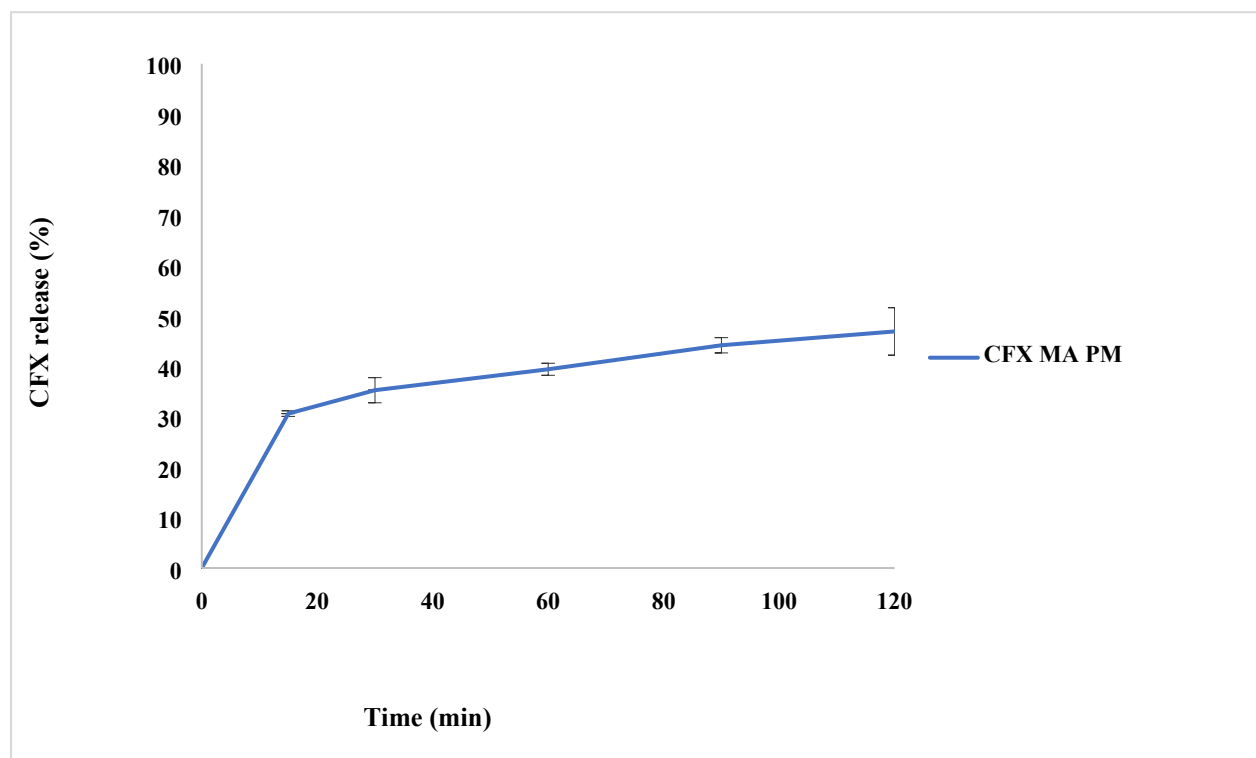


Fig. S5.3†: Dissolution profile for the CFX-MA physical blend at pH 6.8

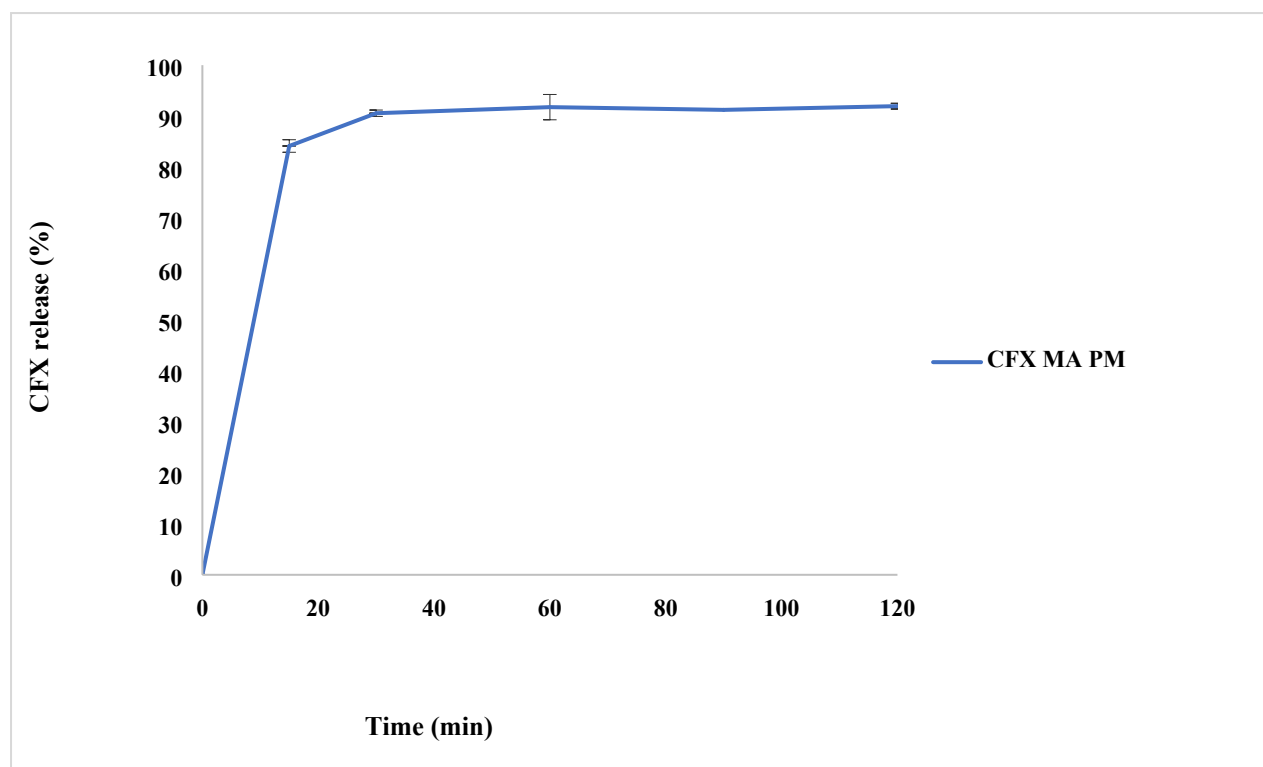


Fig. S5.4†: Dissolution profile for the CFX-MA physical blend at pH 1.2