

Supplementary Information

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

Alex N. Manin <sup>1</sup>, Artem O. Surov <sup>1,2</sup>, Andrei V. Churakov <sup>3</sup> and German L. Perlovich <sup>1,2,\*</sup>

<sup>1</sup> Department of Physical Chemistry of Drugs, Institution of Russian Academy of Sciences, G.A. Krestov Institute of Solution Chemistry of the Russian Academy of Sciences, Ivanovo 153045, Russia; E-Mails: alexnmanin@gmail.com (A.N.M.); artsurov@yandex.ru (A.O.S.)

<sup>2</sup> Department of Chemistry, Lomonosov Moscow State University, 1-3 Leninskiye gory, Moscow 119991, Russia

<sup>3</sup> Department of Crystal Chemistry and X-ray Diffraction Analysis, Kurnakov Institute of General and Inorganic Chemistry of the Russian Academy of Sciences, Leninskii Prospekt 31, Moscow 119991, Russia; E-Mail: churakov@igis.ras.ru

\* Author to whom correspondence should be addressed; E-Mail: germanper@yandex.ru; Tel.: +7-4932-533-784; Fax: +7-4932-336-237.

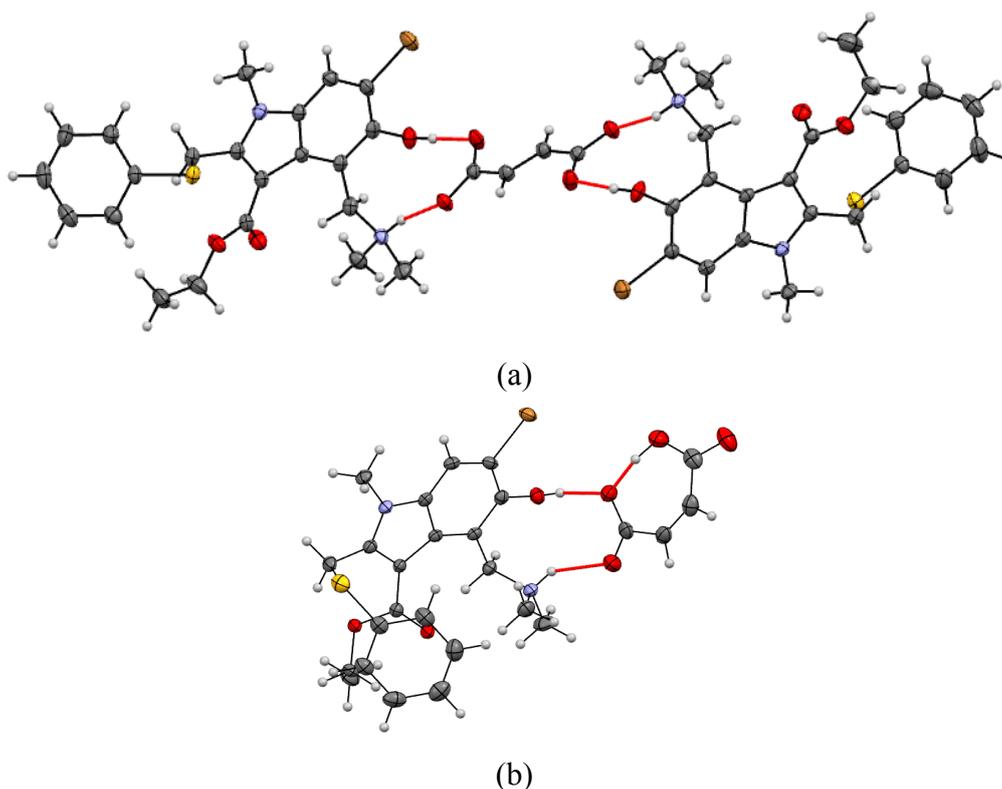
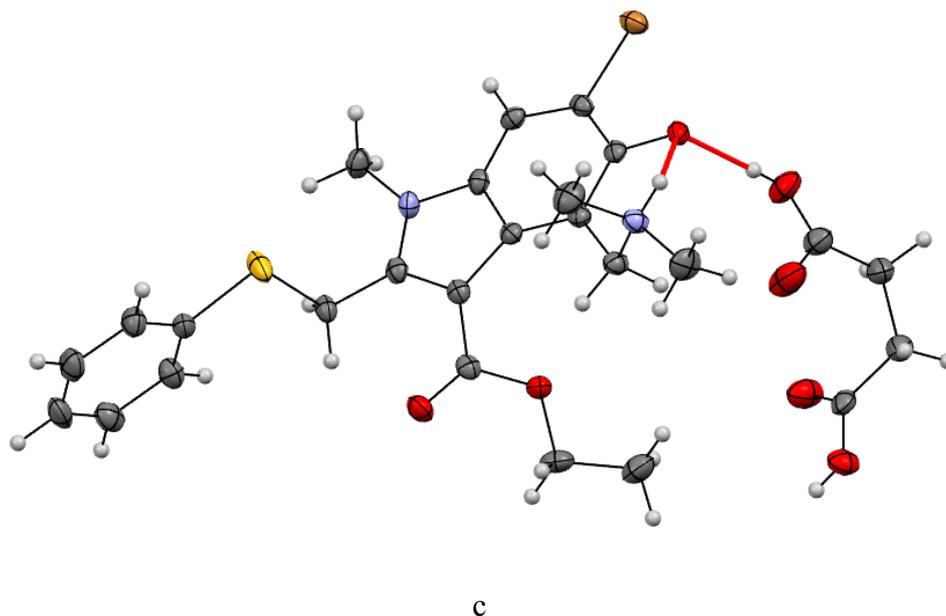
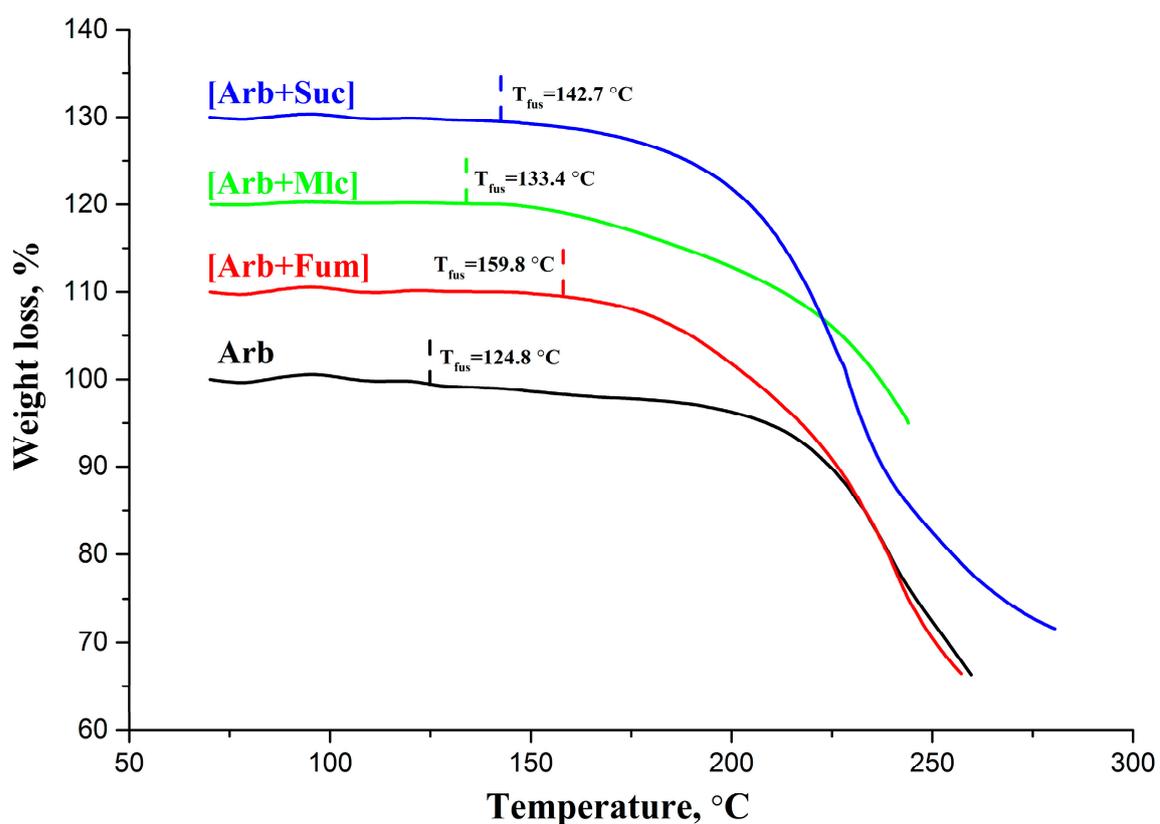


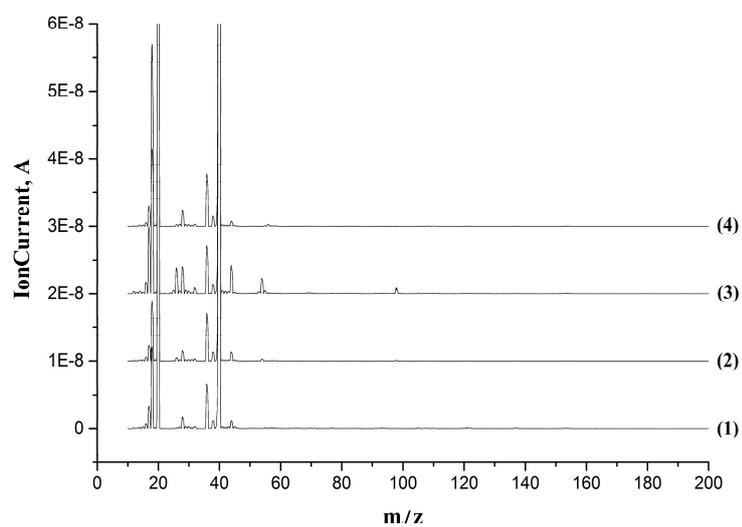
Figure S1. Cont.



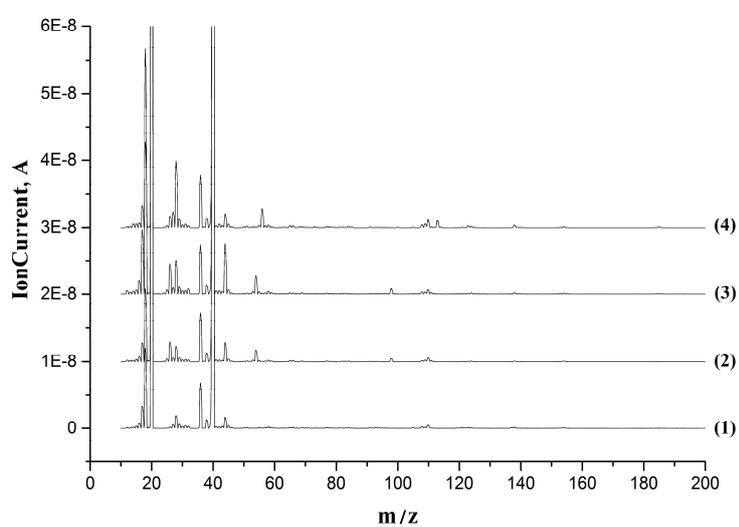
**Figure S1.** Asymmetric units in (a) [Arb + Fum], (b) [Arb + Mlc], (c) [Arb + Suc]. Displacement ellipsoids are shown at 50% probability.



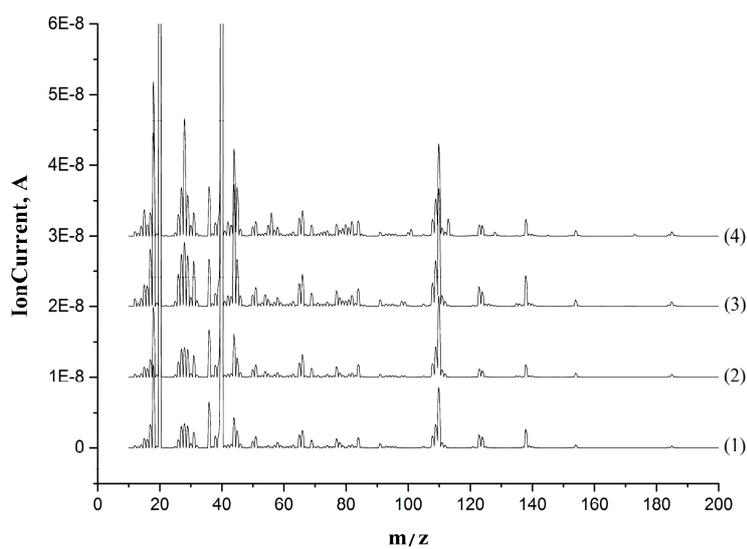
**Figure S2.** TG thermograms of Arb (black line); [Arb + Fum] (red line); [Arb + Mlc] (green line) and [Arb + Suc] (blue line)



(a)

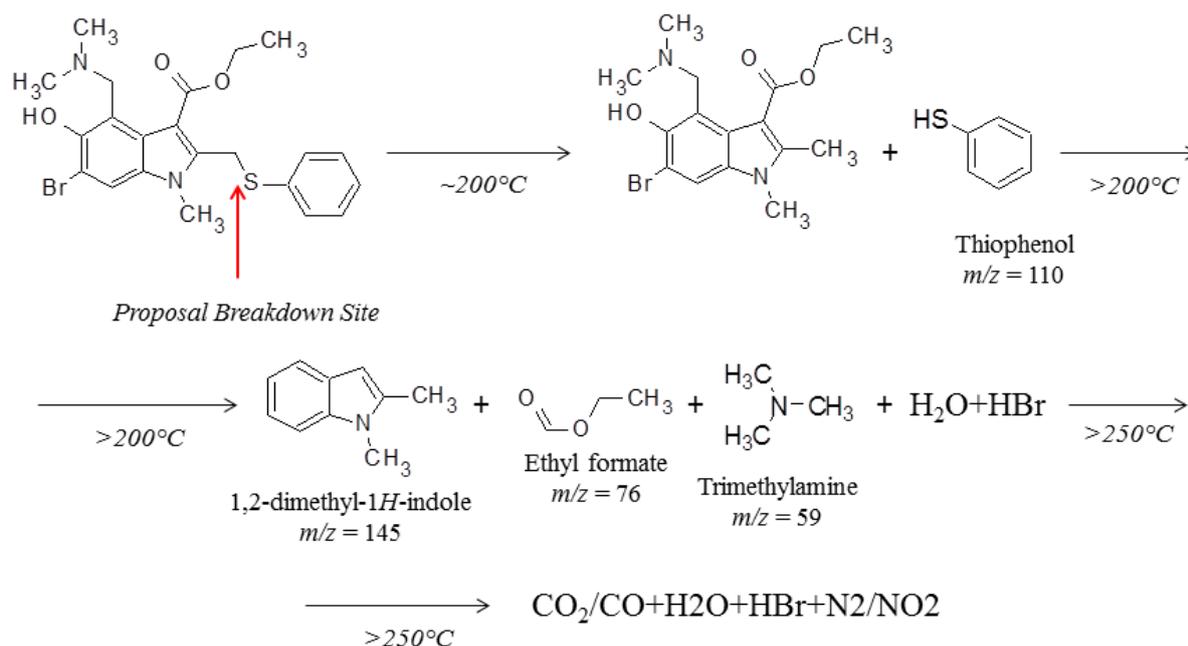


(b)

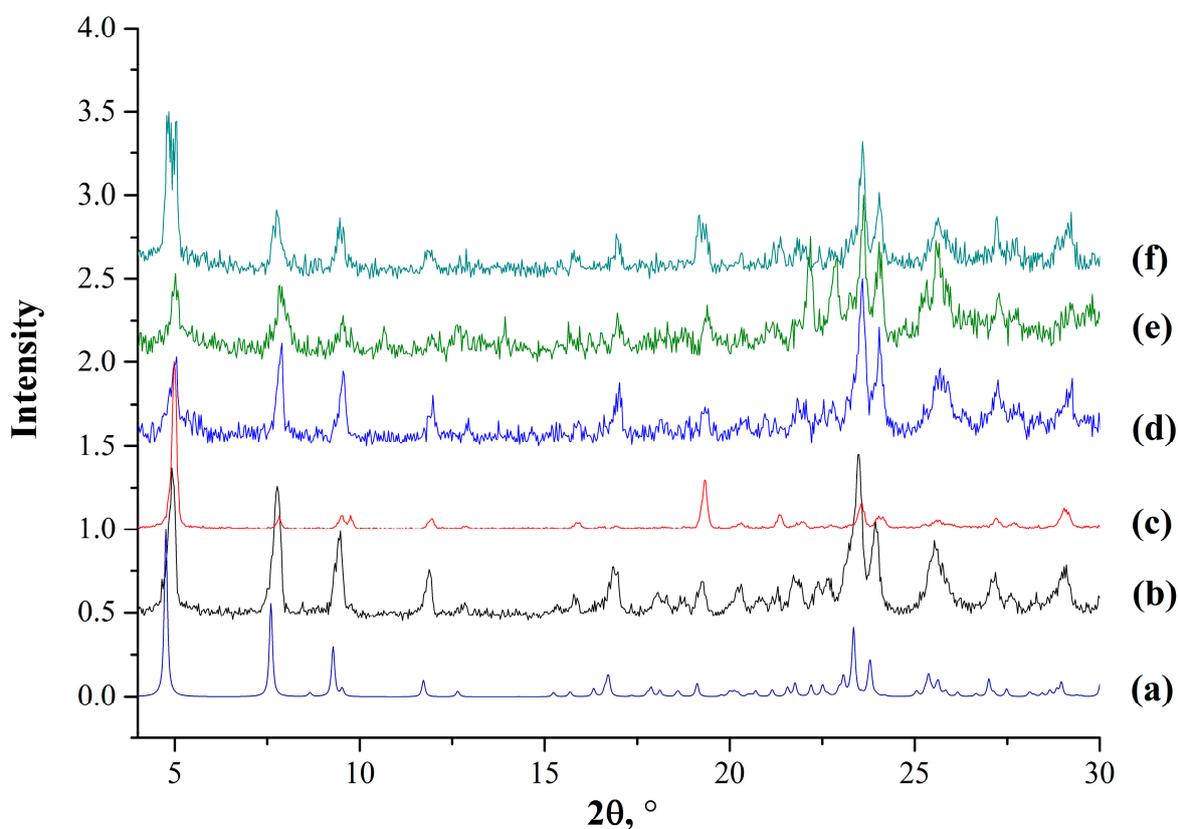


(c)

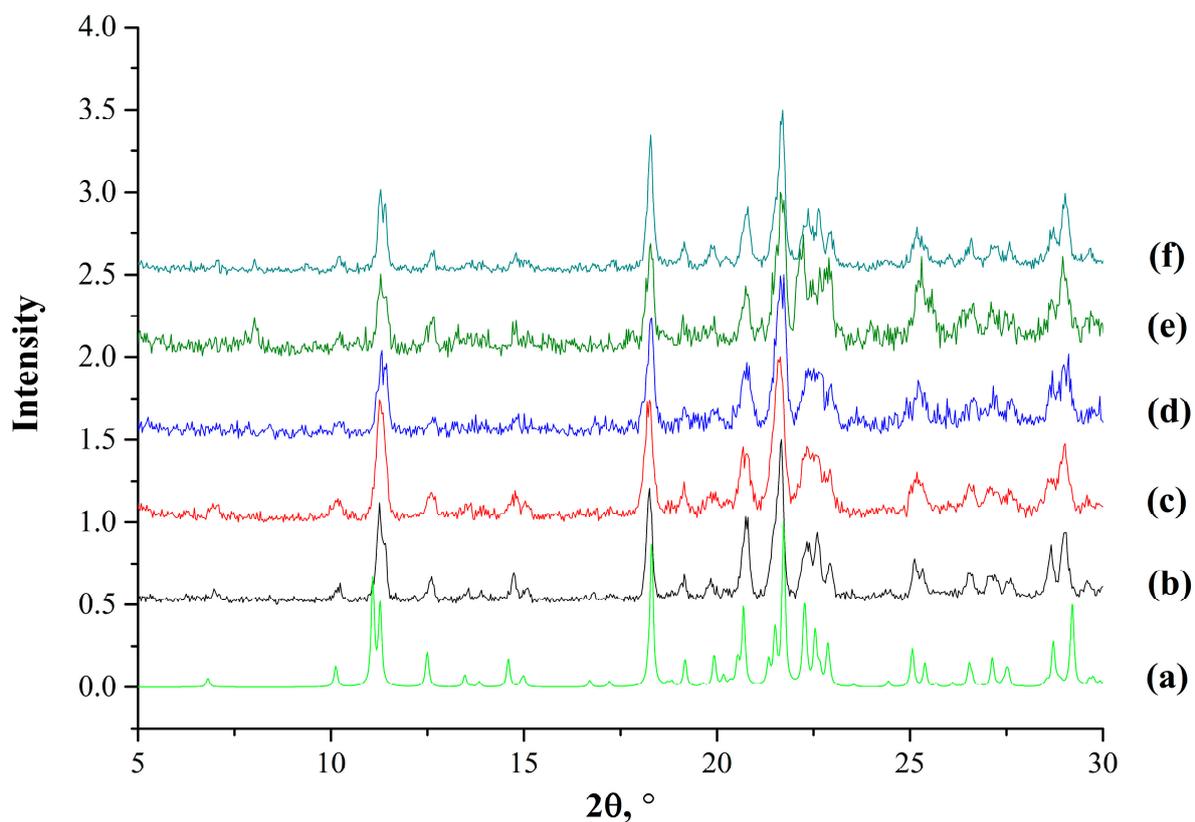
**Figure S3.** Mass spectrometric curves for **Arb (1)**; [**Arb + Fum**] (**2**); [**Arb + Mlc**] (**3**); [**Arb + Suc**] (**4**) at different temperature: **(a)** at 175 °C; **(b)** at 205 °C; and **(c)** at 245 °C.



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.

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