# SWEET SULFAMETHAZINE ACESULFAMATE CRYSTALS

# WITH IMPROVED COMPACTION PROPERTY

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## Abstract

Sulfamethazine (SMT) is a sulfonamide antibacterial drug used to treat or prevent infections in both humans and animals. However, SMT has an unfavorable taste and poor compaction behavior. To overcome these problems, a 1:1 complex with an artificial sweetener, acesulfame (Acs), was prepared and characterized. The single crystal structure suggests that the new complex, SMT-Acs, is a salt. This was confirmed by analysis of C-N bond length and comparison to multicomponent SMT crystals with known ionization states of SMT and Fourier transformation infrared spectroscopy. The applicability of the  $\Delta p K_a$  rule in multicomponent crystals of SMT is discussed. SMT-Acs exhibits better tabletability than SMT, which is attributed to its greater plasticity as shown by Heckel and Kuentz -Leuenberger analysis. The greater plasticity of SMT-Acs is consistent with the presence of slip planes identified by combined energy framework and topological analysis of the crystal structure.

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CHAPTER 1.

**INTRODUCTION** 

# **1.1 General Introduction**

The tablet is the most commonly used dosage form for oral drug delivery.<sup>1</sup> It has advantages over other dosage forms, including 1) accurate dose; 2) generally better chemical, physical and microbiological stability; 3) lower cost for manufacturing and distribution; 4) ease of administration; and 5) better patient compliance.<sup>2-4</sup> In order to develop a successful tablet product, several quality standards have to be met to ensure the efficacy and safety of the drug product. Adequate tablet mechanical strength is one of the key factors that needs to be considered.<sup>5</sup>

Crystal engineering is the rational design of fundamental molecular solids.<sup>6</sup> It is of both fundamental and practical interest to scientists in multiple fields, such as chemistry, material science, and pharmaceutical sciences.<sup>7</sup> Specifically in pharmaceutical sciences, crystal engineering has gradually become a successful design strategy to obtain novel multicomponent complexes of active pharmaceutical ingredients (APIs) with desired physicochemical properties but without affecting their pharmacological performances.<sup>8</sup> Numerous recent examples have amply shown the applicability of crystal engineering in improving mechanical properties, stability, solubility, and bioavailability of APIs.<sup>9-12</sup>

Understanding the relationship between crystal packing and the mechanical properties of the crystals is of interest in the fields of solid-state chemistry and crystal engineering.<sup>9, 10, 12, 13</sup> In pharmaceutical sciences, the structure-property relationship provides a foundation for understanding different tableting behaviors of

APIs.<sup>12, 14-16</sup> Pursuit of greater applicability of this relationship is now aided by computational tools, such as energy framework calculation and topology analysis, which can bring the goal of accurately predicting the compaction behaviors of crystals closer.<sup>17, 18</sup>

# 1.2 The Tablet

The tablet originated in mid 1800s and has been extensively investigated and widely used for many APIs.<sup>19</sup> Based on the intended clinical use, several types of tablets have been designed, including: 1) immediate release; 2) sustained/controlled release; 3) delayed release; 4) buccal and sublingual; 5) chewable; 6) effervescent, and 7) orally disintegrating tablets.<sup>2</sup> A tablet product may belong to more than one of the specific types.

Tablets exhibit advantages in both manufacturing and administration. With respect to manufacturing, tablets generally exhibit better chemical, physical and microbiological stability, lower cost for manufacturing and distribution, as well as more accurate dose.<sup>2-4</sup> The ease of administration is another major benefit that leads to better patient compliance and therapeutic outcomes.<sup>2-4</sup>

In order to develop a successful tablet product, several quality standards have to be met to ensure efficacy and safety. Adequate mechanical strength is essential as a tablet cannot exist in its absence.<sup>5</sup> Parameters, such as particle size and shape, surface texture, plasticity, brittleness, viscoelasticity, and moisture content, can all influence the tableting behavior of APIs.

#### **1.2.1** Tablet compaction

Tablets are typically prepared with a tablet press, where the volume of a powder is reduced under a compaction force.<sup>20</sup> The compaction process, which takes place in a tooling set (a die and two punches), can be divided into four phases involving the specific mechanisms of 1) die filling and particle slippage, 2) compression, 3) decompression and 4) ejection.<sup>21, 22</sup> When the punches come in contact with the powder in the die, a low pressure is applied and particles undergo rearrangement through sliding and rotation. This results in a more densely pack powder bed in the die. Displacement of trapped air in bulk powder occurs, and the powder bed porosity decreases. At a critical porosity, the loose powder transitions into a rigid body,<sup>22</sup> marking the initiation of the compression stage. The continuously increasing compaction pressure and volume reduction of the powder causes deformation of the particles, including particle fracture, elastic deformation, and plastic deformation. Elastic deformation is temporary and reversible, but plastic deformation is permanent and irreversible.<sup>21</sup> The compaction pressure reaches the maximum at the end of the compression phase. The decompression phase, where the punches move away from each other, ensures the stress is reduced to zero. In the ejection phase, the compact is pushed out of the die. Tablets gain their mechanical strength from the compaction process, and the mechanical strength is governed by the total area of contact (bonding area) and the strength of particle interactions

(bonding strength).<sup>21</sup> This bonding area – bonding strength (BABS) interplay model will be further discussed in 1.2.3.

#### **1.2.2 Tablet mechanical strength**

The mechanical strength of tablets obtained is dependent upon a number of factors, e.g., physicochemical, particulate and mechanical properties of the bulk powder, as well as the conditions used for powder compression, such as compaction pressure and speed.<sup>23, 24</sup> Tablet mechanical strength can be quantified by a number of standard measurements, including breaking force <sup>25</sup>, tensile strength<sup>21</sup>, fracture toughness<sup>25</sup>, indentation hardness,<sup>26</sup> and tablet friability<sup>27</sup>.

Tablet tensile strength is one of the most commonly used parameter for quantifying mechanical strength.<sup>28</sup> Unlike breaking force, tablet size and geometry are taken into account in the determination of tensile strength. The tensile strength ( $\sigma$ ) of a cylindrical tablet can be obtained from the diametral compression test<sup>29</sup>, using *eqn. 1* 

$$\sigma = \frac{2F}{\pi Dh} \tag{1}$$

Where *F* is the breaking force, *D* is tablet diameter, and *h* is tablet thickness.

Adequate tablet tensile strength is required for successful tablet product development and manufacturing. On one hand, sufficient strength is needed for tablets to maintain their integrity during packaging, shipping and handling. Weak tablets may lead to problems, including unacceptable friability, reduced dose and a lack of pharmaceutical elegance (good quality in tablet size, shape and color). On the other hand, very strong tablets are difficult to break, as may be needed for patients with difficulty in swallowing. Furthermore, it can also result in slower rate of tablet disintegration and drug dissolution.<sup>21</sup>

#### **1.2.3** The bonding area – bonding strength (BABS) model

Tabletability is the ability of a material to form tablets with adequate mechanical strength by powder compaction, which is of critical importance to successful tablet product manufacturing.<sup>21</sup> Tabletability profiles can be obtained by plotting tablet tensile strength as a function compaction pressure. To explain the diverse tableting behaviors of pharmaceutical powders,<sup>15, 30-37</sup> a theoretical framework has been developed.

The BABS model is a qualitative model based on the concept of interparticulate boding area (BA) and boding strength (BS).<sup>21</sup> For molecular solids, van der Waals interactions as well as hydrogen bonding are primariy responsible for BS. During the consolidation process, the powder undergoes particle rearrangement, slippage and deformation, resulting in formation of BA. BA varies with mechanical and particle properties (e.g., size and shape). The various combinations of BA and BS lead to the diverse tabletability behaviors of pharmaceutical powders, i.e., tabletability is the result of the interplay between BA and BS. A clear understanding of the contributions of BA and BS to tensile strength can guide effective formulation and process development to overcome problems due to low tabletability.

# **1.3 Crystal Engineering**

Crystal engineering is the rational design of fundamental molecular solids.<sup>6</sup> It is of both fundamental and practical interest to researchers in multiple fields such as chemistry, material science, and pharmaceutical sciences.<sup>7</sup> Today, this subject is actively pursued by at least 150 independent research groups worldwide, covering topics such as 1) intermolecular interactions; 2) packing modes, in the context of these interactions and with the aim of defining a design strategy; and 3) crystal structure - properties relationship and their applications to improve properties through crystal structure modifications.<sup>7</sup>

In pharmaceutical sciences, the crystal engineering approach has progressed to become a successful designing strategy to obtain novel multicomponent complexes of the APIs exhibiting desired physicochemical properties without affecting their pharmacological performances.<sup>8</sup> Numerous examples in recent times have shown its application in attaining desirable mechanical and physicochemical properties, such as stability, solubility, and bioavailability.<sup>9-12</sup> In this line of research, it is of fundamental importance to solve the crystal structure of the API, which facilitates the understanding of the structure – property relationship. These information augments the Quality by Design (QbD) objective (a systematic approach to development that begins with predefined objectives and emphasizes product and process understanding and process control, based on sound science and quality risk management) of pharmaceuticals in solid state.<sup>38</sup>

## **1.3.1** The cocrystal - salt continuum and pK<sub>a</sub> rule

Both cocrystals and salts have been used to improve pharmaceutical properties of APIs.<sup>11, 39-41</sup> However, cocrystals are not regarded as new pharmaceutical APIs whereas salts are. This means that a complete set of characterization data is required for FDA approval of pharmaceutical products containing different salts.<sup>42</sup> This regulatory distinction makes it important to identify the ionization state of each new multicomponent complex intended for further pharmaceutical development.

The criterion that distinguishes a cocrystal drug from a salt is proton transfer. In multicomponent crystal systems, the position of the proton between the two components can be continuous based on their chemical nature, ranging from covalently bonded to the donor molecule to covalently bonded to the acceptor molecule.<sup>43, 44</sup> However, sometimes, it is difficult to apply this seemingly simple criterion to identify a cocrystal or salt, because of several phenomena: 1) partial proton transfer<sup>45</sup>; 2) position shift of a proton with temperature between the same pair of acid and base<sup>46</sup>; 3) the extent of proton transfer depends on the packing patterns of molecules in different polymorphs and hydrates<sup>47</sup>.

The empirical  $\Delta pK_a$  rule, which was developed on the basis of an analysis of 6465 crystalline complexes, has been used to predict the ionization state of crystalline complexes.<sup>48, 49</sup> A larger  $\Delta pK_a$  is correlated with a higher tendency of proton transfer. Generally, when the  $\Delta pK_a$  (=  $pK_a$  [base] –  $pK_a$  [acid]) is greater than 3, proton transfer is highly likely to occur.<sup>49</sup> When  $\Delta pK_a < 0$ , proton transfer is unlikely and

cocrystal would be expected. In the cases of  $\Delta pK_a$  between 0 and 3, either salt or cocrystal could form.<sup>43</sup> However, the U.S. FDA proposed  $\Delta pK_a = 1$  as the boundary between cocrystals and salts, i.e., when  $\Delta pK_a > 1$ , substantial proton transfer is likely to occur, resulting in salt formation; and when  $\Delta pK_a < 1$ , cocrystals are expected to be obtained.<sup>42</sup>

#### **1.3.2 Impact on drug properties and performance**

Each new multicomponent complex of any drug is expected to exhibit a distinct set of properties because of the unique structure.<sup>50</sup> Thus, forming multicomponent complex crystals is an effective strategy for improving manufacturability and clinical performance of drug products. Demonstrated advantages of this strategy include: 1) higher solubility, dissolution rate, and bioavailability;<sup>51-54</sup> 2) greater physical and chemical stability;<sup>55-57</sup> 3) better mechanical properties;<sup>11, 12, 58</sup> and 4) improved taste profiles.<sup>59-61</sup>

# **1.4 Structure – property relationship**

Understanding the relationship between crystal packing and the mechanical properties of the crystals is of interest to researchers in the fields of solid – state chemistry and crystal engineering.<sup>9, 10, 12, 13</sup> For example, such structure – property relationships help explain the different tableting behavior of the material.<sup>12, 14-16</sup> In addition, computational methods, such as energy framework calculation and topology analysis, can be used to further gain insight and possibly predict the compaction behavior of

different crystals.<sup>17,18</sup> Acetaminophen is a good example of the efforts in developing the structure – property relationship. The orthorhombic form II was initially observed to exhibit greater tabletability compared with the monoclinic form  $I.^{62}$ Later, an analysis of the crystal structures gave clues to their distinct tableting behavior: 1) pleated sheets along b – axis for form I, resulting in relatively stiff structure and poor compaction behavior, and 2) parallel hydrogen – bonded sheets along c – axis for form II, serving as slip planes to facilitate plastic deformation.<sup>63</sup> More recently, energy framework calculations showed that form II molecule has lower intermolecular interaction energy between neighboring sheets (-106 kJ/mol<sup>-1</sup>) compared with a form I molecule (-143 kJ/mol<sup>-1</sup>), indicating easier slide between sheets for form II than form I. This explains the greater plasticity and tabletability of form II.<sup>64</sup> CHAPTER 2.

# SWEET SULFAMETHAZINE ACESULFAMATE CRYSTALS WITH IMPROVED COMPACTION PROPERTY

# **Overview**

Sulfamethazine (SMT) is a sulfonamide antibacterial drug used to treat or prevent infections in both humans and animals. However, SMT exhibits unfavorable taste and poor compaction behavior. To overcome these problems, a 1:1 complex with an artificial sweetener, acesulfame (Acs), was prepared and characterized. The single crystal structure suggested that the new complex, SMT-Acs, is a salt. This was confirmed by an analysis of C-N bond length in comparison to those of multicomponent SMT crystals with known ionization states of SMT and Fourier transformation infrared spectroscopy. The applicability of the  $\Delta pK_a$  rule in multicomponent crystals of SMT was discussed. SMT-Acs exhibits better tabletability than SMT, which is attributed to its greater plasticity as shown by Heckel and Kuentz – Leuenberger analysis. The greater plasticity of SMT-Acs is consistent with the presence of slip planes identified by combined energy framework and topological analysis of the crystal structure.

# **2.1 Introduction**

Crystal engineering, "the rational design of functional molecular solids", is of interest in multiple research fields, including pharmaceutical sciences.<sup>1, 2</sup> Numerous examples have shown the ability of attaining desirable physicochemical and mechanical properties (i.e., stability, solubility and bioavailability) of active pharmaceutical ingredients (APIs).<sup>3-6</sup> Salt formation is an effective crystal engineering approach, where charge assisted intermolecular interactions play an important role in directing molecule packing in crystals. Hence, salt formation has been routinely applied to improve pharmaceutical properties of APIs.<sup>7-9</sup>

Sulfamethazine (SMT, Scheme 2.1a) is a sulfonamide antibacterial drug, which is used to treat or prevent infections in both humans and animals, with a broad spectrum of activity against gram-positive bacteria, gram-negative bacteria, and some protozoa.<sup>10, 11</sup> SMT can exist in two neutral tautomers, amidine<sup>11-23</sup> and imidine, <sup>11, 17, <sup>24</sup> cation (protonation of the -NH<sub>2</sub> group,  $pK_{a,1} = 2.65$ ),<sup>24-26</sup> and anion (deprotonation of the sulfonamide -NH- group,  $pK_{a,2} = 7.65$ ).<sup>16</sup> Correspondingly, various organic salts and pharmaceutical cocrystals of SMT were prepared and characterized with a focus on crystal structures and solid – state properties, except for compaction properties. Taste masking and compaction property enhancement of SMT are desired for developing high quality oral tablet products.<sup>27</sup></sup>

Acesulfame (Acs, Scheme 2.1b), a relatively strong acid ( $pK_a = 2.0$ ), is a calorie-free sugar substitute that is 200 times sweeter than sucrose.<sup>28</sup> Because of its safety profile,<sup>29</sup> Acs is a good candidate for masking the bitterness of SMT through forming a crystalline complex, as demonstrated in other drugs.<sup>30, 31</sup> The multiple

functional groups (amide and sulfonyl amide) of Acs allows interaction with SMT hydrogen donating groups of amine (-NH<sub>2</sub>) and sulfonamide (-NH-) and accepting groups of sulfoxy -O-, amine -N-, and pyrimidine -N-.<sup>11, 32</sup> In this work, we investigated the possibility of forming a sweet SMT-Acs complex to improve the poor mechanical properties and taste of SMT.

#### 2.2 Material and methods

#### **2.2.1 Materials**

Sulfamethazine (Sigma-Aldrich, St. Louis, MO), acesulfame potassium (Tokyo Chemical Industry co., Ltd., Japan), and acetonitrile (Sigma-Aldrich, St. Louis, MO) were used as received. Acs free acid was prepared from acesulfame potassium using a reported method.<sup>32</sup>

#### **2.2.2 Preparation of single crystals**

Equimolar SMT (556.7 mg, 2 mmoL) and Acs (326.3 mg, 2 mmoL) were suspended in 10 mL of acetonitrile. The suspension was stirred with a magnetic stirring bar for 24 h at room temperature. After 24 h, the suspension was filtered through filter paper with a Brinell funnel, and the filtrate was collected and passed through a 0.22 mm PTFE syringe filter membrane into a vial. The vial was left open in a fume hood to allow solvent evaporation. Crystals suitable for single crystal X-ray diffraction study were harvested within one week.

#### 2.2.3 Synthesis of SMT-Acs salt bulk powder

Bulk powders of SMT-Acs salt for compaction study were prepared by suspending SMT (5.57 g, 20 mmoL) and Acs (3.26 g, 20 mmoL) in 30 mL of acetonitrile and stirred for 24 h at room temperature. After 24 h, the solid phase was isolated by filtration using a Brinell funnel, and the dried at room temperature overnight.

#### **2.2.4 Single Crystal X-Ray Diffraction (SCXRD)**

SCXRD was conducted on a Bruker-AXS Venture Photon-II diffractometer (Bruker AXS Inc., Madison, Wisconsin), with a Photon-II (CMOS) detector. Data was collected at 100 K using Mo K $\alpha$  radiation (graphite monochromator). Various software packages from Bruker, including APEX3, SADABS, and SAINT, were used for data analysis. The structure was solved using SHELXT 2014 program and refined using SHELXL 2018 program.<sup>33, 34</sup> The space group P2<sub>1</sub>/n was determined based on systematic absences and intensity statistics. A direct-methods solution was calculated to provide most non-hydrogen atoms from the E-map. Full-matrix least-squares/difference Fourier cycles were performed to locate the remaining non-hydrogen atoms. All non-hydrogen atoms were refined with anisotropic displacement parameters. All hydrogen atoms were placed in ideal positions and refined as riding atoms with relative isotropic displacement parameters.

# 2.2.5 Powder X-ray Diffractometry (PXRD)

PXRD experiments were performed on a powder X-ray diffractometer (PANalytical

X'pert pro, Westborough, MA), using Cu K $\alpha$  radiation (1.54056 Å). Samples were scanned with a step size of 0.02 ° and 1 s/step dwell time from 5 to 35 ° two theta. The tube voltage and amperage were 45 kV and 40 mA, respectively.

#### **2.2.6 Thermal Analysis**

Hot stage microscopy (HSM) was carried out using a polarized light microscopy (Eclipse e200; Nikon, Tokyo, Japan), fitted with a hot stage. Images were captured by a DS-Fi1 microscope digital camera. Crystals were heated at a rate of 5 °C/min with a temperature controller (Linksys 32; V.2.2.0, Linkam Scientific Instruments, Ltd., Waterfield, UK).

The thermal properties collected on a differential scanning calorimeter (DSC) (Q1000, TA Instruments, New Castle, DE, USA). Powder samples (2~4 mg) were hermetically sealed in aluminum pans and heated with a heating rate of 10 °C/min under continuous dry nitrogen purge at a flow rate of 25 mL/min. Thermal gravimetric analysis (TGA) data were collected using a thermogravimetry analyzer (Model Q500, TA Instruments, New Castle, DE, USA). Powder samples (3~5mg) were loaded in open aluminum pans, and heated from room temperature up to 300 °C at 10 °C/min under a continuous dry nitrogen purge at 60 mL/min.

#### 2.2.7 Fourier Transformation Infrared Spectroscopy (FT-IR)

FT-IR spectra of all solid samples were obtained using a high resolution FT-IR spectrometer (VERTEX 70, Bruker Optics Inc., Billerica, MA. USA). A total of 32

scans for each sample were collected at 4 cm<sup>-1</sup> resolution, and an averaged spectrum ranged from 4000 - 600 cm<sup>-1</sup> was reported. The spectra were processed using OPUS software (v5.5, Bruker Optics Inc., Billerica, MA, USA).

#### 2.2.8 Dynamic Water Vapor Sorption Isotherm (DVS)

Water sorption isotherms were determined at 25  $^{\circ}$ C on an automated vapor sorption analyzer (DVS 1000, Surface Measurement Systems Ltd., Alperton, Middlesex, UK), with continuous 50 mL/min nitrogen flow. The RH values were increased from 0% to 95% using a step size of 5%. At each step, the sample was equilibrated until one of the two criteria was met, dm/dt  $\leq 0.003\%$  or 6 h equilibration period.

#### **2.2.9** Tabletability

Powder compaction studies were conducted on a material testing machine (model 1485; Zwick/ Roell, Ulm, Germany) at 4 mm/min speed. To minimize the effect of particle size, powder samples were ground in a mortar with a pestle and passed through a #60 mesh ( $\leq 250 \mu$ m) sieve before compaction. Comparable particle sizes of SMT and SMT-Acs were confirmed by polarized light microscopy. Compaction was carried out over the pressure range of 20 - 320 MPa using a die and flat-faced round punches (8 mm diameter). A suspension of magnesium stearate in ethanol (5% w/v) was applied onto the punch tips and die wall using a brush, and then dried by a fan prior to filling the die with powder. Tablets were left in sealed glass vials at room temperature for at least 24 h to allow relaxation. The diametrical breaking

force was determined on a texture analyzer (TA-XT2i; Texture Technologies Corporation, Scarsdale, New York). Tablet tensile strength ( $\sigma$ ) was calculated using eqn. 1.<sup>35</sup>

$$\sigma = \frac{2F}{\pi Dh} \tag{1}$$

Where F is the breaking force, D is tablet diameter, and h is tablet thickness.

True density of powder samples was measured by helium pycnometry (Quantachrome Instruments, ultrapycnometer 1000e, Boynton Beach, Florida). Samples  $(1\sim2g)$  were accurately weighed and placed into a sample cell. Measurements were repeated to a maximum of 100 or a <0.005% coefficient of variation of five consecutive measurements was obtained. The mean of the last five measurements was reported as the true density of the sample.

#### 2.2.10 In-Die Heckel Analysis

Three tablets (about 200 mg) were prepared at peak compaction pressures of 280, 300 and 320 MPa at 4 mm/min speed using a material testing machine (model 1485; Zwick/ Roell, Ulm, Germany). The force and nominal strain were exported for calculating pressure and porosity ( $\varepsilon$ ), which were analyzed using the Heckel equation,<sup>36</sup> eqn. 2

$$-\ln\left(\varepsilon\right) = KP + A \tag{2}$$

Where *P* is the compaction pressure applied, the slope *K* and y-axis intercept *A* are calculated using the linear portion of the Heckel plot. The mean yield pressure  $(P_y)$ , which is the reciprocal of *K*, is a plasticity parameter of the powder. A lower  $P_y$ 

value suggests higher powder plasticity.

#### **2.2.11 Out-of-Die Kuentz–Leuenberger (KL) analysis**

The out-of-die pressure – porosity data were analyzed using the Kuentz–Leuenberger (KL) method (Eqn. 3).<sup>37</sup>

$$P = \frac{1}{c} \left[ \varepsilon - \varepsilon_c - \varepsilon_c \ln \left( \frac{\varepsilon}{\varepsilon_c} \right) \right]$$
(3)

Here, the parameter 1/C is another parameter for quantifying material plasticity. Higher plasticity is indicated by a lower 1/C value. The KL analysis is superior to Heckel analysis in analyzing compressibility data, because the former uses the entire set of data whereas the latter only uses data that appears linear over a narrower range of compaction pressure.<sup>38</sup> *P* is the compaction pressure applied,  $\varepsilon$  is the porosity of the powder, and the critical powder porosity,  $\varepsilon_c$ , is a parameter that describes the porosity at which the powder starts to gain rigidity or strength.<sup>37</sup>

## 2.3 Results and discussion

#### **2.3.1** Crystal structure

SMT-Acs complex (CCDC: 1821691) crystallizes in the monoclinic P2<sub>1</sub>/n space group with four asymmetric units in unit cell (Z = 4, Figure 2.2a). Each asymmetric unit contains one SMT<sup>+</sup> cation and one Acs<sup>-</sup> anion (Z' = 1). Key crystallographic parameters are summarized in Table 2.1, and hydrogen bonds are shown in Table 2.2. Proton transfer occurs between SMT N<sub>2</sub> site and Acs sulfonyl amide -NH- group. SMT<sup>+</sup> cation and Acs<sup>-</sup> anion are connected through two hydrogen bonds, namely,  $N_1$ -H<sup>...</sup> $N_{.3}$  and  $N_2^+$ -H<sup>...</sup> $O_1 = C_2$  (Figure 2.2b). Each SMT<sup>+</sup> cation is connected to four more SMT<sup>+</sup> through N-H<sup>...</sup>O = S (2.995 Å) and one Acs<sup>-</sup> through the above mentioned  $N_1$ -H<sup>...</sup> $N_{.3}$  (2.796 Å) and  $N_2^+$ -H<sup>...</sup> $O_1 = C_2$ , (2.627 Å) hydrogen bonds, forming a  $R_2^2$ (8) heterosynthon (Figure 2.2b). The Acs<sup>-</sup> also connects to another SMT<sup>+</sup> through N-H<sup>...</sup>O = C (3.015 Å). SMT<sup>+</sup> molecules, linked by N-H...O = S with bond length of 2.995 Å, form a zigzag chain along b-axis (Figure 2.2c). The three-dimensional packing pattern of SMT-Acs is shown in Figure 2.2d.

#### **2.3.2 Ionization state analysis**

#### 2.3.2.1 $\Delta pK_a$ analysis

An empirical  $\Delta pK_a$  rule has been proposed to predict the ionization state of crystalline complexes, which have been tested in 6465 crystalline complexes.<sup>39, 40</sup> Generally, when the  $\Delta pK_a$  (=  $pK_a$  [base] –  $pK_a$  [acid]) is greater than 3, proton transfer is highly likely to occur since a larger  $\Delta pK_a$  suggests higher tendency of proton transfer.<sup>40</sup> When  $\Delta pK_a < 0$ , proton transfer is unlikely and a cocrystal would be expected. In the cases of  $\Delta pK_a$  between 0 and 3, either salt or cocrystal could form.<sup>41</sup> On the other hand, the U.S. FDA proposed  $\Delta pK_a = 1$  as the boundary between cocrystals and salts, i.e., when  $\Delta pK_a > 1$ , substantial proton transfer is likely to occur, resulting in salt formation; and when  $\Delta pK_a < 1$ , cocrystals are expected to be obtained.<sup>42</sup>

A search of CSD returned a total of 32 multicomponent crystals of SMT, including 26 cocrystals and 6 salts.<sup>43</sup> Out of the 26 cocrystals, SMT can be present in the form of

amidine (19) or imidine (6), with one crystal contains both amidine and imidine forms. Out of the 6 salts, 4 are SMT cations and 2 are anions. However, the application of the  $\Delta p K_a$  rule in SMT multicomponent crystals requires caution, since SMT can exist in different molecular forms (amidine, imidine, cation, and anion) with different  $pK_{as}$ . The correct calculation of  $\Delta p K_a$  should consider functional groups that are directly involved in the interacting hydrogen/proton donating and accepting groups. The simple use of SMT amine -NH<sub>2</sub> group  $pK_a$  value would have led to an erroneous  $\Delta pK_a$ , since the hydrogen/proton donating and accepting groups in many known SMT multicomponent structures do not directly involve that group. By taking account of the actual interacting groups, the distribution map of ionization state versus  $\Delta p K_a$  is obtained (Figure 2.3, Table 2.3). Salts could form at  $\Delta p K_a$  well below 0, and cocrystals could effectively form at  $\Delta p K_a$  about 3. A similar result was observed in fumaric acid multicomponent crystals recently, in which cocrystals distributed around 53% in  $\Delta p K_a > 0$  region and 47% in  $\Delta p K_a < 0$  region.<sup>44</sup> Thus, the  $\Delta p K_a$  rule cannot be reliably used to predict ionization state in SMT multicomponent crystals. Therefore, bond length analysis and infrared spectroscopy were applied to establish the ionization state of SMT-Acs.

#### **2.3.2.2 Bond length analysis**

Bond lengths often give clues to the complex's ionization state. Bond length diagrams containing different forms of SMT and Acs in known crystal structures are shown in Figure 2.4. For SMT-Acs, the SMT  $C_1$ - $N_1$  and  $C_1$ - $N_2$  bond lengths (1.366)

and 1.354 Å, respectively, Figure 2.4a). Thus, SMT-Acs falls into the SMT cation region, whereas Acs fall into the anionic group based on  $C_2$ - $N_3$  and  $C_2$ - $O_1$  bond lengths (1.352 and 1.256 Å, respectively, Figure 2.4b).<sup>32, 45</sup> Thus, bond length analysis indicates salt formation for SMT-Acs.

#### 2.3.2.3 Fourier Transformation Infrared Spectroscopy

Characteristic FTIR peaks of SMT and Acs-K can be observed in the spectrum of SMT-Acs (Figure 2.5). For example, the C=O peak (1652 cm<sup>-1</sup> in Acs-K) shows up at 1647 cm<sup>-1</sup> in SMT-Acs, and the N–H stretching frequency of SMT is shifted from 3440, 3339 cm<sup>-1</sup> to 3490 and 3389 cm<sup>-1</sup> in SMT-Acs. Additionally, a 2636 cm<sup>-1</sup> broad peak is observed in SMT-Acs, which corresponds to the NH<sup>+</sup> functional group stretching (between 2700 and 2400 cm<sup>-1</sup>), confirming that both SMT and Acs are in ionic state.<sup>32</sup> Considering these data, complete proton transfer from Acs to SMT is confirmed for SMT-Acs.

## 2.3.3 Solid-State Characterization

#### **2.3.3.1** Phase Purity of the Bulk Powders

Powder X-ray diffractogram (PXRD) of the bulk SMT-Acs powder matched the PXRD calculated from the single crystal structure (Figure 2.6). The experimental PXRD pattern (room temperature) was shifted to slightly smaller  $2\theta$  angles compared to the calculated pattern using structure solved at 100 K due to thermal expansion

effect.<sup>46, 47</sup> The good PXRD pattern match confirms the phase purity of the bulk powder.

#### 2.3.3.2 Thermal Analysis

Both SMT and Acs exhibited single endotherms in their DSC traces (Figure 2.7a), with onset temperatures of 197 °C ( $\Delta H_f = 144.7 \text{ J/g}$ ) and 116 °C ( $\Delta H_f = 130.8 \text{ J/g}$ ), respectively. Acs is volatile as significant weight loss occurred after melting (Figure 2.7c). SMT did not undergo detectable weight loss until about 20 °C above melting temperature (Figure 2.7c). An exothermal peak 198 °C was observed in SMT-Acs after the endothermal peak. These thermal events corresponded to bubbling at 198 °C without any sign of melting (Figure 2.7b), indicating dissociation of SMT and Ace and subsequent evaporation of Ace gave rise to the weight loss near 200 °C (Figure 2.7c). Importantly, SMT-Acs had negligible weight loss below 150 °C, suggesting thermal stability acceptable for tablet development.<sup>48</sup>

#### 2.3.3.3 Moisture Sorption Behavior

Low hygroscopicity is preferred as the moisture content could alter the flowability, tabletability, and cause stability issue.<sup>49-51</sup> The Acs free acid takes up nearly 30% of water when RH increased from 70% to 95% (Figure 2.8). Thus, Acs is very hygroscopic according to the classification in the European Pharmacopeia.<sup>52</sup> SMT is non-hygroscopic, since it only takes up negligible amount of water, 0.6%, at 95% RH. Moisture uptake by SMT-Acs, 0.2% at 95% RH, is the lowest, which also supports its

use in tablet manufacturing.

#### 2.3.3.4 Tabletability

A tensile strength of 2 MPa is considered sufficient for maintaining tablet integrity during normal handling of most materials.<sup>53</sup> The tabletability of SMT-Acs was significantly better than SMT (Figure 2.9a), where tensile strength of SMT-Acs tablets was approximately two times that of SMT tablet at a given pressure despite similar particle sizes (Figure 2.10). Importantly, SMT-Acs tablet reached 2 MPa tensile strength at about 150 MPa, whereas SMT could not reach 2 MPa over the entire range of compaction pressure. Thus, the improved tabletability of SMT-Acs also supports its use in tablet manufacturing.

The higher tabletability can be attributed to either larger BA, higher BS, or both.<sup>54, 55</sup> To better understand the underlying mechanism of tabletability improvement, the plasticity of SMT and SMT-Acs was assessed, because more plastic materials can deform more easily to result in larger bonding area (BA). SMT-Acs is more plastic than SMT, since both the  $P_y$  and 1/C values of SMT-Acs were significantly lower than those of SMT (Table 2.4). The greater plasticity explains the steeper decrease in tablet porosity of SMT-Acs with increasing pressure (Figure 2.9b). It is interesting to note that tablet porosity of SMT-Acs was higher than that of SMT tablets at pressure < 300 MPa, despite the similar particle sizes (Figure 2.10). Thus, the bonding area may not account for the higher tabletability of SMT-Acs despite its higher plasticity. However, the tensile strength extrapolated to zero porosity, which is a parameter for assessing the apparent bonding strength (BS), was comparable between SMT-Acs and SMT (Figure 2.9c). Thus, BS cannot explain the superior tabletability of SMT-Acs. Consequently, the larger BA is responsible for the better tabletability of SMT-Acs despite the larger tablet porosity below 300 MPa compaction pressures (Figure 2.9b). This is possible only when the plasticity of SMT-Acs is significantly higher than SMT, which is supported by its significantly lower  $P_y$  and 1/C (Table 2.4).<sup>55</sup>

# 2.4 Structure-properties relationship

The origin of the significantly greater plasticity of SMT-Acs than SMT can be understood by quantitatively examining their crystal structures using a computational tool kit.<sup>56</sup> The most likely two-dimensional layers in SMT run along the *ab* plane (Figure 2.11a). Neighboring molecules in this layer interact through N-H<sup>...</sup>O=S (2.947 Å) and N-H<sup>...</sup>N (3.092 Å) hydrogen bonds as well as strong  $\pi$ - $\pi$  interaction (3.393 Å) (Figure 2.11a). Adjacent layers are fortified through weaker hydrogen bonds of C-H<sup>...</sup>O=S (3.331 Å) (Figure 2.11a). The energy framework of SMT (Figure 11b) suggested that (0 0 1) planes exhibit higher intralayer bonding energy (-261.1 kJ/mol) than the interlayer bonding energy (-80.6 kJ/mol), which meets the anisotropic energy requirement for slip planes. However, the molecular layers sliding along the (0 0 1) is hindered by the interdigitation between layers (negative interlayer distance of -1.80 Å) (Figure 2.11a).

In comparison, the SMT-Acs exhibits a columnar structure running along the a axis

(Figure 2.11c). Within the columns, molecules are linked by N-H<sup> $\cdots$ </sup>N (2.796 Å), N-H<sup> $\cdots$ </sup>O (2.627 Å), and N-H<sup> $\cdots$ </sup>O=S (3.011 Å) hydrogen bonds. Each column interacts with four neighboring columns via N-H<sup> $\cdots$ </sup>O=S (2.995 Å) hydrogen bonds (Figure 2.11c). The intracolumn bonding energy (-256.7 kJ/mol) is much higher than the intercolumn bonding energy (-92.8 kJ/mol) (Figure 2.11d), suggesting energetically favored molecular movement through slippage of columns, when the crystal is stressed.<sup>6</sup> Both (0 1 2) and (0 0 1) are possible slip planes in SMT-Acs based on energy framework alone. However, the (0 1 2) plane is likely the primary slip plane because of the lowest degree of interdigitation between adjacent layers (-0.22 Å) based on the topology analysis. Therefore, the crystal structure of SMT-Acs supports its higher plasticity than SMT established by bulk powder compaction data (Table 2.4).<sup>54</sup>

## **2.5 Conclusions**

A sweet 1:1 salt between sulfamethazine and acesulfame was obtained. Characterization of solid-state and compaction properties suggests suitability of SMT-Acs for tablet formulation and manufacturing. The better tabletability of SMT-Acs was attributed to its higher plasticity, which in turn is explained by the favorable crystal structure, featuring facile slip planes.

Name	SMT-Acs
Formula	$C_{16}H_{19}N_5O_6S_2$
Formula weight	441.48
Temperature / K	100 (2)
Crystal system	Monoclinic
Space group	$P2_1/n$
<i>a</i> , Å	8.3209 (2)
<i>b</i> , Å	10.8674 (3)
<i>c</i> , Å	21.7899 (5)
a, deg	90
b, deg	99.9810 (10)
g, deg	90
Volume, Å <sup>3</sup>	1940.56 (8)
Ζ	4
$Dc / g cm^{-3}$	1.511
F(000)	920
GOF	1.085
<i>R</i> 1 [ <i>I</i> >2sigma( <i>I</i> )]	0.0371
<i>wR</i> 2 [ <i>I</i> >2sigma( <i>I</i> )]	0.0909

 Table 2.1 Crystallographic data for the SMT-Acs crystal.

**Table 2.2** Hydrogen bonds in SMT-Acs crystal.

D-HA	<b>D-H</b> (Å)	<b>HA</b> (Å)	<b>DA</b> (Å)	D-HA (deg)	Symmetry codes
C1-H1CO1	0.980	2.643	3.593	163.23	-x+1, -y+2, -z+1
N3-H3AN1	0.881	1.916	2.796	175.99	
N2-H2BO5	0.884	2.334	2.995	131.56	-x+3/2, y-1/2, -z+3/2
N2-H2AO6	0.833	2.204	3.011	163.05	x+1, y, z
N5-H5O4	0.947	1.683	2.627	174.10	
С9-Н9О1	0.956	2.609	3.562	174.88	
C10-H10O6	0.944	2.588	3.344	137.34	x+1, y, z
C13-H13O2	0.951	2.579	3.246	127.52	x, y-1, z
C15'-H15DO6	0.980	2.640	3.420	136.72	-x+1/2, y-1/2, -z+3/2
C15'-H15EO2	0.980	2.466	3.353	150.36	x, y-1, z

Table 2.3 Realistic interaction sites and corresponding  $\Delta p K_a$  in multicomponent crystals of SMT

coformer	coformer SMT SMT interaction		Coformer	$\Delta pK_a$
	conformation site & pK <sub>a</sub>		pKa	
2,4-dihydroxybenzoic	amidine	pyrimidine -N-	3.11	-4.08
acid		(base), -0.97		
3,4-dichlorobenzoic acid	amidine	pyrimidine -N-	3.6	-4.57
		(base), -0.97		
sorbic acid	amidine	pyrimidine -N-	4.76	-5.73
		(base), -0.97		
1-hydroxy-2-naphthoic	amidine	pyrimidine -N-	10.94	-11.91
acid		(base), -0.97		
benzene-1,2,4,5-tetracar	amidine	pyrimidine -N-	1.87	-2.84
boxylic acid		(base), -0.97		
2-hydroxybenzoic acid	amidine	pyrimidine -N-	2.97	-3.94
		(base), -0.97		
2-nitrobenzoic acid	amidine	pyrimidine -N-	2.16	-3.13
		(base), -0.97		
4-nitrobenzoic acid	amidine	pyrimidine -N-	3.41	-4.38
		(base), -0.97		
2-aminobenzoic acid	amidine	pyrimidine -N-	2.14	-3.11
		(base), -0.97		
4-aminobenzoic acid	amidine	pyrimidine -N-	2.38	-3.35
		(base), -0.97		
1,4-diazabicyclo[2.2.2]o	amidine	sulfonamide	8.7	1.71
ctane		-NH- (acid),		
		6.99		
acetylsalicylic acid	amidine	pyrimidine -N-	3.49	-4.46
		(base), -0.97		
4-aminosalicylic acid	amidine	pyrimidine -N-	3.25	-4.22
		(base), -0.97		
indole-2-carboxylic acid	amidine	pyrimidine -N-	4.44	-5.41
		(base), -0.97		
2,4-dinitrobenzoic acid	amidine	pyrimidine -N-	1.43	-2.4
		(base), -0.97		
4-chlorobenzoic acid	amidine	pyrimidine -N-	3.98	-4.95
		(base), -0.97		
4,4'-ethane-1,2-diyldipyr	amidine	pyrimidine -N-	6.13	7.1
idine		(base), -0.97		
niflumic acid	amidine	pyrimidine -N-	1.88	-2.85
		(base), -0.97		

flufenamic acid	flufenamic acid amidine pyrimidine -N-		3.88	-4.85
A hydroxybonzoia acid imidina		sulfonamide -N-	1 51	2 47
	minume	(base) 7.01	4.54	2.47
hanzamida	imidina	(Uase), 7.01	2	0.01
benzannde	imaine	sufformating -IN-	-2	-9.01
		(base), 7.01	1.00	0.00
picolinamide	imidine	sulfonamide -N-	-1.08	-8.09
		(base), 7.01		
4-hydroxybenzamide	imidine	sulfonamide -N-	-3.01	-10.02
		(base), 7.01		
3-hydroxy-2-naphthoic	imidine	sulfonamide -N-	2.79	4.22
acid		(base), 7.01		
saccharine	imidine	sulfonamide -N-	1.31	5.7
		(base), 7.01		
theophylline	Amidine &	sulfonamide	7.82 (acid)	-0.81 &
	imidine	-NH- (acid),	& -0.78	-7.77
		6.99 &	(base)	
		sulfonamide -N-		
		(base), 7.01		
4-aminopyridine	anion	sulfonamide	8.95	1.96
1,5		-NH- (acid).		
		6.99		
piperazine	anion	sulfonamide	9.56	2.57
r-r		-NH- (acid)		
		6 99		
saccharin	cation	pyrimidine -N-	1 31	-2.28
saccharm	cation	(base) = 0.97	1.51	2.20
5 nitro soliculio acid		nyrimidine N	2.2	3 17
J-IIIIO Sancyne aciu	Cation	(base) 0.07	2.2	-3.17
		(base), -0.97	0.42	1.20
picric acid	cation	pyrimidine -N-	0.42	-1.39
		(base), -0.97	1.50	2.5
3,5-dinitrosalicylic cation		pyrimidine -N-	1.53	-2.5
		(base), -0.97	-	
acesulfame	cation	pyrimidine -N-	2	-2.97
		(base), -0.97		

**Table 2.4** Plasticity parameters of SMT and SMT-Acs from in-die Heckel analysisand out-of-die KL analysis.

	SMT	SMT-Acs
$P_{y}$ (MPa)	77.1 $\pm$ 1.2 (R <sup>2</sup> = 0.99)	57.2 $\pm$ 3.3 (R <sup>2</sup> = 0.98)
<i>1/C</i> (MPa)	$3345 \pm 422$ (R <sup>2</sup> = 0.98)	$818 \pm 101$ (R <sup>2</sup> = 0.98)



Figure 2.1 Molecular structure of (a) sulfamethazine (b) acesulfame.



**Figure 2.2** Crystal structure of SMT-Acs: (a) unit cell viewed along *a*-axis, (b) asymmetric unit, (c)  $SMT^+$  cations formed a chain along the *b*-axis, (d) three-dimensional packing diagram.



Figure 2.3 Ionization state distribution of SMT multicomponent crystals versus  $\Delta p K_a$  between interacting groups. Saccharin (blue line) forms both cocrystal and salt with SMT, while theophylline (orange line) forms cocrystal with amidine and imidine SMT simultaneously.



**Figure 2.4** Bond length distributions of (a)  $C_1$ - $N_2$  vs.  $C_1$ - $N_1$  for SMT in crystal structures containing SMT, (b)  $C_2$ - $N_3$  vs.  $C_2$ - $O_1$  for Acs in known crystal structures containing Acs



**Figure 2.5** IR spectra for SMT, Acs-K, and SMT-Acs. The 2636  $\text{cm}^{-1}$  broad peak corresponds to NH<sup>+</sup> stretching.



Figure 2.6 Calculated and experimental PXRD patterns of SMT-Acs



Figure 2.7 Thermal analysis of SMT, Acs, and SMT-Acs: (a) DSC, (b) HSM, and (c)

TGA



Figure 2.8 Moisture sorption curves of SMT, Acs and SMT-Acs.



**Figure 2.9** (a) Tabletability (b) compressibility, and (c) compactability profiles of SMT and SMT-Acs, where the shaded bands indicate 90% confidence interval.



Figure 2.10 Particle size of (a) SMT and (b) SMT-Acs.



**Figure 2.11** Crystal packing patterns (left) and corresponding energy framework (right) for SMT (a, b) and SMT-Acs salt (c, d) viewed along c axis. The thickness of each cylinder (in blue) represents the relative strength of interaction.

# CHAPTER 3.

# **RESEARCH SUMMARY AND FUTURE WORK**

# **Research Summary**

A sweet 1:1 salt between sulfamethazine and acesulfame, SMT-Acs, was successfully prepared and characterized. The solid-state and compaction properties of SMT-Acs suggests its suitability for tablet formulation and manufacturing. The better tabletability of SMT-Acs was attributed to its higher plasticity, which in turn is explained by the presence of facile slip planes in the crystal structure that favors plastic deformation.

### **Future Work**

The thermal behavior of SMT-Acs captured by DSC and HSM are consistent. We also demonstrated the thermal stability of SMT-Acs from the TGA data. However, the nature of the thermal events between 190 and 200 °C remains elusive. Thus, further investigations are needed to fully understand the thermal properties of SMT-Acs. DSC experiments at different heating rates and variable temperature XRD may be of help to elucidate these observations. In addition, additional confirmation of the sweetness of SMT-Acs by either a taste panel or E – tongue test is needed.

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