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How elastically flexible can molecular crystals be? – a new record

During the gradient-cooling melt crystallization process, we obtained single crystals of CEL form I, which exhibited a record-high elastic flexibility of up to 8.7% when bent along the (001) crystal face, attributed to the surface tension of the melt. This exceptional flexibility is driven by the presence of multiple long-range dispersive interactions between (001) slip planes, as well as the rough topology of these planes, which hinders the onset of plastic slip during bending.

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### How elastically flexible can molecular crystals be? – a new record<sup>+</sup>

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The elastic strain limit, which quantifies the elastic flexibility of a material, is critical for technological applications of functional materials in a number of fields. Although the elastic flexibility of molecular crystals has been recognized, the extent of elastic flexibility of such materials remains to be defined. Here, we report a molecular crystal, *i.e.*, form I polymorph of celecoxib (CEL), exhibiting exceptional elastic flexibility with an elastic strain of at least 8.70%. The record high elastic strain is accompanied by low Young's modulus ( $E = 3.18 \pm 1.01$  GPa) and hardness ( $H = 39.8 \pm 15.6$  MPa), as determined by single crystal nanoindentation, along with the high plasticity of the bulk powder observed in in-die Heckel analysis.

Extensive efforts have been devoted to designing and synthesizing mechanically flexible molecular crystals,1-13 because of their potential applications in a wide range of technological areas, including pharmaceuticals,<sup>14,15</sup> sensors,<sup>16,17</sup> electronics,18,19 optical waveguides,20,21 and wearable devices.22,23 It was projected that the flexible electronics market would reach a value of 70 billion US dollars by 2026.24 An elastically deformed crystal spontaneously regains its original shape once the applied stress is removed. However, irreversible deformation, either plastic deformation or brittle fracture, takes place once a certain elastic strain is exceeded. Thus, the maximum elastic strain determines the elastic flexibility of a single crystal. which ultimately determines the maximum flexibility of the device employing such crystals. Hence, knowledge of the elastic limit of an elastic crystal is critically important for its successful application in various areas.25

Elastic strain can be calculated from specimen thickness and the radius of curvature using Euler–Bernoulli beam-bending theory.<sup>26,27</sup> The maximum elastic strain of known elastically flexible organic crystals typically ranges from 1% to 3%, with only a handful of cases reaching 6%.<sup>28-33</sup> The most commonly adopted three-point bending method, performed using forceps and a needle for studying the elastic flexibility of molecular crystals, requires large crystals (typically mm in length) and physical contact with a stiff tool, which likely introduce defects and cause premature failure before reaching the theoretical elastic limit of the crystal.<sup>34,35</sup> Here, using a contact-free bending method performed during melt crystallization, we show that an elastic strain of at least 8.70% could be attained for the form I polymorph of a nonsteroidal anti-inflammatory drug (NSAID), celecoxib (CEL), setting a new record of elastic strain for molecular crystals.

CEL has been widely used to treat acute pain since 1998.<sup>36</sup> To date, four polymorphs of CEL, forms I, II, III, and IV, have been discovered.<sup>37</sup> Form III is the polymorph present in commercial capsule products due to its thermodynamic stability under ambient conditions. Despite the extensive research performed on CEL, only the form III single crystal structure has been reported in the crystal structure database (CSD), likely because of the difficulty of growing sufficiently large single crystals of other polymorphs of CEL using the solution crystallization process. Form I CEL can crystallize from an amorphous film when stored overnight at 80 °C (Fig. 1a).<sup>38</sup> However, the form I crystals thus obtained were too small with low quality for structure elucidation by single crystal X-ray diffraction (SCXRD) analysis. Hence, we attempted the melt crystallization method to grow large form I CEL single crystals for structure elucidation by SCXRD.

The melt crystallization technique, which is material-sparing (~10 µg per experiment), has been used for preparing polymorphs that are thermodynamically unstable near ambient temperature, *e.g.*, indomethacin, ROY, and piroxicam.<sup>39–43</sup> The temperature suitable for single crystal growth is usually within the range of 0.97–0.99 times the melting temperature ( $T_{\rm m}$ ).<sup>44</sup> However, in that temperature range, uncontrolled late growth of curled fine CEL form I crystals covered the surface of the initial single crystal before it grew sufficiently large for SCXRD. To overcome this problem, we developed a gradient-cooling melt crystallization method, which involves the following steps: (1)

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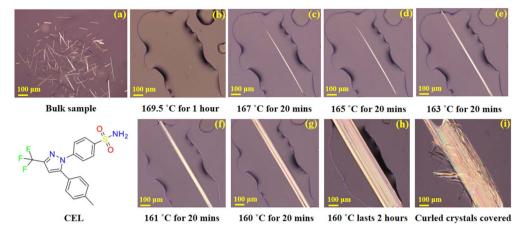


Fig. 1 Single crystal growth of CEL form I by the gradient-cooling melt crystallization method. Scale bars are 100  $\mu$ m. (a) Amorphous CEL with form I seeds, (b) sample held at 169.5 °C till only a single form I CEL seed remaining, (c–h) sample held at gradually lowered temperatures, from 167 °C to 160 °C, to facilitate single crystal growth, (i) a single crystal covered with fine curled crystals, grown from remaining melt, after being held at 160 °C much longer than 2 hours.

a small sample of amorphous CEL containing form I crystal seeds (Fig. 1a) was prepared by storing an amorphous CEL film overnight at 80 °C.38 This was heated on a glass slide at 10 °C  $min^{-1}$  to 169.5 °C, close to the end of the form I melting range observed from a DSC thermogram (Fig. S1<sup>†</sup>), on a hot stage, and held at this temperature until only one seed crystal remained (Fig. 1b). (2) The temperature was lowered to 167 °C and held for 20 min (Fig. 1c), followed by lowering the temperature to 165 °C, 163 °C, and 161 °C sequentially (held for 20 min at each temperature, Fig. 1d-f) and finally to 160 °C (held for 1-2 hours, Fig. 1g and h) to obtain a single crystal with a sufficient size. This cooling program was intended to maintain the growth of the single crystal while avoiding the generation of new crystal nuclei invariably encountered during single step cooling. However, even with this gradient program, the growth of an acicular crystal was very slow once the fast-growing ends of the crystal exceeded the boundary of the melt (Fig. 1h). With a long

holding time, thin crystals would always form and cover the surface of the single crystal before all the liquid was consumed by single crystal growth (Fig. 1i). Directly isolating the single crystal from the melt would still cause crystallization of the residual melt into fine crystals. To overcome this problem, a drop of silicone oil was placed to overlap with the remaining melt at 160 °C, and then the single crystal was isolated from the remaining melt by pushing it into the oil with a needle. By following this procedure, high-quality acicular single crystals (1–2 mm long and 0.05–0.1 mm wide) suitable for crystal structure elucidation by SCXRD were successfully prepared (Fig. S2†).

CEL form I crystallizes in the  $P\bar{1}$  space group with Z' = 3 (Table S1<sup>†</sup>), belonging to the rare class of structures with Z' = 3 (total 6387 crystal structures, corresponding to 0.52% of all structures in the CSD), out of which only 30.5% are in the  $P\bar{1}$  space group.<sup>45</sup> The unit cell parameters and global crystal

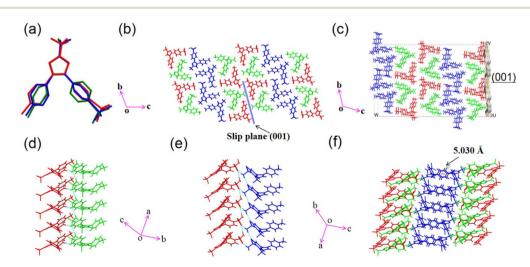


Fig. 2 Crystal structure analysis of CEL form I: (a) the overlay of three molecules in the asymmetric unit showing conformational differences; (b) the molecular arrangement viewed along the *a*-axis; (c) calculated surface contour of the (001) slip plane; (d) hydrogen bonding involving one pair of CEL molecules; (e) hydrogen bonding involving the other pair of molecules; (f) packing of three molecules.

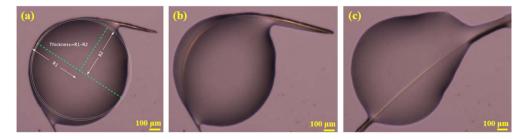


Fig. 3 CEL form I: (a) a bent crystal (two circles were drawn to fit the outer and inner side of the bent region, from which the crystal thickness was calculated); (b) elastic recovery in the process; (c) the crystal with shape recovered.

packing structures of CEL forms I differ from those of form III (Fig. S3 and Table S1<sup>†</sup>). There are three crystallographically unique molecules in the asymmetric unit (labelled with different colors in Fig. 2a), which exhibit significantly different conformations owing to the rotation of 4-methylphenyl (dihedral angles of 25.80°, 29.71°, and 61.12°) and benzenesulfonamide (dihedral angles of 59.18°, 60.80°, and 33.83°) rings about the trifluoromethyl-1*H*-pyrazole ring (Fig. S4a-c<sup>†</sup>). The three molecules in the asymmetric unit stack along the a-axis (Fig. S4d<sup>†</sup>) to form molecular columns, which are fortified through several types of weak intermolecular interactions, such as C-H…F (3.266 Å and 138.30°, Table S2†) and  $\pi \dots \pi$  stacking (centroid-to-centroid distance = 5.030 Å, Fig. 2f). The plane defined by the 4-methylphenyl and trifluoromethyl-1H-pyrazole rings in CEL form III is not observed in the form I structure,46 where the dihedral angles are all greater than 25° in form I but less than 18° in form III. The (001) plane is identified as a slip plane (Fig. 2b) with an uneven surface contour (Fig. 2c). The three conformationally different CEL molecules form six unique

types of hydrogen-bonded tetramers (Fig. S5†), which can be classified into two categories based on their compositions: (a) an asymmetric unit plus one CEL molecule from a stacking asymmetric unit and (b) a pair of CEL molecules from the asymmetric unit with the same pair from a stacking asymmetric unit (Fig. S6†). In contrast, CEL form III only forms one type of dimer in the crystal structure. The different types of tetramers are interconnected through a number of hydrogen bonds (Table S2†). A one-dimensional (1D) chain along the *a*-axis (Fig. S7a†) is formed by stacking tetramers, fortified *via* weak  $\pi \cdots \pi$  (4.953 Å, centroid-to centroid), C–H $\cdots \pi$  (3.347 Å), and N–H $\cdots$ O (*d*/ $\theta$ : 2.961 Å/170.57° and 2.858 Å/160.83°) interactions. Adjacent 1D columns connect *via*  $\pi \cdots \pi$  (5.030 Å, centroid-to-centroid) interactions between two aromatic rings to form a 2D layer of CEL molecules (Fig. S7b†).

The dominant faces of the form I CEL crystal are (001)/(00-1), with the side faces running along the long axis of the crystal being (010)/(0-10) and the capping ends being (100)/(-100) (Fig. S8†). Interestingly, some of the CEL form I single

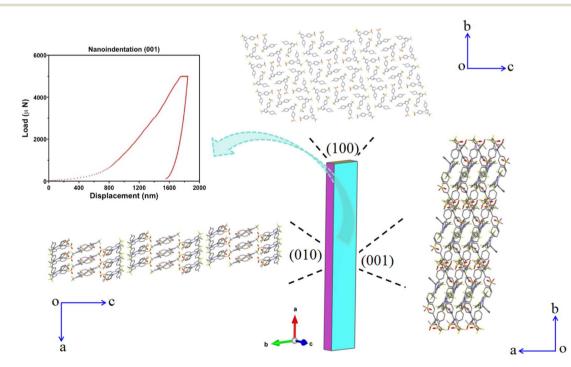


Fig. 4 Packing diagrams of CEL form I viewed along the (100), (001) and (010) crystal faces (with hydrogens omitted for clarity), accompanied by a nanoindentation load-displacement curve for the (001) crystal face.

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crystals bent when growing from a melt at 160–167 °C (Fig. 3a). These crystals straightened up when the viscosity of the melt was sufficiently decreased as the temperature was raised to 167–168 °C (Fig. 3c, Video S1†), indicating the elastic nature of the deformation. The elastic strains of different bent crystals ranged from 0.54 to 8.7% (Fig. S9, S10 and Table S3†), calculated from the crystal thickness and diameter of the curvature arc based on Euler–Bernoulli beam theory. Thus, the maximum elastic strain of CEL form I is at least 8.7%, which is significantly higher than those of all reported elastic molecular crystals underwent brittle fracture when the (010) face was bent during crystal mounting for SCXRD (Fig. S2†). Thus, similar to CEL form III, form I CEL is 1D elastic.

The Young's modulus ( $E = 3.18 \pm 1.01$  GPa) and hardness (H= 39.8  $\pm$  15.6 MPa) of the elastically bendable (001)/(00-1) crystal face determined by nanoindentation (Fig. 4 and S11<sup>†</sup>) are much lower than those of CEL form III ( $E = 16.27 \pm 0.43$  GPa and  $H = 450 \pm 20$  MPa, (001)/(00-1) crystal face).<sup>46</sup> The experimentally measured room temperature true density of form I  $(1.4758 \text{ g cm}^{-3})$  is ~3.28% lower than that of form III (1.526 g cm<sup>-3</sup>).<sup>47</sup> Thus, the greater elastic flexibility of CEL form I is accompanied by both lower stiffness and lower density. Moreover, the form I CEL bulk powder also exhibits greater plasticity, as shown by its lower in-die mean yield pressure (P<sub>y,i</sub> = 41.7  $\pm$ 3.0 MPa) than that of form III powder ( $P_{v,i} = 74.6 \pm 3.6$  MPa) (Table S4 and Fig. S13, S14<sup>+</sup>). The lower stiffness of form I, confirmed by both single crystal nanoindentation and bulk powder Heckel analysis, is consistent with its higher elastic flexibility than form III.

In order to better understand the lower stiffness and higher elastic flexibility of CEL form I, we performed an integrated energy framework and topology analysis.48 CEL molecular columns, running along the a-axis and strengthened by numerous C-H···O, N-H···O, and C-H···F interactions, are stabilized by  $\pi \cdots \pi$  and C-H $\cdots \pi$  inter-columnar interactions, resulting in a highly anisotropic energy framework. The energy frameworks of CEL form I and III differ significantly, where it appears discontinuous in form I (Fig. S15<sup>†</sup>), but continuous in form III (Fig. S16<sup>†</sup>). The (001) slip plane based on visual analysis of the crystal structure (Fig. 2b), i.e., on the basis of flat molecular layers with only van der Waals type inter-layer interactions, is confirmed by the integrated energy framework and full interaction map (FIM, Fig. S17<sup>†</sup>). The (001) slip plane in CEL form III exhibits a rather different FIM than that in form I, corresponding to their different crystal packing patterns and intermolecular interactions. In addition, the attachment energy for the slip plane (001) in form I ( $-3.165 \text{ kJ mol}^{-1}$ ) is significantly weaker than that in form III  $(-28.077 \text{ kJ mol}^{-1})$ , while the lattice energy of form I is also weaker than that of form III (Table S5<sup>†</sup>), all of which is consistent with the lower density, lower stiffness, higher plasticity, and higher elastic flexibility of form I. The existence of multiple long-acting dispersive interactions between (001) slip planes, which is also the elastically bending crystal face, and the rough topology of the slip planes that hinders the onset of plastic slip during bending explain the significantly higher elastic flexibility of form I than form III.49

Among the four known mechanisms for the elastic flexibility of molecular crystals,<sup>50</sup> the reversible molecular rotation model<sup>8</sup> best explains the exceptional elasticity of CEL form I crystals. Its structure lacks key features typically associated with other mechanisms, such as interlocked stacking layers,<sup>51</sup> fibril lamellae,<sup>52</sup> and layered nanorods.<sup>53</sup>

In summary, using the gradient-cooling melt crystallization method, we successfully obtained a high-quality single crystal of CEL form I and solved its structure 27 years after its first discovery. The correspondence between the record-high elastic flexibility and low crystal stiffness, if proven robust with more systems, suggests the possible use of low stiffness as one of the criteria for screening elastically flexible molecular crystals that otherwise could have been overlooked because of the difficulty of forming large acicular crystals required for performing the classical three point bending test.

#### Data availability

The data supporting this article have been included as part of the ESI.†

#### Author contributions

Zhengzheng Zhou: formal analysis, investigation, writing – original draft, visualization, and project administration; Vikram Chandrashekhar Joshi: single crystal X-ray diffraction, nano-indentation data collection, and writing – review & editing; Yiwang Guo: nanoindentation and mean yield pressure  $(P_y)$  data collection; Tianyi Xiang: single crystal X-ray diffraction data collection; Zijian Wang: scanning electron microscope data collection; Changquan Calvin Sun: conceptualization, resources, writing – review & editing, supervision, project administration, and funding acquisition.

#### Conflicts of interest

The authors declare that they have no known competing financial interests or personal relationships that will influence this work.

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