

## ***In situ* non-invasive Raman spectroscopic characterisation of succinic acid polymorphism during segmented flow crystallisation**

Anuradha R. Pallipurath,<sup>1†</sup> Pierre-Baptiste Flandrin,<sup>1</sup> Lois E. Wayment,<sup>1, 2, 3</sup> Chick C. Wilson<sup>1,2</sup> and Karen Robertson<sup>1‡\*</sup>

1. Department of Chemistry, University of Bath, Claverton Down, BA2 7AY, UK
2. CMAC Future Manufacturing Hub, University of Bath, Claverton Down, Bath BA2 7AY, UK
3. Diamond Light Source, Harwell Campus, Didcot, Oxfordshire OX11 0DE, UK

\* Email: Karen.Robertson@nottingham.ac.uk

† Present Address: Department Chemistry, CMAC Future Manufacturing Hub, University of Leeds, Woodhouse Lane, Leeds, LS2 9JT

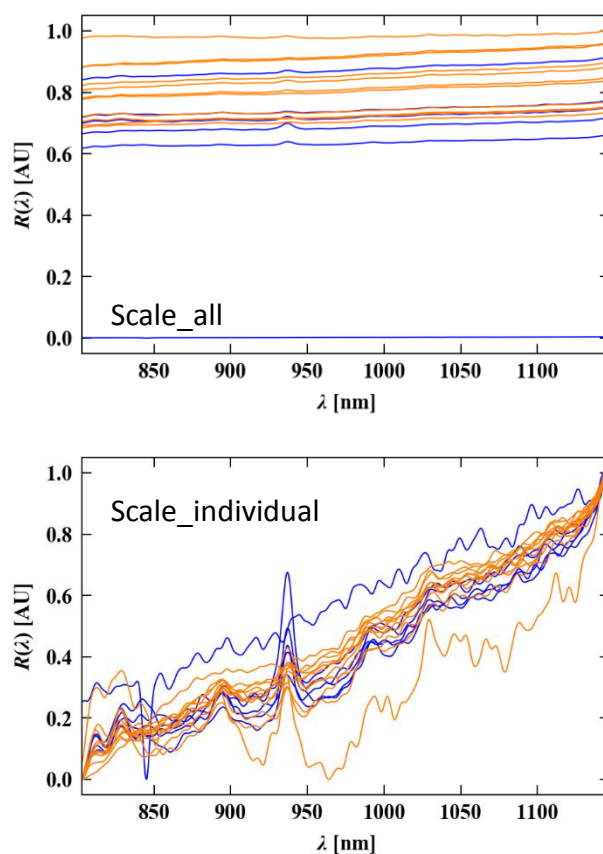
‡ Present Address: Department of Chemical and Environmental Engineering, University of Nottingham, University Park, Nottingham NG7 2RD

## **Supporting information**

## Contents

1. Scale_all algorithm.....	3
2. 2D plots of Raman spectra for slurry of $\beta$ -SA.....	4
3. Offline analysis of SA produced in the KRAIC .....	5
4. 1D plots of Raman data.....	5
5. Tentative assignments of Raman peaks.....	6

## 1. Scale\_all algorithm

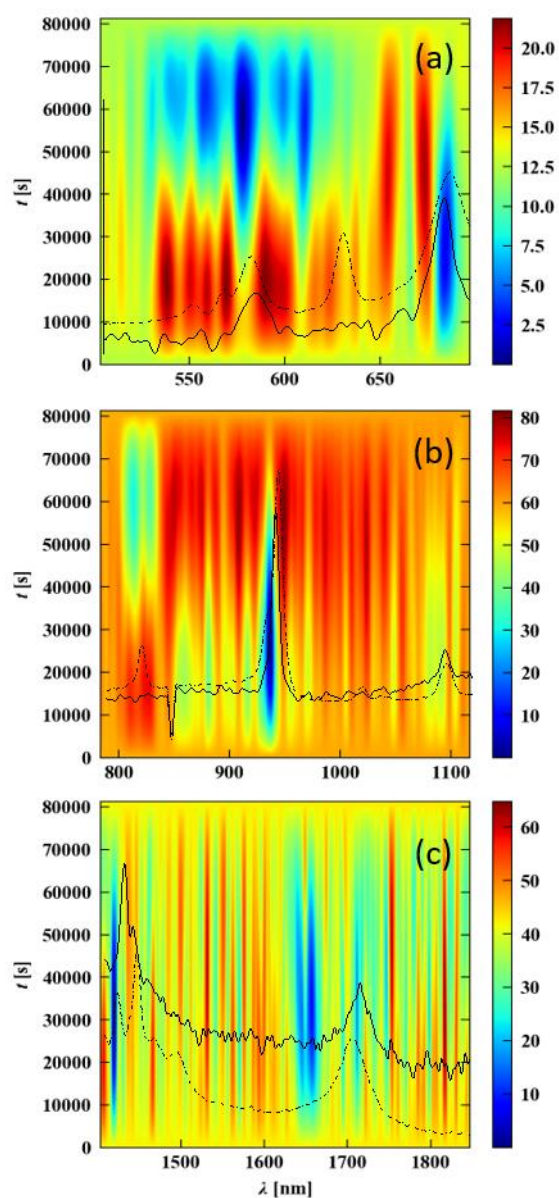


**Figure S1:** Time-resolved Raman spectra of succinic acid form  $\beta$  high density slurry collected after segmentation in the KRAIC set-up. The ‘scale\_all’ algorithm exaggerates the effect of the background thereby enabling distinction between regions of high and low Raman density from solids. The ‘scale\_individual’ algorithm improves the S/N ratio of the features from the solids against the background.

**‘Scale\_all’:** Each spectra in the time-dependent data collection is treated simultaneously. Every data point (intensity) in a spectrum is subtracted by the lowest intensity across all of the spectra and divide by the difference in the maximum and minimum intensity across all of the spectra.

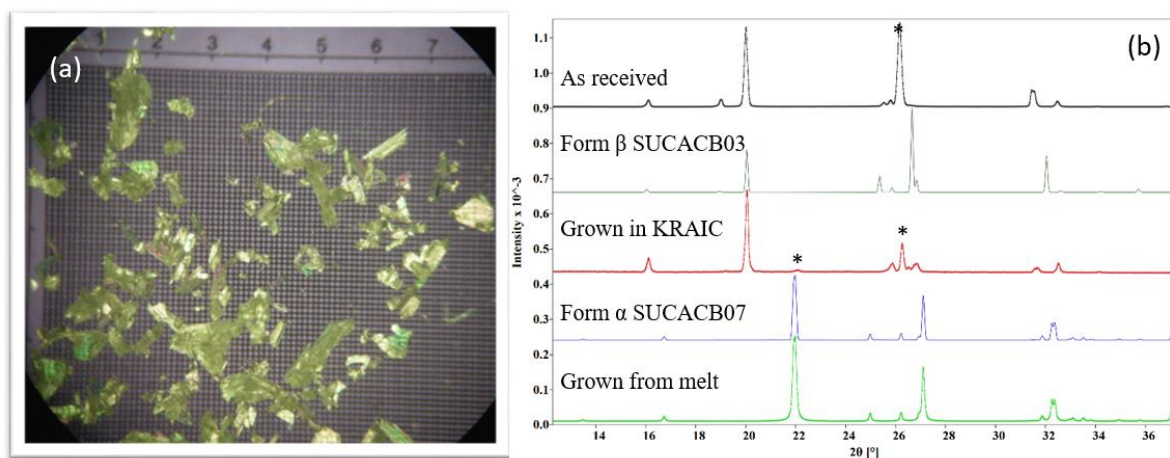
$$I_{Sa} = \frac{I - I_{\min\_all}}{I_{\max\_all} - I_{\min\_all}} \quad \text{eq (1)}$$

## 2. 2D plots of Raman spectra for slurry of $\beta$ -SA



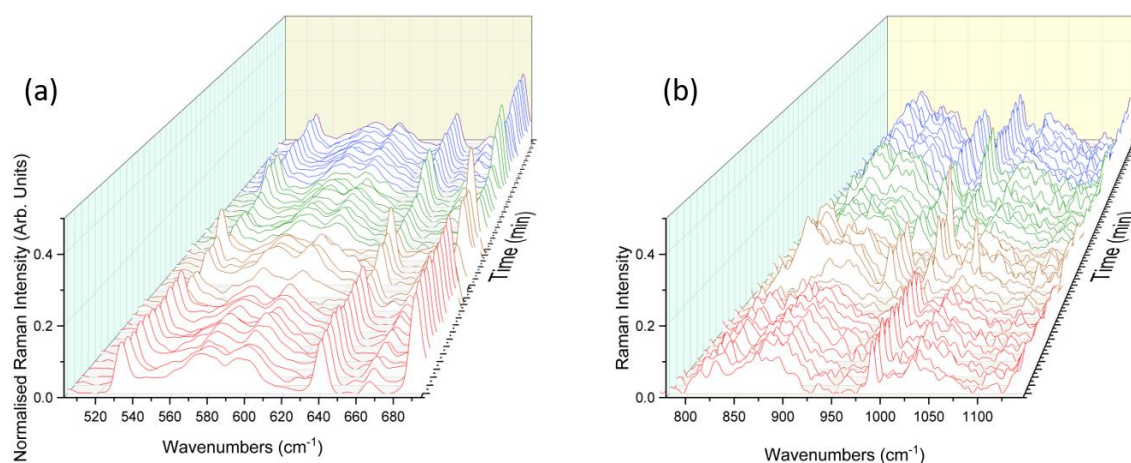
**Figure S2:** Time-dependent Raman spectra collected from pre-prepared high density slurry of succinic acid form  $\beta$ . In the region of (a) 500 – 700  $\text{cm}^{-1}$  (b) 780 – 1120  $\text{cm}^{-1}$  (c) 1400 – 1850  $\text{cm}^{-1}$ . The regions of interest were scaled individually, to enhance the Raman peaks from the solids flowing through. Overlaid spectra are  $\beta$ -SA (red) and  $\alpha$ -SA (grey).

### 3. Offline analysis of SA produced in the KRAIC



**Figure S3:** (a) Microscope image of succinic acid (SA) crystallised from KRAIC, (b) Powder X-ray diffraction of the polymorphic forms of SA as received (form  $\beta$ ), grown from KRAIC (form  $\beta$ , with small  $\alpha$  impurities), form  $\alpha$  grown from melt and simulated patterns of both forms from crystal structures deposited in the Cambridge Structural Database (CSD) – REF code: SUCACB03 and SUCACB07.

### 4. 1D plots of Raman data



**Figure S4:** 1d plots of PCM FI and FII slurry before background subtraction. The major intensities are from the FEP tubing.

## 5. Tentative assignments of Raman peaks

**Table S1:** Tentative assignments for fluoro ethylene propylene reactor and reactor with cyclohexane.

FEP	FEP+solvent	Tentative analysis
208.4961 239.2048 295.2031	293.3967156	C-C stretches
387.3293 19.8444 580.6137 602.2904 643.8375	387.3292999 418.0380294	C-C deformations
734.1573 752.2213	692.6101988 734.1573034 750.414866	C-F deformation
940.0864 1109.888		C-C-C-C Symmetric vibrations
1164.08	1151.434745 1191.175454	C-C-C stretches
1218.271 1247.174		C-F stretching
1299.559	1299.559205	CH <sub>2</sub> -CH <sub>2</sub> twisting
1348.332 1382.653	1326.655143 1351.944685 1382.653414 1435.038894	
1545.229	1543.422645	CH <sub>2</sub> F deformation
	1604.840104 1850.509939 2052.826275 2060.051858 2164.822818 2240.691443 2331.011236 2370.751945	
2601.971 2876.543	2880.155575 2986.73293	C-H stretching

**Table S2:** Tentative assignments for succinic acid polymorphs

$\alpha$ -SA	$\beta$ -SA	Tentative analysis
94.69317	83.8548 94.69317 125.4019	Phonon modes
130.8211	136.2403	
148.885		
175.981	161.5298	
212.1089	266.3008 304.2351	C-C stretches
320.4927	320.4927	
392.7485	387.3293 457.7787	C-C deformation in aliphatic carbons
549.9049		
569.7753		C-O out-of-plane deformation
582.4201	582.4201	
631.1927	663.7079	out-of-plane bending of H-bonded OH
687.191	685.3846	
	703.4486	
	714.2869	
817.2515		
900.3457	887.701	Out-of-plane C-O deformations
940.0864	940.0864	
	963.5696	
	983.4399	
1012.342		monomeric C-O stretching
1032.213	1035.825	
1088.211	1086.404	
1165.886	1160.467 1198.401	C-H deformations
1216.465		
1238.142	1232.723	
	1247.174	
	1265.238	
	1277.882	
1297.753	1295.946	

	1323.042	
1382.653	1371.815	
1400.717	1400.717	C-O ...OH combination band
1413.362	1420.588	
1435.039		
1449.49		CH <sub>2</sub> deformation
1472.973		
1548.842		
1650	1655.419	C=O stretching
	1687.934	
	1725.869	
1816.188	1763.803	
1848.704	1799.931	
1875.799		
1908.315		
1931.798		
2329.205		H-bonded COOH stretches
2553.198		
2591.132	2576.681	
	2645.324	
	2782.61	
2851.253	2865.704	Aliphatic C-H stretching
2865.704	2885.575	
	2910.864	
2934.347	2928.928	Aromatic C-H stretching
2946.992	2948.799	
2966.863	2968.669	
3008.41		
3089.697		O-H vibrations



**Table S3:** Tentative assignments for paracetamol polymorphs

PCM FI	PCM FII	Tentative analysis
89.7		Phonon modes
128.0355	118.774	
	137.2969	
158.9069		
217.5628	205.2142	
328.7001	331.7873	C-C stretching
	365.7459	
393.5303	396.6174	
415.1403	421.3146	
467.6219	452.1861	
501.5805	510.8419	
603.4564	606.5436	out-of-plane bending H-bonded OH
631.2408	628.1536	
652.8508	649.7637	
686.8095		
711.5066	711.5066	out-of-plane N-H deformation in H-bonded secondary amides
739.291		
797.9468	801.034	N-H wagging in secondary amines and C-H out of plane deformation in ortho substituted aromatics
834.9926	838.0798	
858.3	862.777	out-of-plane C=O deformation
967.74	970.8272	
1023.309	1023.309	
	1041.832	C-C-C stretches
1100.487	1109.749	
1168.8	1171.492	
	1220.886	CH <sub>2</sub> deformations + Amide III band
1237.5	1248.671	
1261.019		Amide III band
1279.542	1282.629	
1324.5	1332.024	C-N stretching amide III
1372.157	1378.331	C-O O-H combination bands
	1430.812	
1449.335	1455.51	N-H bonding
1507.991		N-H deformation and C-N stretching amide II
1514.165	1511.078	
1560.473	1560.473	C=C stretches
	1578.996	

1610.7	1612.954	
1618.5	1628.39	C=O stretches amide I
1648.8	1650	
	2733.589	carboxylic acid dimer
2931.9	2841.64	aliphatic C-H stretches
	2884.86	
	2937.341	
	2983.648	
3014.52	3014.52	
3055.2	3060.827	
3066.3	3079.35	aromatic C-H stretches
3110.222	3104.047	
	3178.139	O-H stretches
	3205.923	