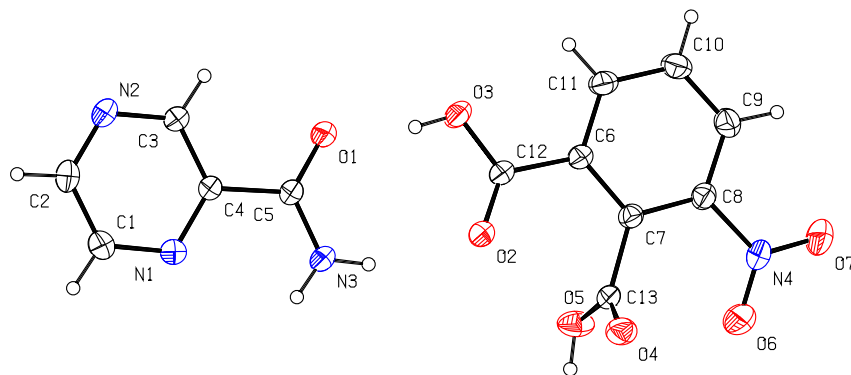


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# The crystal structure of 3-nitrobenzene-1,2-dicarboxylic acid–pyrazine-2-carboxamide(1/1), $C_{13}H_{10}N_4O_7$



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## Abstract

$C_{13}H_{10}N_4O_7$ , monoclinic,  $P2_1/n$  (no. 14),  $a = 10.7017(7)$  Å,  $b = 7.1240(5)$  Å,  $c = 19.0878(13)$  Å,  $\beta = 99.280(3)^\circ$ ,  $V = 1436.19(17)$  Å<sup>3</sup>,  $Z = 4$ ,  $R_{gt}(F) = 0.0586$ ,  $wR_{ref}(F^2) = 0.1746$ ,  $T = 296.15$  K.

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The molecular structure is shown in the figure. Table 1 contains crystallographic data and Table 2 contains the list of the atoms including atomic coordinates and displacement parameters.

## Source of materials

The mixture of pyrazinamide and 3-nitrophthalic acid with a molar ratio of 1:1 was dissolved in deionized

**Table 1:** Data collection and handling.

|                                                       |                                           |
|-------------------------------------------------------|-------------------------------------------|
| Crystal:                                              | Colourless block                          |
| Size:                                                 | 0.48 × 0.35 × 0.29 mm                     |
| Wavelength:                                           | Mo K $\alpha$ radiation (0.71073 Å)       |
| $\mu$ :                                               | 0.13 mm <sup>-1</sup>                     |
| Diffractometer, scan mode:                            | Bruker D8 Venture, $\varphi$ and $\omega$ |
| $\theta_{max}$ , completeness:                        | 27.2°, >99 %                              |
| $N(hkl)_{measured}$ , $N(hkl)_{unique}$ , $R_{int}$ : | 20,755, 3189, 0.034                       |
| Criterion for $I_{obs}$ , $N(hkl)_{gt}$ :             | $I_{obs} > 2\sigma(I_{obs})$ , 2684       |
| $N(param)_{refined}$ :                                | 219                                       |
| Programs:                                             | Bruker [1], Olex2 [2], SHELX [3, 4]       |

water, and the resulting mixed solution was stirred and dissolved in a constant temperature water bath at 30 °C to obtain a clear solution. The solution was filtered and placed in a sample vial, covered with membrane and punctured. The filtrate was slowly evaporated at room temperature to obtain crystals of the title compound.

## Experimental details

Absorption corrections were applied by using multi-scan program [1]. Using Olex2 [2], the structure was solved with the ShelXT [3] structure solution program and refined with the ShelXL [4] refinement package. The H atoms were fixed, fixed  $U_{iso}$  were set to 1.2 times of all C(H) groups and N(H, H) groups; 1.5 times of O(H) groups.

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**Table 2:** Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (Å<sup>2</sup>).

| Atom | x            | y           | z            | <i>U</i> <sub>iso</sub> */ <i>U</i> <sub>eq</sub> |
|------|--------------|-------------|--------------|---------------------------------------------------|
| O1   | 0.56030 (14) | 0.6521 (2)  | 0.80568 (8)  | 0.0554 (4)                                        |
| N1   | 0.39272 (16) | 1.0651 (2)  | 0.83422 (10) | 0.0491 (4)                                        |
| N2   | 0.60064 (18) | 1.1038 (3)  | 0.94184 (10) | 0.0539 (5)                                        |
| N3   | 0.36314 (15) | 0.7425 (2)  | 0.75681 (9)  | 0.0475 (4)                                        |
| H3A  | 0.352312     | 0.647621    | 0.728643     | 0.057*                                            |
| H3B  | 0.303967     | 0.824517    | 0.756137     | 0.057*                                            |
| C1   | 0.4089 (2)   | 1.2159 (3)  | 0.87595 (14) | 0.0618 (6)                                        |
| H1   | 0.349051     | 1.311502    | 0.868907     | 0.074*                                            |
| C2   | 0.5118 (2)   | 1.2347 (3)  | 0.92937 (13) | 0.0597 (6)                                        |
| H2   | 0.518990     | 1.342165    | 0.957370     | 0.072*                                            |
| C3   | 0.58626 (19) | 0.9533 (3)  | 0.89993 (10) | 0.0463 (5)                                        |
| H3C  | 0.647217     | 0.859162    | 0.906400     | 0.056*                                            |
| C4   | 0.48247 (16) | 0.9344 (2)  | 0.84700 (9)  | 0.0371 (4)                                        |
| C5   | 0.47010 (17) | 0.7623 (3)  | 0.80088 (10) | 0.0382 (4)                                        |
| O2   | 0.38517 (13) | 0.3770 (2)  | 0.68504 (8)  | 0.0546 (4)                                        |
| O3   | 0.58489 (14) | 0.3501 (3)  | 0.73787 (10) | 0.0666 (5)                                        |
| H3   | 0.570195     | 0.452311    | 0.754514     | 0.100*                                            |
| O4   | 0.19355 (13) | 0.1258 (2)  | 0.59364 (8)  | 0.0515 (4)                                        |
| O5   | 0.31431 (14) | 0.2893 (2)  | 0.53055 (8)  | 0.0550 (4)                                        |
| H5   | 0.246717     | 0.335789    | 0.512353     | 0.083*                                            |
| O6   | 0.2637 (2)   | −0.0980 (3) | 0.48266 (13) | 0.1094 (10)                                       |
| O7   | 0.3765 (2)   | −0.3365 (3) | 0.48143 (12) | 0.0877 (7)                                        |
| N4   | 0.35091 (17) | −0.1942 (2) | 0.50934 (9)  | 0.0474 (4)                                        |
| C6   | 0.50408 (16) | 0.1122 (3)  | 0.65828 (9)  | 0.0374 (4)                                        |
| C7   | 0.41254 (16) | 0.0484 (2)  | 0.60224 (9)  | 0.0350 (4)                                        |
| C8   | 0.43686 (17) | −0.1239 (3) | 0.57199 (9)  | 0.0383 (4)                                        |
| C9   | 0.54172 (19) | −0.2334 (3) | 0.59650 (11) | 0.0460 (5)                                        |
| H9   | 0.552355     | −0.349845   | 0.576264     | 0.055*                                            |
| C10  | 0.6298 (2)   | −0.1668 (3) | 0.65125 (11) | 0.0507 (5)                                        |
| H10  | 0.701304     | −0.237663   | 0.668069     | 0.061*                                            |
| C11  | 0.61190 (18) | 0.0054 (3)  | 0.68128 (10) | 0.0463 (5)                                        |
| H11  | 0.672964     | 0.050936    | 0.717525     | 0.056*                                            |
| C12  | 0.48513 (17) | 0.2934 (3)  | 0.69422 (10) | 0.0396 (4)                                        |
| C13  | 0.29354 (16) | 0.1587 (2)  | 0.57572 (9)  | 0.0364 (4)                                        |

## Comment

Pyrazinamide (PZA), a white crystalline powder, is the first-line drug for the treatment of pulmonary tuberculosis and the most effective antibacterial drug in short-course chemotherapy. Due to the low water solubility (15 µg/mL) and low bioavailability, its oral absorption effect is severely inhibited [5]. In recent years, drug co-crystal technology, as an emerging technology, enhances the efficacy of drugs by improving the solubility and permeability of drugs [6]. Therefore, PZA can be co-crystallized to improve its solubility and bioavailability. Cocrystals of pyrazinamide were obtained with 3-(4-hydroxyphenyl)prop-2-enoic acid (CCDC: 2026373, JAJYUS), thiophene-2,5-dicarboxylic acid (CCDC: 2026372, JAJZON), 2,6-dichlorobenzoic acid

(CCDC: 2026371, JAJZED), 5-hydroxybenzene-1,3-dicarboxylic acid (CCDC: 2069727, JAJZIH) [7], hydrochlorothiazide (CCDC: 1510703, EGENIP01) [8], 4-nitrobenzoic acid (CCDC: 963629, MUDVUE) [9] and so on.

In the title crystal, the main intermolecular force is the formation of hydrogen bonds. The O2 atom on the carboxyl groups of 3-nitrophthalic acid is a hydrogen bond acceptor. The H3A and H3B atoms on the amide group (N3–H3) of the pyrazinamide molecule are hydrogen bond donors, forming the intermolecular hydrogen bond N3–H3A...O2 [length 2.969(2) Å, angle 138.0°]. The O1 atom on the amide group of the pyrazinamide molecule is a hydrogen bond acceptor, and the H2 atom on the carboxyl group (O3–H3) of the 3-nitrophthalic acid molecule is a hydrogen bond donor, forming the intermolecular hydrogen bond O3–H3...O1 [length 2.546(2) Å, angle 167.8°]. The N1 atom on the pyrazine ring of the pyrazinamide molecule is a hydrogen bond acceptor, the H5 atom on the carboxyl group (O5–H5) of the 3-nitrophthalic acid molecule is a hydrogen bond donor, forming an intermolecular hydrogen bond O3–H3...O1 (symmetry code:  $-1/2 + x, 3/2 - y, -1/2 + z$ ) [length 2.725(2) Å, angle 160.2°].

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