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# **Supplementary information**

# Polymorphism of Vanillin Revisited: The Discovery and Selective Crystallization of a Rare Crystal Structure

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## X-ray Crystallography information

The X-ray diffraction data were measured on a Bruker D8 Venture PHOTON 100 CMOS system equipped with a Cu K $\alpha$  INCOATEC Imus micro-focus source ( $\lambda$  = 1.54178 Å). The crystal was kept at 100(2) K during data collection. Indexing was performed using APEX2.<sup>1</sup> (Difference Vectors method). Data integration and reduction were performed using SaintPlus 6.01.<sup>2</sup> Absorption correction was performed by multi-scan method implemented in SADABS.<sup>3</sup> Space groups were determined using XPREP implemented in APEX2.<sup>1</sup> The structure was solved using SHELXS-97 (direct methods) and refined using SHELXL-2013<sup>7</sup> (full-matrix least-squares on F2) contained in APEX2<sup>1,7</sup>, WinGX v1.70.01<sup>4-7</sup> and OLEX2.<sup>7,8</sup> All non-hydrogen atoms were refined anisotropically. H9 Hydrogen atom of hydroxyl group has been found from difference Fourier map and was freely refined with Uiso(H) = 1.5Ueq(-OH). The remaining hydrogen atoms of -OH groups as well as -CHO groups have been found from difference Fourier map and were freely refined. All other hydrogen atoms (of -CH, -CH2 and -CH3 groups) were placed in geometrically calculated positions and included in the refinement process using riding model with isotropic thermal parameters: Uiso(H) = 1.2Ueq(-CH,-CH2) and Uiso(H) = 1.5Ueq(-CH3).

Table S1. Hydrogen Bonds for vanillin Form II.

D	Н	A	d(D-H)/Å	d(H-A)/Å	d(D-A)/°	D-H-A/°
O9	Н9	O7i	1.07 (4)	1.69 (4)	2.681 (3)	151 (3)
O6	Н6	O3 <sup>ii</sup>	1.03 (5)	1.81 (5)	2.712 (3)	144 (4)
О8	Н8	O9 <sup>iii</sup>	0.97 (6)	1.96 (6)	2.806 (3)	144 (5)
O10	H10	O6iv	0.85 (5)	2.10 (5)	2.794 (3)	139 (4)

Symmetry operations: (i) -1/2+X,1/2-Y,+Z; (ii) 1/2+X,3/2-Y,+Z; (iii) 1/2-X,1/2+Y,1/2+Z; (iv) +X,-1+Y,+Z

Table S2. Crystal data and structure refinement for vanillin Form II

Empirical formula	C <sub>8</sub> H <sub>8</sub> O <sub>3</sub>		
Formula weight	152.14		
Temperature/K	100 (2)		
Crystal system	Orthorhombic		
	$Pna2_1$		
Space group a/Å	16.397 (7)		
b/Å			
c/Å	3.810 (1)		
	45.645 (2)		
α/°	90		
β/°	90		
γ/°	90		
Volume/Å <sup>3</sup>	2851.61 (19)		
Z	16		
Z'	4		
ρ calc mg/mm <sup>3</sup>	1.418		
$\mu/\text{mm}^{-1}$	0.918		
F(000)	1280.0		
Crystal size/mm <sup>3</sup>	$0.11 \times 0.04 \times 0.02$		
Radiation	$CuK\alpha (\lambda = 1.54178)$		
2θ range for data collection	7.748 to 144.68°		
Index ranges	$-20 \le h \le 20, -4 \le k \le 4, -56 \le 1 \le 54$		
Reflections collected	14594		
Independent reflections	5340 [Rint = 0.0521, Rsigma = 0.0485]		
Data/restraints/parameters	5340/1/432		
Goodness-of-fit on F2	1.028		
Final R indexes [I>=2σ (I)]	R1 = 0.0387, $wR2 = 0.0887$		
Final R indexes [all data]	R1 = 0.0465, $wR2 = 0.0925$		
Largest diff. peak/hole / e Å <sup>-3</sup>	0.18/-0.23		
Flack parameter 0.12 (10)	Flack parameter 0.12 (10)		

Fig. S1: Comparison of calculated PXRD patterns of vanillin Form I crystal structures reported by Natarajan et al. (YUHTEA) and Nieger (YUHTEA01).

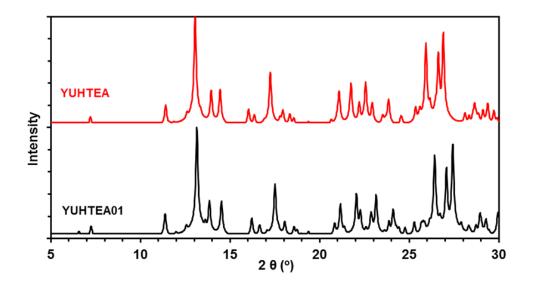


Fig. S2: PXRD patterns of calculated FI and FII of vanillin and it comparision to the PXRD patterns of vanillin crystals obtained from Isopropy alcohol, water, ethanol, acetonitrile, acetone.

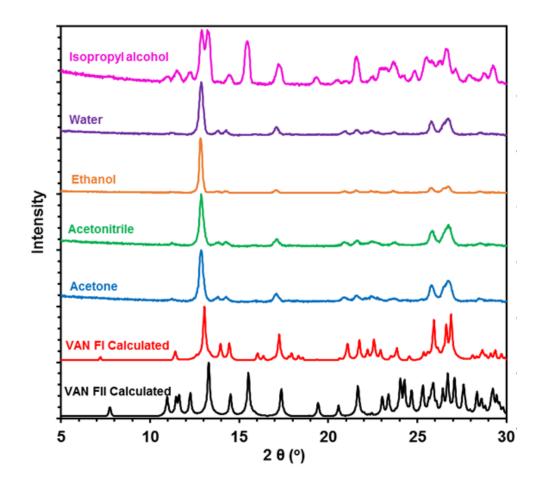


Fig. S3: Illustration of (a) hydrogen bonding in vanillin Form I; (b) lateral interactions in vanillin tapes in Form I (CSD ref code: YUHTEA).

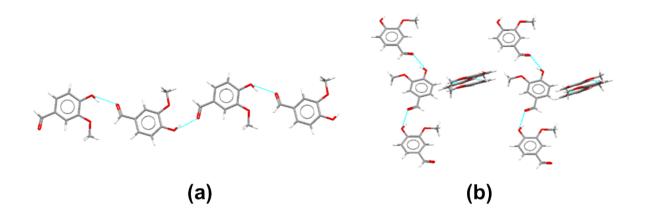


Fig. S4: DSC traces showing melting points of vanillin Forms I (red) and Form II (green).

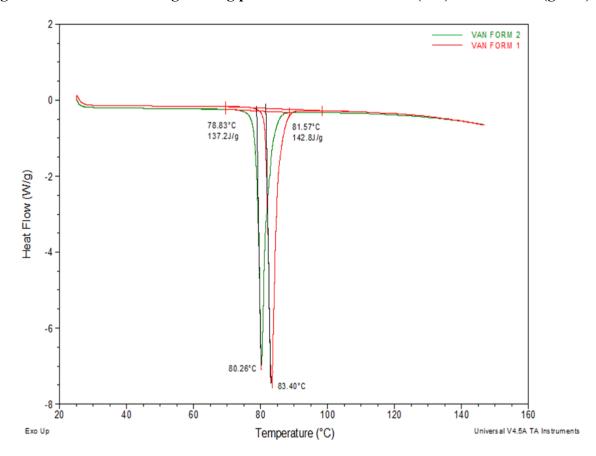


Fig. S5: PXRD patterns of vanillin crystals obtained from slow evaporation of Isopropy alcohol in presence of PAA, PMMA, and PEG.

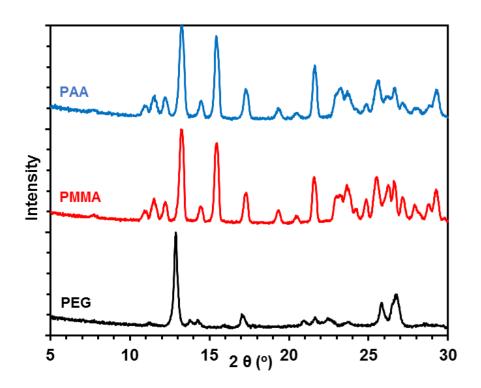


Fig. S6: SEM image of vanillin Form I crystallized in presence of PEG by slow evaporation

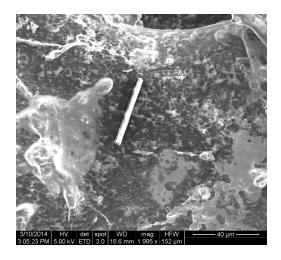
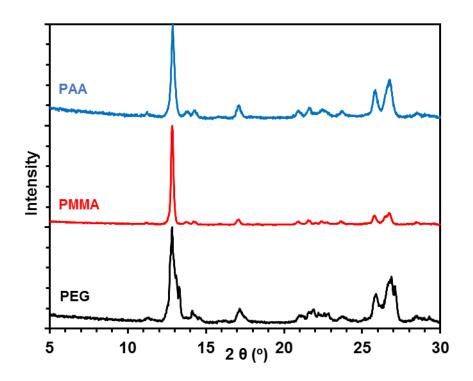


Fig. S7: PXRD patterns of vanillin crystals obtained by cooling crystallization from Isopropy alcohol in presence of PAA, PMMA, and PEG.



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