

Article

Exploiting Supramolecular Synthons in Cocrystals of Two Racetams with 4-Hydroxybenzoic Acid and 4-Hydroxybenzamide Coformers

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Abstract: Structures of three cocrystals of nootropic racetams were studied. They included two cocrystals of phenylpiracetam (PPA) with 4-hydroxybenzoic acid (HBA) with different stoichiometries, PPA·HBA and PPA·2HBA, and cocrystal of 2-(4-phenyl-2-oxopyrrolidin-1-yl)-N'-isopropylideneacetohydrazide (PPAH) with 4-hydroxybenzamide (HBD), PPAH·HBD·(acetone solvate). X-ray study of the pure forms of PPA and PPAH was also carried out to identify variations of molecular synthons under the influence of conformers. The cocrystal structures revealed the diversity of supramolecular synthons namely, amide-amide, amide-acid, acid-acid, and hydroxyl-hydroxyl; however, very similar molecular chains were found in PPA and PPA·2HBA, and similar molecular dimers in PPAH and PPAH·HBD. In addition, conformational molecular diversity was observed as disorder in PPA·2HBA as it was observed earlier for *rac*-PPA that allows for the consideration that cocrystal as an example of partial solid solution. Quantum chemical calculations of PPA and PPAH conformers demonstrated that for most conformers, energy differences do not exceed 2 kcal/mol that suggests the influence of packing conditions (in this case *R*- and *S*-enantiomers intend to occupy the same molecular position in crystal) on molecular conformation.

Keywords: racetams; phenylpiracetam; X-ray; supramolecular synthon; hydrogen bond; conformation



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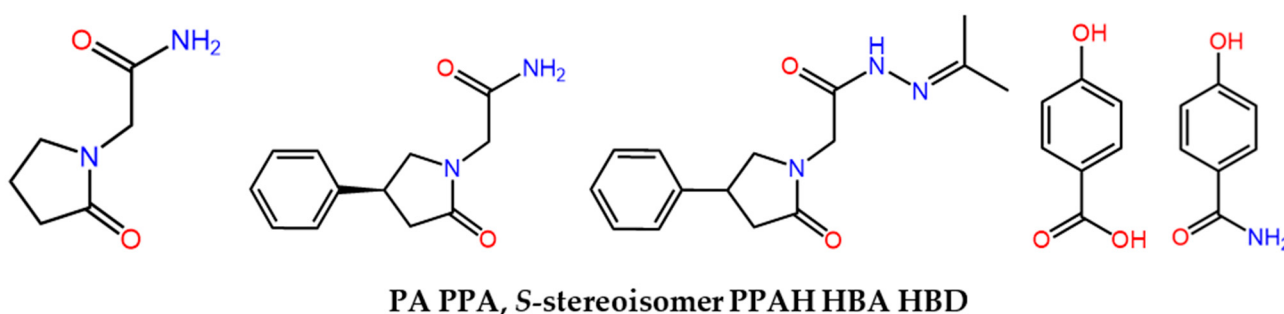
1. Introduction

Currently there is a significant interest in pharmaceutical materials that can be used for treatment of central nervous system (CNS) disorders. One of the groups of such materials is nootropics that are prescribed as drugs or supplements to improve memory, focus, and cognitive performance. Racetams (piracetam and its derivatives) are nootropic agents known since the 1970s [1–4]. Depending on their substituents, they have different degrees of pharmacological activity.

Nootropil (2-oxo-1-pyrrolidinyl-acetamide, piracetam, PA, Scheme 1) was marketed by the United Collection Bureau (UCB) to treat memory and balance problems [5]. The physiological action and structural peculiarities of PA have been studied thoroughly. This compound of one hydrate and five anhydrous crystalline forms was documented by Fabbiani et al. [6].

In search for nootropics with improved physicochemical properties, significant efforts were addressed to the PA cocrystals [7–11]. Zaworotko et al. [11] demonstrated the importance of supramolecular homo- and heterosynthons in crystal engineering of active pharmaceutical ingredients (APIs) for an explanation and prediction of polymorphic and cocrystalline forms of APIs. This paradigm looks significantly important for nonionized

drugs like racetams since it depicts the relative ease with which pharmaceutical cocrystals can be prepared and diversified from the view point of their composition and physical properties. Several examples revealed the successful exploiting of carboxylic acid—primary amide supramolecular heterosynthon in crystal engineering of cocrystals including those that contain two APIs [12–15]. Among published examples of **PA** cocrystals containing two APIs, to mention a few, are carbamazepine with **PA** [8], 2,6-diaminopyridine with **PA** [9], myricetin with **PA** [16]. Viertelhaus et al. obtained six **PA** cocrystals with different organic acids [10]. Cocrystal screening through structural resemblance for eleven **PA** cocrystals with ten different acids was reported by Leyssens et al. [17]. Those examples demonstrated that the amide-carboxylic acid heterosynthon was rarely found in the crystal structures in the presence of additional hydrogen bond donors and acceptors. An example of significant change of properties by cocrystallization is **PA**—*L*-tartaric acid cocrystal. It was found that a reversible water adsorption with a maximum water uptake of 3% was observed for this cocrystal, while for pure **PA** it was more than 40% [10]. Such a drop of hygroscopicity is important for the pharmaceutical industry. An interesting peculiarity of **PA**-hydroquinone cocrystal [8] was that both coformers produced separate parallel 1D motifs of H-bonded **PA** and hydroquinone molecules. In paper [18] multi-component crystallization conditions of several APIs including **PA** allowed for obtaining new polymorph modifications and solvates of APIs.



Scheme 1. Structural formulas of piracetam (**PA**), *S*-phenylpiracetam (**S-PPA**), 2-(4-phenyl-2-oxopyrrolidin-1-yl)-*N'*-isopropylideneacetohydrazide (**PPAH**), and coformers 4-hydroxybenzoic acid (**HBA**) and 4-hydroxybenzamide (**HBD**).

Another member of racetams family, 2-(2-oxo-4-phenyl-1-pyrrolidinyl)acetamide (phenotropil, carphedon, phenylpiracetam, **PPA**, Scheme 1) demonstrates, apart from nootropic properties, strong antidepressant and anxiolytic activity [19]. The molecule of **PPA** contains one chiral center in the pyrrolidine ring. So, **PPA** exists in racemic and two enantiomeric, *R*- and *S*-forms (Scheme 1) [19,20]. Crystallographic data (CSD version 5.43, last update November 2022) are available for *rac*-**PPA** and polymorphs of *R*- and *S*-enantiomers [20,21]. *Rac*-**PPA** (refcodes QELNEB01, QELNEB) crystallizes in the triclinic *P*-1 space group. Stereoisomers crystallize as orthorhombic (refcode SAMZOY, *P*₂₁₂₁₂₁ space group) and triclinic (refcode SANBAN, *P*1 space group) polymorphs [19,20]. It should be mentioned that pharmacological activities of *rac*-**PPA** and its *R*- and *S*-enantiomers are different, and the memory-improving activity is only characteristic of *R*-enantiomer [22].

To mask their bitter taste, some racetams including **PA** and **PPA** were cocrystallized with saccharine and zinc saccharinate [7]. Particularly for **PPA**, these attempts resulted in preparation of two crystalline complexes, ZnSac₂Car (Car = carphedon in [7], refcode NEFLOB) as a 1D coordination chain where both carbonyl oxygens of pyrrolidone and amide groups coordinate to two different zinc atoms, and [ZnSac₂(H₂O)₂] $\cdot$ Car $\cdot$ EtOH $\cdot$ H₂O (refcode NEFMOE) as a discrete inclusion compound with carphedon interacting with the [ZnSac₂(H₂O)₂] complex through hydrogen bonding [7]. Apart from the **PPA** pure forms, these are the only two known examples of structurally studied multicomponent **PPA** crystals.

For 2-(4-phenyl-2-oxopyrrolidin-1-yl)-N'-isopropylideneacetohydrazide (**PPAH**, Scheme 1), a relatively new member of racetams family [19], to the best of our knowledge, no crystallographic information is available either for pure **PPAH**, or for its cocrystals.

The goal of this study was a search for new forms of nootropics **PPA** and **PPAH**, and a comparison of molecular synthons found in pure materials, different cocrystalline, and polymorphic forms. As coformers for **PPA** and **PPAH**, we used compounds with molecular structures similar to those which were successfully used for formation of **PA** cocrystals [10,11], such as 4-hydroxybenzoic acid (**HBA**) and 4-hydroxybenzamide (**HBD**). The **HBA** in particular has performed well in cocrystal design with APIs [11,23–26]. Structural, and computational studies were carried out for the following pure and cocrystalline materials: **PPA**, **PPA·HBA**, **PPA·2HBA**, **PPAH**, and **PPAH·HBD**·(acetone solvate).

2. Materials and Methods

2.1. Chemicals and Solvents

Phenylpiracetam powder (100%) was purchased from Purenootropics (Purenootropics.net), and Phenylpiracetam as a supplement in capsules was purchased from BrainMedics. 4-Hydroxybenzoic acid, 4-hydroxybenzamide, and acetone solvent were purchased from Sigma Aldrich and used without purification.

2.2. Synthesis

2.2.1. Synthesis of **PPA·HBA** (1:1) Cocrystal

An amount of 22.0 mg (0.1 mmol) of **PPA** and 14.0 mg (0.1 mmol) of **HBA** were dissolved in 5 mL of acetone at room temperature in a test tube covered with parafilm with a hole. After a week, the solvent had evaporated and crystals were observed.

2.2.2. Synthesis of **PPA·2HBA** (1:2) Cocrystal

An amount of 44.0 mg (0.2 mmol) of **PPA** and 42.0 mg (0.3 mmol) of **HBA** were dissolved in 5 mL of acetone at room temperature in a test tube covered with parafilm with a hole. After a week the solvent had evaporated and crystals were observed.

2.2.3. Synthesis of **PPAH·HBD** Cocrystal

Phenylpiracetam produced by BrainMedics was removed from capsules and used for cocrystallization. After cocrystal was obtained and characterized with single crystal X-ray diffraction analysis, we realized that instead of **PPA** we had its derivative, 2-(4-phenyl-2-oxopyrrolidin-1-yl)-N'-isopropylideneacetohydrazide (**PPAH**) presented in Scheme 1. To be sure that powder from capsules was indeed **PPAH**, we recrystallized it from acetone and the single crystal X-ray study unambiguously confirmed that the substance was **PPAH** not **PPA**. In addition, HPLC analysis was performed to check the purity of the substance from capsules. It demonstrated only one peak that stated the compound from capsules did not have any contaminations.

2.2.4. Synthesis of **PPAH·HBD** Cocrystal

To obtain the cocrystal, **PPAH** (27.0 mg, 0.1 mmol) and **HBD** (14.0 mg, 0.1 mmol) in 1:1 molar ratio were separately dissolved in 5 mL of acetone, filtered, and mixed. The resulting solution was covered with parafilm with small holes for slower evaporation and left at room temperature. Small colorless needles showed up on the vial walls after 24 h. Melting points for **PPAH** (144 °C), **HBD** (161 °C) and 1:1 cocrystal **PPAH·HBD**(acetone solvate) (108 °C) were measured on SRS MeltTemp apparatus. All samples were heated with a heat rate 1 °C/min until the substance was completely liquid.

2.3. Single-Crystal X-ray Diffraction Analysis

X-ray diffraction data for **PPA**, **PPAH**, and **PPA·2HBA** cocrystal, were collected on a SMART APEX II CCD diffractometer (Bruker AXS, Madison, WI [27] using graphite-monochromatized Mo K α radiation ($\lambda = 0.71073$ Å). For **PPA·HBA** and **PPAH·HBD** cocrysts-

tals, X-ray diffraction data were collected on a Bruker D8 VENTURE diffractometer with microfocus sealed tube using graphite-monochromatized Cu K α radiation ($\lambda = 1.54178 \text{ \AA}$). SADABS program was used for absorption correction and scaling of observed data [28]. The structures were solved by direct methods and refined by full-matrix least-squares on F^2 for all data, using SHELXS97 [29] and OLEX 2.0 [30] software suites. The nonhydrogen atoms were refined in anisotropic approximation. The disorder was resolved for **PPA** in the pure form and in the 1:1 cocrystal **PPA·HBA**. The C–H atoms were fixed at idealized positions and refined with a riding-model approximation: C–H = 0.95–1.00 $\text{\AA}$ with Uiso(H) = 1.5 Ueq (C-methyl) and 1.2 Ueq (C) for other H atoms. The N–H and O–H hydrogens were located from the difference Fourier map or constrained. In cocrystal **PPAH·HBD** acetone solvent was disordered and the SQUEEZE procedure [31] was used to treat the diffused solvent in voids. The electron counts and the volume of associated voids indicated approximately one acetone solvent molecule per asymmetric unit. The sum formula reflected this ration. A check of the final CIF files (CCDC no.2247133–2247137) using PLATON did not show any missed symmetry. MERCURY2022.3.0 program (Cambridge Crystallographic Data Centre, Cambridge, UK) was used to make the molecular graphics [32].

2.4. Quantum Chemical Calculations

For the comparison of different enantiomers and their conformers quantum chemical calculations of total energy of **PPA** molecules in pure form (4 conformers), and in two cocrystals with **HBA**, **PPAH** in pure form and in cocrystal with **HBD** were carried out using GAUSSIAN 09 program package [33] and the B3LYP/6-311++G(d,p) level of theory [34]. The experimental X-ray coordinates were used for calculations. Correction on zero-point energy (ZPE) was included into molecular energy. A summary of results is presented in Table S1.

3. Results

Single crystal X-ray diffraction data, data collection, and structure refinement details are summarized in Table 1. Hydrogen bonding interactions were analyzed and finally calculated using the HTAB instruction in SHELXL [29], see Table 2.

Table 1. Crystal data for pure forms **PPA** and **PPAH** and cocrystals **PPA·HBA**, **PPA·2HBA**, and **PPAH·HBD**·(acetone solvate).

Compound	PPA	PPA·HBA	PPA·2HBA	PPAH	PPAH·HBD ^a
CCDC number	2247135	2247137	2247134	2247133	2247136
Empirical formula	C ₁₂ H ₁₄ N ₂ O ₂	C ₁₉ H ₂₀ N ₂ O ₅	C ₂₆ H ₂₆ N ₂ O ₈	C ₁₅ H ₁₉ N ₃ O ₂	C ₂₅ H ₃₂ N ₄ O ₅
Formula weight	218.25	356.37	494.49	273.337	468.54
Temperature/K	150(2)	293(2)	150(2)	100(2)	173(2)
Crystal system	triclinic	monoclinic	triclinic	monoclinic	triclinic
Space group	<i>P</i> -1	<i>P</i> 2 ₁ / <i>c</i>	<i>P</i> -1	<i>P</i> 2 ₁ / <i>c</i>	<i>P</i> -1
<i>a</i> / $\text{\AA}$	6.0917(17)	18.9326(3)	7.9851(8)	5.5613(8)	5.2285(3)
<i>b</i> / $\text{\AA}$	10.782(3)	5.58410(10)	12.6900(12)	14.190(2)	13.2516(7)
<i>c</i> / $\text{\AA}$	17.502(5)	18.7881(4)	13.6456(13)	18.170(3)	18.4515(9)
α /°	76.700(4)	90	107.579(3)	90	104.612(4)
β /°	85.742(5)	118.9220(10)	104.040(3)	91.034(5)	92.217(3)
γ /°	81.827(5)	90	107.749(3)	90	92.539(3)

Table 1. Cont.

Compound	PPA	PPA·HBA	PPA·2HBA	PPAH	PPAH·HBD ^a
Volume/Å ³	1106.3(5)	1738.57(6)	1167.0(2)	1433.7(4)	1234.22(12)
Z	4	4	2	4	2
ρ_{calc} g/cm ³	1.310	1.361	1.407	1.266	1.261
μ /mm ^{−1}	0.091	0.825	0.105	0.086	0.726
F(000)	464	752	520	584	500
Crystal size/mm ³	0.6 × 0.2 × 0.2	0.34 × 0.34 × 0.06	0.3 × 0.2 × 0.2	0.3 × 0.15 × 0.05	0.21 × 0.1 × 0.08
Reflections collected	52871	39609	34754	2220	16835
Independent reflections	6646 [$R_{\text{int}} = 0.0958$]	2646 [$R_{\text{int}} = 0.0595$]	4124 [$R_{\text{int}} = 0.0842$]	2220	4272 [$R_{\text{int}} = 0.0451$]
Data/restraints/parameters	6646/254/406	2646/0/239	4124/190/452	2220/36/184	4272/0/277
Goodness-of-fit on F^2	1.028	1.059	1.044	1.034	0.857
Final R indexes R_1, wR_2 [$I \geq 2\sigma(I)$]	0.0484, 0.1090	0.0399, 0.1081	0.0438, 0.1033	0.0473, 0.1263	0.0519, 0.1426
Final R indexes [all data] R_1, wR_2	0.0957, 0.1300	0.0422, 0.1102	0.0787, 0.1189	0.0767, 0.1379	0.0582, 0.1523
Largest diff. peak/hole/e Å ^{−3}	0.25/−0.25	0.32/−0.19	0.30/−0.20	0.356/−0.196	0.64/−0.28

^a In the refinement, scattering from the disordered acetone molecule was taken into account with the “SQUEEZE” procedure.

Table 2. Hydrogen bonds in the studied cocrystals and in the PPAH pure form.

D-H...A	H...A/Å	d(D...A)/Å	∠D-H...A/°	Symmetry Transformation for Acceptor
PPA·HBA				
O(5)-H(5)...O(1)	1.84	2.6638(18)	178.9	$x, 5/2 - y, z + 1/2$
N(2)-H(2A)...O(4)	2.10	2.936(2)	163.6	$x, y + 1, z$
N(2)-H(2B)...O(4)	2.23	2.9751(19)	144.9	$-x, 1 - y, 1 - z$
O(3)-H(3)...O(2)	1.83	2.6329(17)	166.0	$x, y - 1, z$
PPA·2HBA				
N(2)-H(2A)...O(1)	1.97(3)	2.906(3)	174(2)	$2 - x, 1 - y, 1 - z$
N(2)-H(2B)...O(2)	2.12(3)	2.999(3)	169(2)	$1 - x, 1 - y, 1 - z$
O(3)-H(3B)...O(2)	1.78(3)	2.671(2)	176(2)	$3 - x, 2 - y, 1 - z$
O(5)-H(5B)...O(6)	1.710(17)	2.599(2)	175(2)	x, y, z
O(7)-H(7B)...O(4)	1.745(17)	2.636(2)	174(2)	$3 - x, 2 - y, 1 - z$
O(8)-H(8)...O(3)	1.95(3)	2.820(2)	174(2)	$1 - x, 1 - y, -z$
C(11)-H(11A)...O(6)	2.58	3.407(3)	142.6	$2 - x, 1 - y, 1 - z$
PPAH				
N(2)-H(2)...O(2)	2.09	2.948(2)	163.3	$2 - x, 1 - y, 1 - z$
C(10)-H(10B)...O(2)	2.53	3.474(3)	159.7	$x - 1, y, z$
C(15)-H(15B)...O(1)	2.53	3.387(3)	146.1	$2 - x, 1/2 + y, 3/2 - z$

Table 2. Cont.

D-H...A	H...A/Å	d(D...A)/Å	∠D-H...A/°	Symmetry Transformation for Acceptor
PPAH·HBD				
N(1)-H(1A)...O(2)	2.07	2.940(4)	172.5	$2 - x, 1 - y, 2 - z$
N(1)-H(1B)...O(2)	2.26	3.064(4)	151.2	$x - 1, y, z$
N(3)-H(3)...O(4)	2.07	2.924(4)	161.9	$1 - x, 1 - y, 1 - z$
O(1)-H(6)...O(3)	1.83(5)	2.665(4)	170(5)	x, y, z
C(2)-H(2)...O(3)	2.53	3.217(5)	129.2	x, y, z
C(17)-H(17B)...O(4)	2.53	3.423(4)	150.2	$x - 1, y, z$

3.1. Crystal Structures of PPA Pure Form, PPA·HBA and PPA·2HBA Cocrystals

Figure 1 reveals the contents of asymmetric units for PPA, PPA·HBA, and PPA·2HBA (the molecules with labelling schemes are summarized in Figure S1). To estimate if purchased PPA material represents a racemic mixture or enantiopure compound we carried out its diffraction study. It was found that this material was a racemate with two symmetrically independent molecular positions in the unit cell, each of which was occupied by either *R*- or *S*-enantiomers with almost equal ~50% occupation factor (Figure 1a). These results are very close to data previously published by Rekis et al. for *rac*-PPA [20]. The *rac*-PPA structure allowed the authors of the above mentioned paper to describe the molecular arrangement in this crystal as a solid solution of enantiomers. Such a situation is common for structures of enantiomers when *R*- and *S*-enantiomers are interchangeable due to their isosterism, or significant similarity of their molecular shapes [35–39]. It was also mentioned that conformational flexibility of enantiomers helps them to adjust to their positions in crystal lattice. For room-temperature crystal structure PPA·HBA, the disorder was not resolved during structure refinement (Figure 1b); however, thermal ellipsoids of the phenyl and pyrrolidone rings demonstrate hints of such disorder. For low-temperature structure PPA·2HBA the disorder model was found (Figure 1c) with occupancy factors of 0.510(6) and 0.490(6) for two disordered positions similar to the PPA pure form.

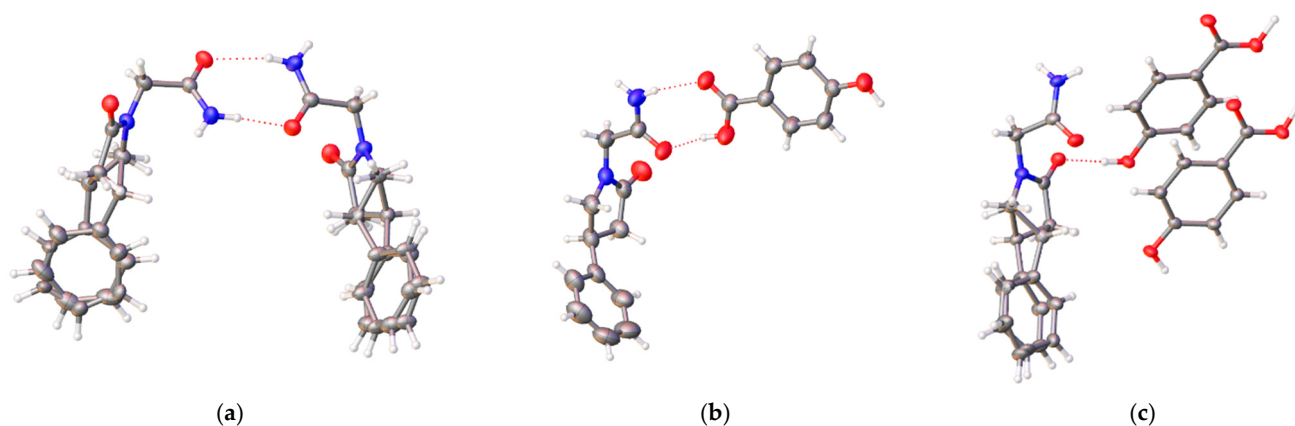


Figure 1. Contents of asymmetric units in: (a) PPA pure form, (b) PPA·HBA, and (c) PPA·2HBA cocrystals. Two disordered positions in PPA molecules are shown in Figure (a,c). Colorings scheme: oxygen—red; nitrogen—blue; carbon—grey; hydrogen—white, as used here and further.

Cocrystals PPA·HBA and PPA·2HBA were grown using the different ratios of starting materials (1:1 and 2:3, see Experimental Section). It is worth mentioning that variable stoichiometric ratios are not uncommon for API cocrystals [11,40,41].

The PPA molecule in PPA, PPA·HBA, and PPA·2HBA crystals demonstrates conformational flexibility. Figure 2 presents three examples of overlay of the molecular structures of the same enantiomers found in the above mentioned crystals. Absolute and relative molecular

energies of the enantiomers with different conformations are given in Table S1. It is possible to see that energy differences between these structures are not large and related mostly to different orientations of amide and phenyl groups attached to the oxopyrrolidine ring.

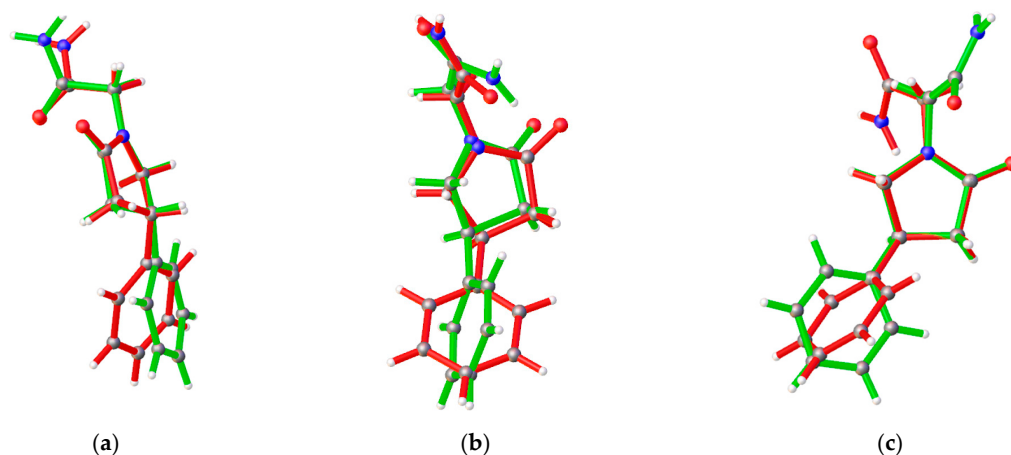


Figure 2. Selected examples of comparison of **PPA** molecular structure in crystals of **PPA·HBA**, and **PPA·2HBA**: (a) Overlay of *R*-enantiomers of **PPA** from **PPA·2HBA** (green) and from **PPA·HBA** (red); (b) Overlay of *S*-enantiomer of **PPA** from **PPA·2HBA** (green) and from **PPA·HBA** (red); (c) Overlay of *S*-enantiomer of **PPA** from **PPA·HBA** (green) and from **PPA·2HBA** (red).

In the 1:1 cocrystal, **PPA** and **HBA** molecules are interlinked via amide-carboxylic acid supramolecular heterosynthon ($N\cdots O = 2.936(2)$ Å, $O\cdots O = 2.6329(17)$ Å, Table 2) and dimerize in the centrosymmetric tetramer ($N\cdots O = 2.9751(19)$ Å) via fusion of three cyclic motifs in the sequence $R_2^2(8)R_4^2(8)R_2^2(8)$ (Figure 3a) [42]. Such tetrameric motif was previously registered in the **PA·HBA** cocrystal and in some other cocrystals with **HBA** coformer [11,43,44]. The H-bonded tetramers are further interlinked via the hydroxyl group of **HBA** and carbonyl of **PPA** molecules ($O\cdots O = 2.6638(18)$ Å), thus affording the H-bonded layer situated parallel to the *bc* plane. All H-bonds are concentrated within this layer. The layers stack along the crystallographic *a*-axis with interdigitating **PPA** phenyl rings (Figure 3b).

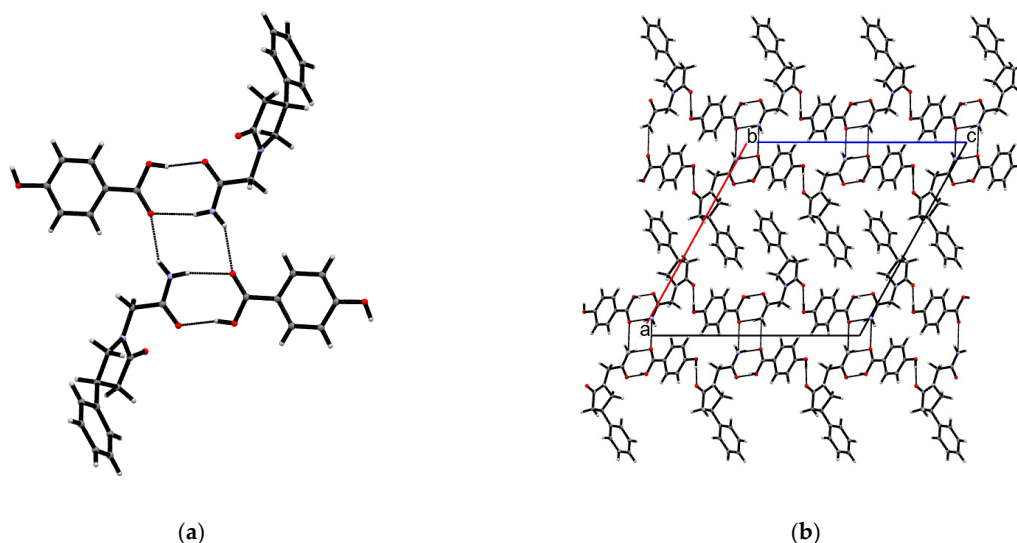


Figure 3. 1:1 cocrystal **PPA·HBA**: (a) View of $(PPA)_2(HBA)_2$ H-bonded tetramer. (b) Packing of two H-bonded layers in the crystal, view along the *b*-axis.

In the 1:2 cocrystal, **PPA** and **HBA** molecules are self-assembled in two separate homomeric H-bonded motifs that are interconnected. The **PPA** H-bonded chain is formed

via alternation of two centrosymmetric supramolecular homosynths, namely the typical amide-amide $R_2^2(8)$ homosynthon ($N\cdots O = 2.906(3) \text{ \AA}$) and the less common larger $R_2^2(14)$ homosynthon ($N\cdots O = 2.999(3) \text{ \AA}$) (Figure 4a). The similar infinite H-bonded motif was registered for **PA** in the **PA**·**HBA**·**H₂O** cocrystal [8] and for *rac*-**PPA** and *S*-**PPA** homomolecular forms (Figure 4b), except in the latter cases the amide-amide $R_2^2(8)$ homosynths are non-centrosymmetric [20]. In the reported cocrystal, the **PPA** homomeric chains interdigitate forming the layer parallel to the (0 1-1) plane (Figure 4c) and stabilized by significantly less directional dispersion interactions ($\pi(\text{Ph})\cdots\pi(\text{Ph})$, $\text{CH}(7)\cdots\pi(\text{Ph}) = 2.1 \text{ \AA}$) and weak $\text{C-H}\cdots\text{O}$ hydrogen bond ($\text{C}\cdots\text{O} = 3.42(2) \text{ \AA}$). In turn, two chemically identical but crystallographically distinct **HBA** cofomers are dimerized via robust $R_2^2(8)$ acid-acid supramolecular homosynthon ($\text{O}\cdots\text{O} = 2.671(2) \text{ \AA}$ and $2.599(2) \text{ \AA}$) and form the tail-to-tail H-bonded chain via hydroxyl-groups ($\text{O}\cdots\text{O} = 2.636(2) \text{ \AA}$) (Table 2). These carboxylic chains were also packed in the layers (Figure 4d). Similar to the 1:1 cocrystal, **HBA** and **PPA** interconnect through hydroxyl and carbonyl groups ($\text{O}\cdots\text{O} = 2.820(2) \text{ \AA}$) that results in the 3D hydrogen bonded network where the homomeric **PPA** and **HBA** layers alternate (Figure S2). In addition to the already mentioned **PA**-hydroquinone cocrystal [8], **PA** and citric acid in cocrystal also form homomeric H-bonded domains that interconnect [10].

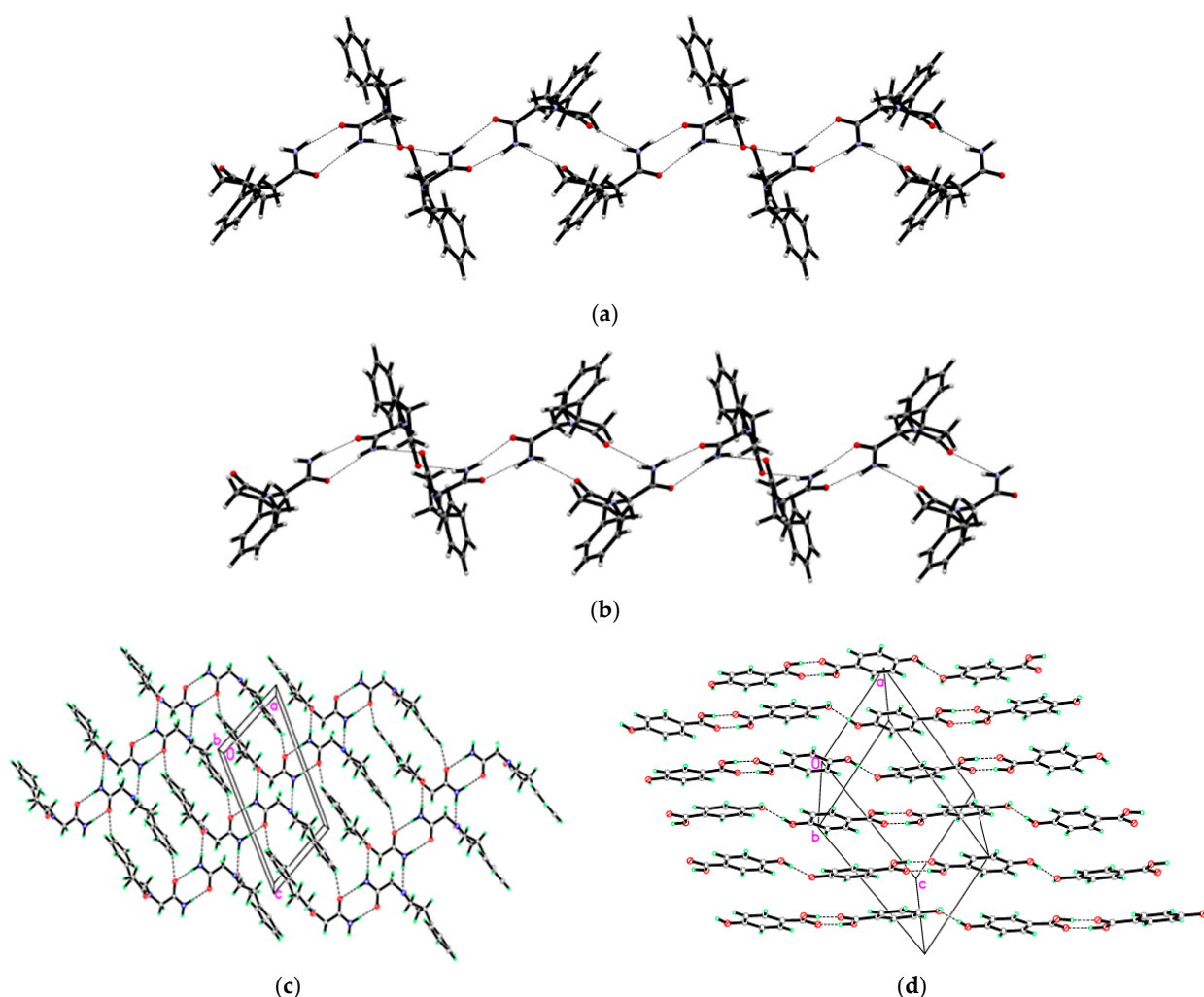


Figure 4. Comparison of molecular synthons in *rac*-**PPA** and 1:2 cocrystal **PPA**·**2HBA**: (a) View of the **PPA** homomeric chain in **PPA**·**2HBA**; (b) **PPA** homomeric chain in *rac*-**PPA**; (c) Packing of **PPA** chains in the layer in cocrystal via $\text{CH}\cdots\text{O}$ short contacts and π - π stacking interactions between the phenyl rings. (d) Packing of **HBA** chains in stacking layer.

3.2. Crystal Structures of PPAH Pure Form and PPAH·HBD Cocrystal

Asymmetric units of **PPAH** and **PPAH·HBD** (the disordered acetone solvent molecule was excluded) crystals and overlapping diagram for **PPAH** from these two solids are shown in Figure 5a–c (for numbering see Figure S1). The geometrical parameters of the **PPAH** molecule are similar in the pure form and in the cocrystal. These similarities are evident from the overlay of these molecules presented in Figure 5c. Molecular energies of the **PPAH** molecule in pure crystal and in cocrystal are very close (Table S2).

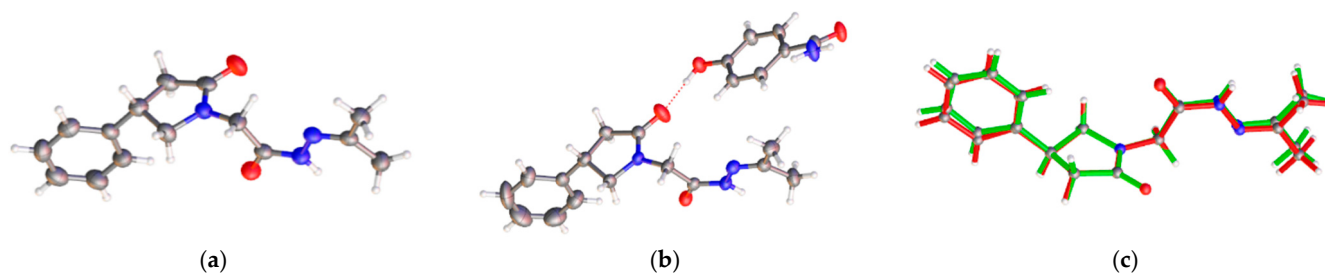


Figure 5. Content of asymmetric units in: (a) **PPAH**, (b) **PPAH·HBD** (see Supplementary Materials for structures with labels), and (c) overlay diagram for **PPAH** from pure material (green) and from cocrystal with **HBD** (red).

PPAH crystallizes in the monoclinic $P2_1/c$ space group with one molecule in the asymmetric unit. The expected dimerization of molecules occurs via amide-amide homosynthon, $N\cdots O = 2.948(2)$ Å (Figure 6a). The dimers pack in stacks along the shortest a -axis with only weak intermolecular $CH\cdots O$ interactions between stacks (Table 2, Figure 5b).

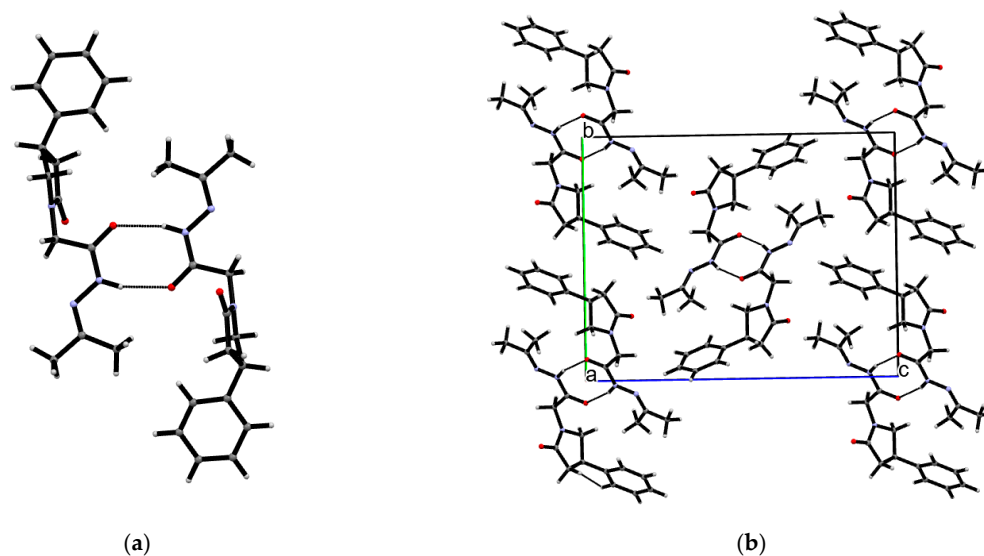


Figure 6. (a) View of centrosymmetric dimer in **PPAH**, (b) packing of **PPAH** dimers in crystal, view along the a -axis.

In the **PPAH·HBD** cocrystal, the self-association of each component occurs via amide-amide homosynthon (Figure 7a,b). Similar to the pure form, the **PPAH** molecules in cocrystal form the centrosymmetric dimers ($N\cdots O = 2.924(4)$ Å), while **HBD** coformers form the H-bonded double chain where the amide-amide homosynthon are further interlinked via synthon's sequence $\dots R_2^2(8)R_4^2(8)R_2^2(8)\dots$ ($N\cdots O = 2.940(4)$, $3.064(4)$ Å, Table 2) identical with that found for $(PPA)_2(HBA)_2$ tetramers (Figure 2a). The discrete **PPAH** dimers and roughly planar **HBD** chains interconnect through **HBD** hydroxyl and **PPAH** carbonyl groups ($O\cdots O = 2.665(4)$ Å) resulting in the H-bonded layer (Figure 7c). An interesting feature is that the stacking pattern of **PPAH** in the pure form was conserved in

cocrystal strengthening by the **PPAH**⋯**HBD** coformer H-bond (Figure S3). The packing of the layers remains voids of 241 Å³ filled by the disordered acetone solvent (Figure S4).

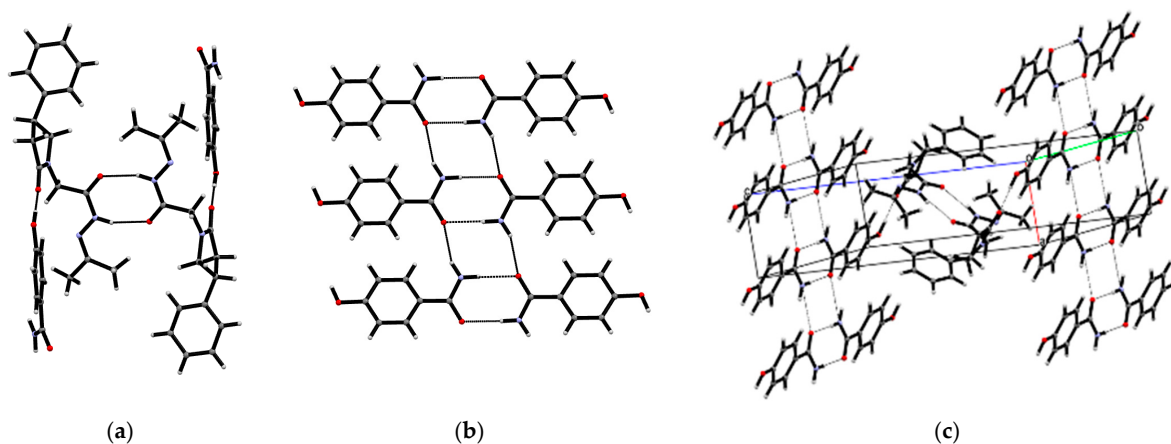


Figure 7. Cocrystal **PPAH·HBD**: (a) View of **PPAH** dimerization via amide-amide homosynthon with two **HBD** cofomers attached to dimer via OH⋯O hydrogen bonds; (b) H-bonded double chain of **HBD** cofomers; (c) Fragment of **PPAH·HBD** H-bonded layer, view along the *b*-axis.

Quantum chemical calculations of **PPA** and **PPAH** conformers found in the pure materials and cocrystals (see Supplementary Materials) demonstrated that energy differences for most conformers do not exceed 2 kcal/mol. Such a difference suggests that conformational variations of these molecules are mostly defined by influence of packing conditions (in this case *R*- and *S*-enantiomers intend to occupy same molecular position in crystal) on molecular conformation.

4. Conclusions

Two cocrystals of phenylpiracetam (**PPA**) with 4-hydroxybenzylcarboxylic acid (**HBA**) coformer with different stoichiometry and a cocrystal of 2-(4-phenyl-2-oxopyrrolidin-1-yl)-*N'*-isopropylideneacetohydrazide (**PPAH**) with 4-hydroxybenzamide (**HBD**) coformer were obtained by cocrystallization experiments. Despite the fact that both racetams contain a primary amide as a main functional group for hydrogen bonding, the structures of reported cocrystals are significantly different and dependent on the interplay of amide-amide, amide-carboxylic acid, carboxylic acid-carboxylic acid supramolecular homo- and heterosynthons. Otherwise, the hydroxyl–carbonyl heterosynthon was found in all three cocrystal structures with both cofomers. The H-bonded homomeric chain found in the **PPA** *rac*- and enantiopure forms, was also present in the 1:2 cocrystal resulting in alternation of **PPA** and **HBA** homomeric regions. The dimerization of **PPAH** via amide-amide supramolecular homosynthon led to the identical stacking of this homodimer as a dominating motif both in the cocrystal and in the racetam pure form. **PPA** in two cocrystals, **PPA·HBA** and **PPA·2HBA** demonstrated conformational freedom originating from the lack of strong intermolecular interactions with participation of phenyl and oxopyrrolidine rings that was reflected in disorder registered for these moieties in the crystals. The described situation allowed us to consider these cocrystals as examples of partial solid solutions.

Supplementary Materials: The following supporting information can be downloaded at: <https://www.mdpi.com/article/10.3390/chemistry5020074/s1>, Figure S1. ORTEP drawings with labelling schemes; Figure S2. Alternation of homomeric **PPA** and **HBA** H-bonded layers in cocrystal **PPA·2HBA**; Figure S3. Similar packing motifs in **PPAH** and **PPAH·HBD** cocrystal; Figure S4. Packing of **PPAH·HBD** H-bonded layers with voids filled by the disordered acetone solvent molecules; Table S1. Relative energies of **PPA** enantiomers with different conformations according to DFT calculations on B3LYP/6-311++(d,p) level of theory, kcal/mol; Table S2. Relative energies of **PPAH** enantiomers with different conformations according to DFT calculations on B3LYP/6-311++(d,p) level of theory, kcal/mol.

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