

## Supplementary Information

## Crystal Structures, Thermal Analysis and Dissolution Behavior of New Solid Forms of the Antiviral Drug Arbidol with Dicarboxylic Acids

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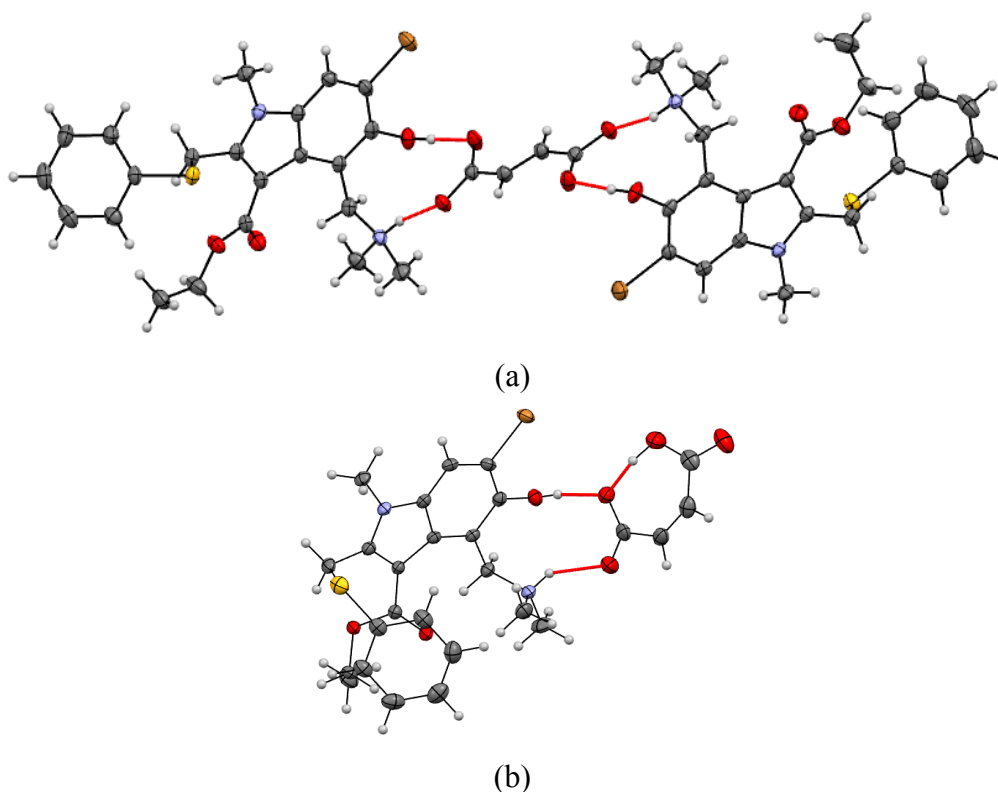
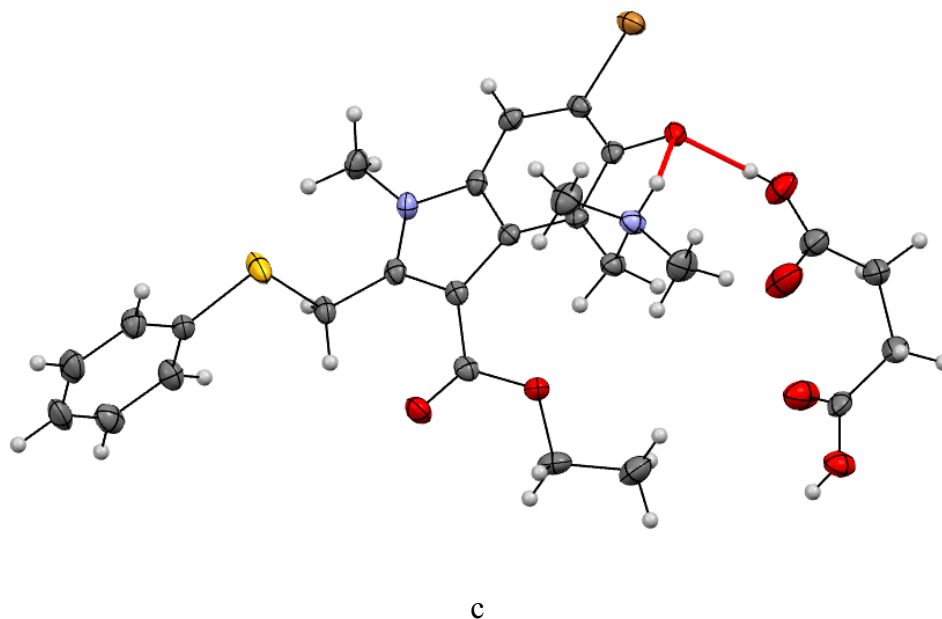
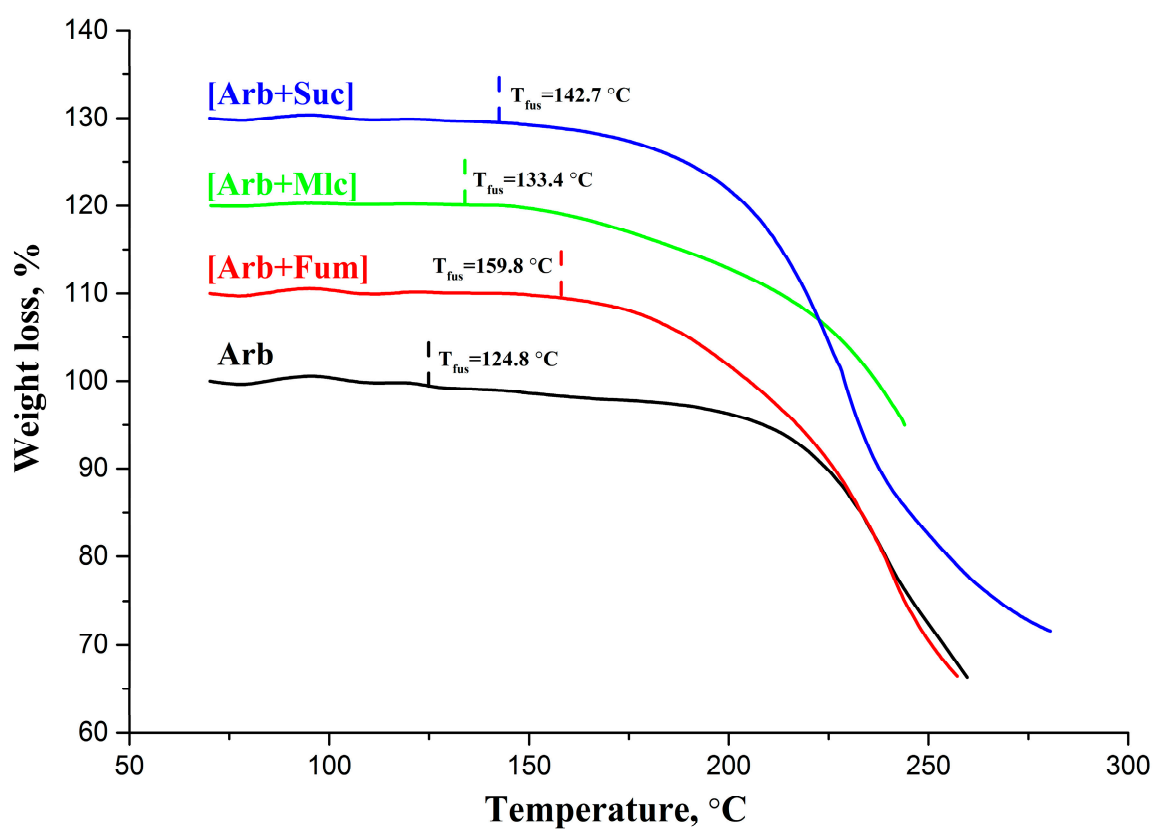


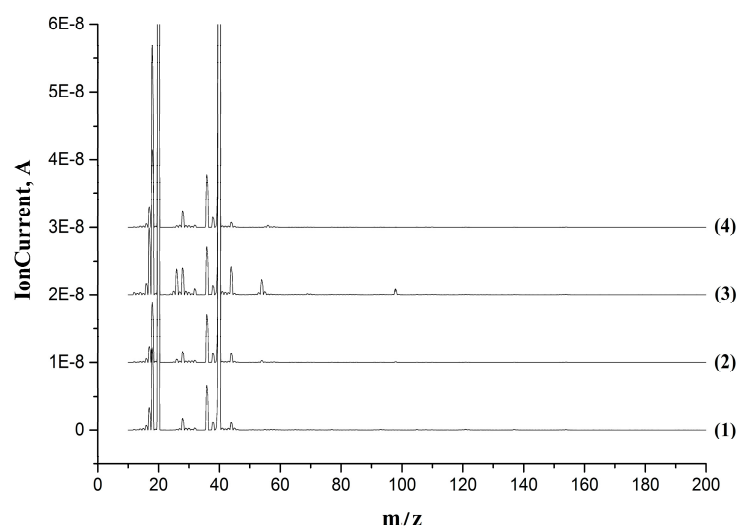
Figure S1. Cont.



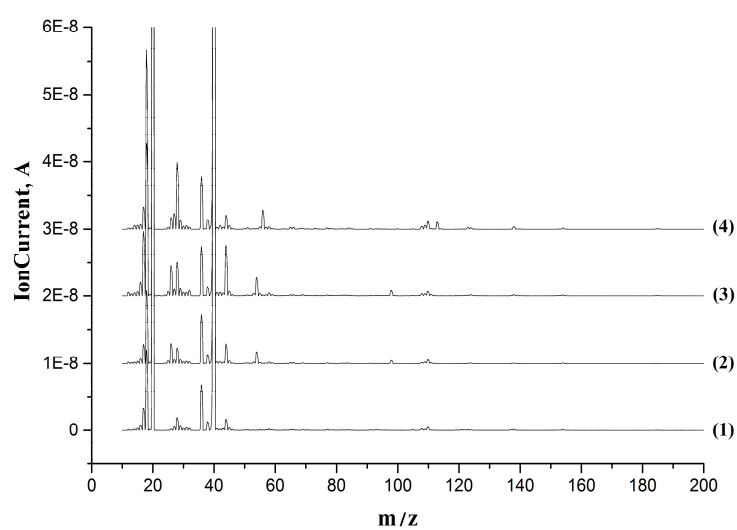
**Figure S1.** Asymmetric units in (a) [Arb + Fum], (b) [Arb + Mlc], (c) [Arb + Suc]. Displacement ellipsoids are shown at 50% probability.



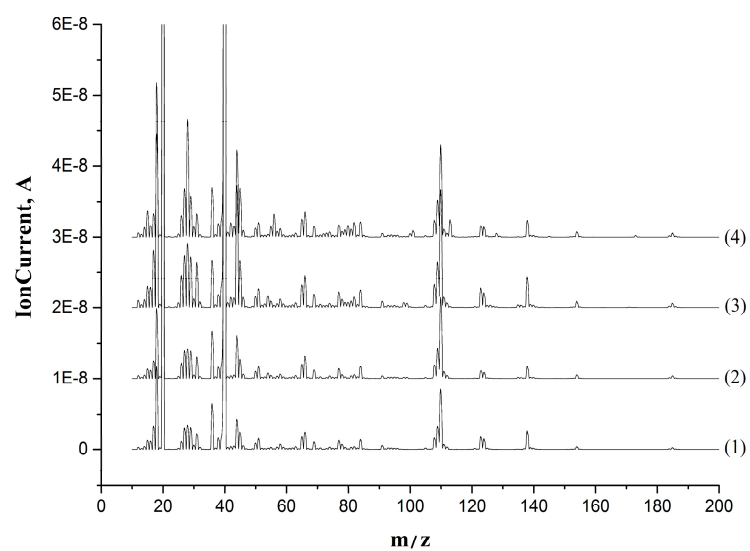
**Figure S2.** TG thermograms of **Arb** (black line); [**Arb + Fum**] (red line); [**Arb + Mlc**] (green line) and [**Arb + Suc**] (blue line)



(a)

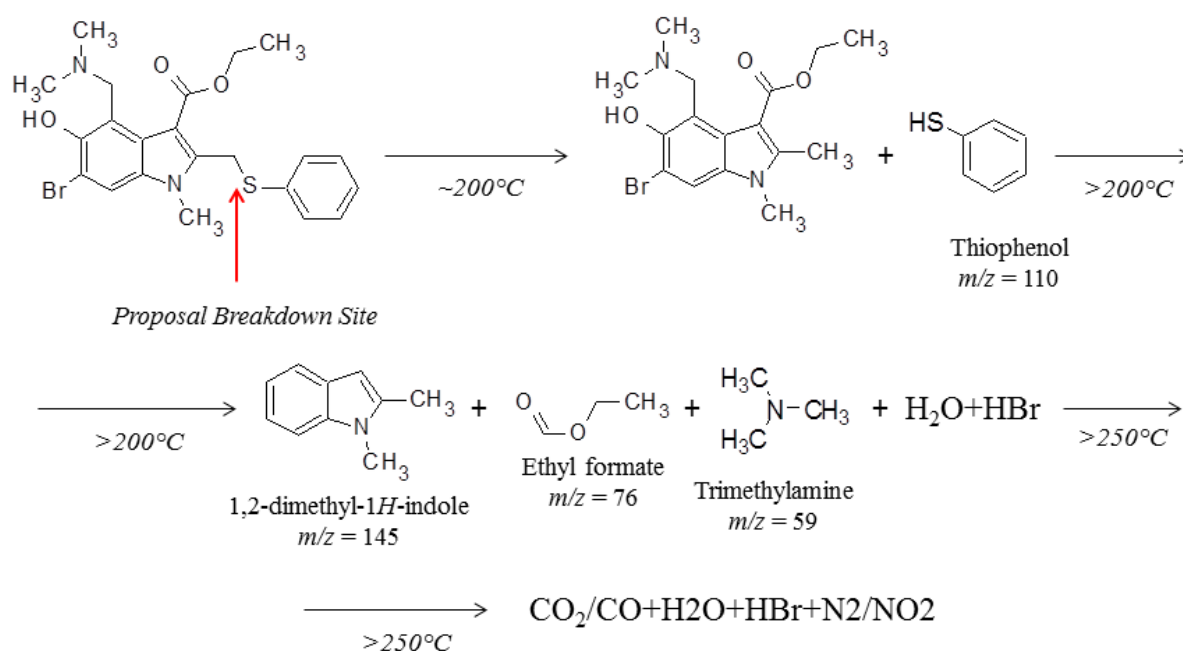


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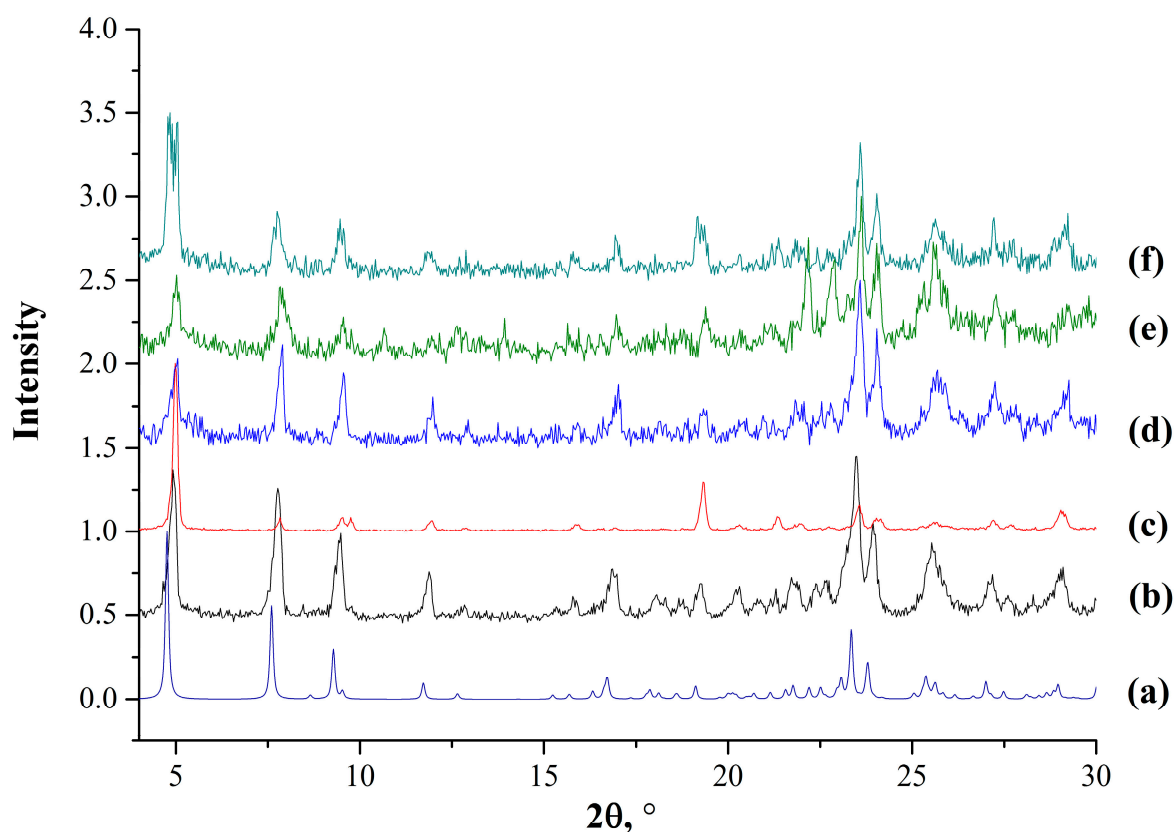


(c)

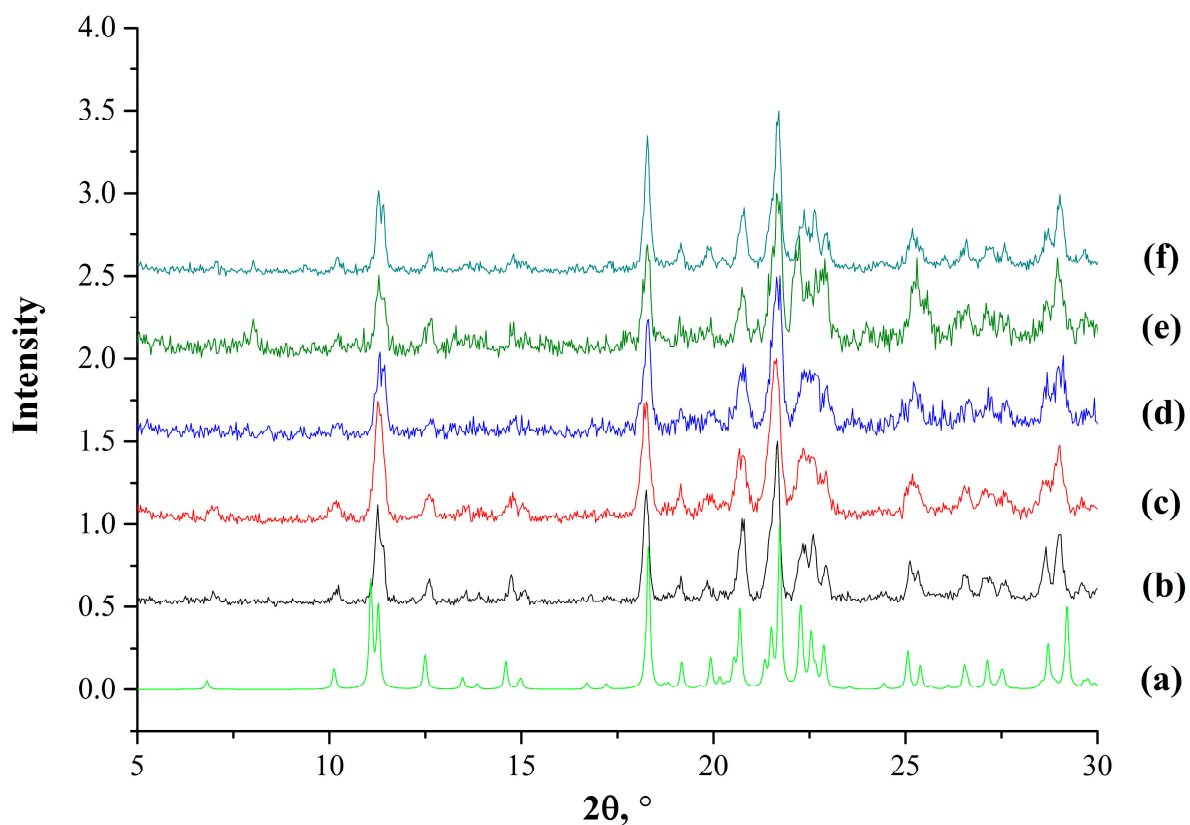
**Figure S3.** Mass spectrometric curves for **Arb** (1); [**Arb** + **Fum**] (2); [**Arb** + **Mlc**] (3); [**Arb** + **Suc**] (4) at different temperature: (a) at 175 °C; (b) at 205 °C; and (c) at 245 °C.



**Scheme S1.** Proposed thermal degradation of Arb.



**Figure S4.** XRPD analysis of: (a) simulated pattern of [Arb + HCl + H<sub>2</sub>O] based on the SAJPUQ crystal structure; the residual materials after solubility of (b) arbidol base, (c) [Arb + HCl + H<sub>2</sub>O], (d) [Arb + Fum], (e) [Arb + Mlc], and (f) [Arb + Suc] in pH 1.2 solution.



**Figure S5.** XRPD analysis of: (a) simulated pattern of arbidol base (ref. code SAJPOK) and the residual materials after solubility of (b) arbidol base, (c) [Arb + HCl + H<sub>2</sub>O], (d) [Arb + Fum], (e) [Arb + Mlc], and (f) [Arb + Suc] in pH 6.8 solution.