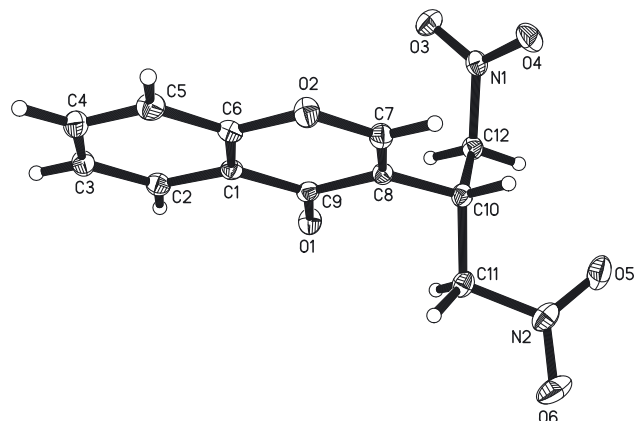


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# Crystal structure of 3-(1,3-dinitropropan-2-yl)-4*H*-chromen-4-one, C<sub>12</sub>H<sub>10</sub>N<sub>2</sub>O<sub>6</sub>



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

|                                                                            |                                                    |
|----------------------------------------------------------------------------|----------------------------------------------------|
| Crystal:                                                                   | Yellow block                                       |
| Size:                                                                      | 0.13 × 0.12 × 0.10 mm                              |
| Wavelength:                                                                | Mo K $\alpha$ radiation (0.71073 Å)                |
| $\mu$ :                                                                    | 0.13 mm <sup>-1</sup>                              |
| Diffractometer, scan mode:                                                 | Bruker APEX-II, $\varphi$ and $\omega$             |
| $\theta_{\max}$ , completeness:                                            | 25.0°, >99%                                        |
| $N(hkl)_{\text{measured}}$ , $N(hkl)_{\text{unique}}$ , $R_{\text{int}}$ : | 5022, 2091, 0.028                                  |
| Criterion for $I_{\text{obs}}$ , $N(hkl)_{\text{gt}}$ :                    | $I_{\text{obs}} > 2 \sigma(I_{\text{obs}})$ , 1740 |
| $N(\text{param})_{\text{refined}}$ :                                       | 181                                                |
| Programs:                                                                  | SHELX [1], Bruker [2]                              |

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## Abstract

C<sub>12</sub>H<sub>10</sub>N<sub>2</sub>O<sub>6</sub>, monoclinic,  $P2_1/c$  (no. 14),  $a = 14.6017(10)$  Å,  $b = 6.8095(4)$  Å,  $c = 13.0101(8)$  Å,  $\beta = 113.075(7)^\circ$ ,  $V = 1190.10(14)$  Å<sup>3</sup>,  $Z = 4$ ,  $R_{\text{gt}}(F) = 0.0405$ ,  $wR_{\text{ref}}(F^2) = 0.0976$ ,  $T = 293$  K.

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Table 1 contains crystallographic data and Table 2 contains the list of the atoms including atomic coordinates and displacement parameters.

## Source of material

The mixture of 4-oxo-4*H*-chromene-3-carbaldehyde (261 mg, 1.5 mmol), di-*tert*-butyl malonate (368 mg, 1.7 mmol) and 4-dimethylaminopyridine (DMAP, 12 mg, 0.1 mmol) was stirred in nitromethane (50 mL) at room temperature for 40 min. The reaction mixture was poured into saturated NaCl

solution (100 mL), and a yellow precipitate appeared. The precipitate was filtered and washed with saturated NaCl solution. The yellow precipitate was recrystallized from ethyl acetate to give 3-(1,3-dinitropropan-2-yl)-4*H*-chromen-4-one.

## Experimental details

All hydrogen atoms were identified in difference Fourier syntheses and placed in geometrically idealized positions. The  $U_{\text{iso}}$  values of the hydrogen atoms of methyl groups were set to  $1.5U_{\text{eq}}(\text{C})$  and the  $U_{\text{iso}}$  values of all other hydrogen atoms were set to  $1.2U_{\text{eq}}(\text{C})$ .

## Comment

Chromone derivatives are naturally occurring compounds spread in the plant kingdom [3], and therefore present in representative amounts in the normal human diet [4]. Some chromone derivatives possess biological activities including antiinflammatory [5], anti-HIV [6], anticancer [7], antibacterial [8], antimalarial [9], and antitumor [10], as well as treatment of Alzheimer's disease [11].

The molecular structure of title compound (*cf.* the figure) consists of a benzene ring A (C1–C6), a pyranone ring B (C1/C6–C9/O2) and a 1,3-dinitropropyl group. Ring A and ring B constitute the chromone skeleton. The chromone skeleton is essentially planar, with mean deviation from this plane of 0.0054(3) Å. The dihedral angle between rings A and B is 0.72(2)°, similar to the value found in literature [12, 13]. In the crystal structure of title compound, the chromone skeleton is stabilized by three weak intramolecular

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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> <sup>*</sup> / <i>U</i> <sub>eq</sub> |
|------|---------------|------------|--------------|---------------------------------------------------------------|
| C1   | 0.44312 (13)  | 0.7713 (3) | 0.54597 (14) | 0.0166 (4)                                                    |
| C2   | 0.51132 (13)  | 0.7369 (3) | 0.49654 (15) | 0.0202 (4)                                                    |
| H2   | 0.488382      | 0.703780   | 0.421264     | 0.024*                                                        |
| C3   | 0.61178 (14)  | 0.7517 (3) | 0.55845 (16) | 0.0228 (4)                                                    |
| H3   | 0.656617      | 0.727432   | 0.525213     | 0.027*                                                        |
| C4   | 0.64678 (14)  | 0.8033 (3) | 0.67131 (16) | 0.0248 (5)                                                    |
| H4   | 0.714913      | 0.813757   | 0.712690     | 0.030*                                                        |
| C5   | 0.58145 (14)  | 0.8387 (3) | 0.72175 (15) | 0.0235 (5)                                                    |
| H5   | 0.604702      | 0.873016   | 0.796866     | 0.028*                                                        |
| C6   | 0.48016 (14)  | 0.8223 (3) | 0.65837 (15) | 0.0187 (4)                                                    |
| C7   | 0.31896 (13)  | 0.8486 (3) | 0.65442 (15) | 0.0202 (4)                                                    |
| H7   | 0.277923      | 0.874945   | 0.692265     | 0.024*                                                        |
| C8   | 0.27491 (13)  | 0.8022 (3) | 0.54564 (14) | 0.0163 (4)                                                    |
| C9   | 0.33550 (13)  | 0.7581 (3) | 0.48188 (14) | 0.0175 (4)                                                    |
| C10  | 0.16258 (13)  | 0.7980 (3) | 0.48815 (15) | 0.0183 (4)                                                    |
| H10  | 0.134810      | 0.841982   | 0.541668     | 0.022*                                                        |
| C11  | 0.12853 (13)  | 0.9405 (3) | 0.38950 (15) | 0.0213 (4)                                                    |
| H11A | 0.174095      | 1.050908   | 0.407157     | 0.026*                                                        |
| H11B | 0.131285      | 0.874663   | 0.324655     | 0.026*                                                        |
| C12  | 0.12402 (14)  | 0.5901 (3) | 0.44964 (14) | 0.0198 (4)                                                    |
| H12A | 0.052132      | 0.592338   | 0.411119     | 0.024*                                                        |
| H12B | 0.152104      | 0.541700   | 0.398153     | 0.024*                                                        |
| N1   | 0.15299 (11)  | 0.4580 (2) | 0.54878 (12) | 0.0208 (4)                                                    |
| N2   | 0.02524 (12)  | 1.0153 (3) | 0.36135 (13) | 0.0260 (4)                                                    |
| O1   | 0.29803 (9)   | 0.7139 (2) | 0.38197 (10) | 0.0224 (3)                                                    |
| O2   | 0.41817 (9)   | 0.8598 (2) | 0.71304 (10) | 0.0229 (3)                                                    |
| O3   | 0.20439 (10)  | 0.3135 (2) | 0.55230 (11) | 0.0290 (4)                                                    |
| O4   | 0.12560 (10)  | 0.5022 (2) | 0.62346 (11) | 0.0295 (4)                                                    |
| O5   | −0.03394 (11) | 0.9161 (2) | 0.38404 (13) | 0.0362 (4)                                                    |
| O6   | 0.00517 (12)  | 1.1751 (2) | 0.31429 (13) | 0.0418 (4)                                                    |

hydrogen bonds, with C11—H11B⋯O1 to be 2.502(4) Å, C12—H12B⋯O1 to be 2.512(5) Å and C12—H12A⋯O5 to be 2.493(3) Å. Additionally, hydrogen bonds C2—H2⋯O2 and C7—H7⋯O7 connect neighboring molecules to form chain structure along *c* axis. Neighboring molecules were linked together through C11—H11⋯O3, C11—H11B⋯O6 and C12—H12A⋯O4 to form another chain structure along *b* axis.

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**Conflict of interest statement:** The authors declare no conflicts of interest regarding this article.

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