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Sodium 2-nitrocinnamate dihydrate: a one-dimensional hydrogen-bonded coordination polymer

Graham Smith* and Urs D. Wermuth

School of Physical and Chemical Sciences, Queensland University of Technology, GPO Box 2434, Brisbane, Qld 4001, Australia
Correspondence e-mail: g.smith@qut.edu.au

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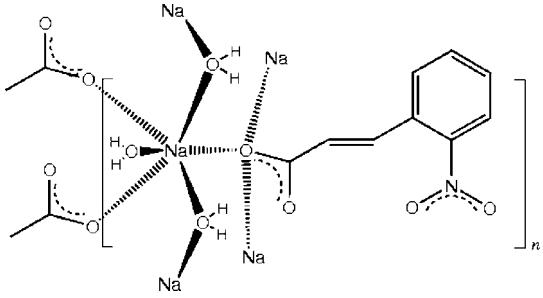
Key indicators: single-crystal X-ray study; T = 297 K; mean σ(C–C) = 0.002 Å; R factor = 0.038; wR factor = 0.113; data-to-parameter ratio = 12.4.

The title compound *catena*-poly[aquasodium-μ_2-aqua-μ_3-2-nitrocinnamato], $[Na(C_9H_6NO_4)(H_2O)_2]_n$, the sodium salt of *trans*-2-nitrocinnamic acid, is a one-dimensional coordination polymer based on six-coordinate octahedral NaO_6 centres, comprising three facially related monodentate carboxylate O-atom donors from separate ligands (all bridging) [Na—O = 2.4370 (13)–2.5046 (13) Å], and three water molecules (two bridging and one monodentate) [Na—O = 2.3782 (13)–2.4404 (17) Å]. The structure is also stabilized by intra-chain water–carboxylate and water–nitro O—H···O hydrogen bonds.

Related literature

For literature on similar compounds, see: Crowther *et al.* (2008); Kariuki *et al.* (1995); Kula *et al.* (2007); Schmidt (1964); Smith *et al.* (2006); Trividi *et al.* (2005).



Experimental

Crystal data

$[Na(C_9H_6NO_4)(H_2O)_2]$
M_r = 251.17
Monoclinic, $P2_1/c$
a = 19.4179 (7) Å
b = 3.6899 (2) Å
c = 14.8738 (7) Å
β = 92.239 (4)°
V = 1064.90 (9) Å^3
Z = 4
Mo $K\alpha$ radiation
μ = 0.17 mm^{-1}
T = 297 K
0.40 × 0.30 × 0.13 mm

Data collection

Oxford Diffraction Gemini-S CCD-detector diffractometer
Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996)
T_{min} = 0.93, T_{max} = 0.98
6531 measured reflections
2100 independent reflections
1626 reflections with $I > 2\sigma(I)$
R_{int} = 0.019

Refinement

$R[F^2 > 2\sigma(F^2)]$ = 0.038
$wR(F^2)$ = 0.113
S = 1.09
2100 reflections
170 parameters
H atoms treated by a mixture of independent and constrained refinement
$\Delta\rho_{max}$ = 0.30 e Å^{-3}
$\Delta\rho_{min}$ = −0.19 e Å^{-3}

Table 1
Hydrogen-bond geometry (Å, °).

D—H···A	D—H	H···A	D···A	D—H···A
O1W—H11W···O32^{i}	0.78 (3)	2.14 (3)	2.8871 (17)	162 (2)
O1W—H12W···O32^{ii}	0.89 (2)	1.90 (2)	2.7852 (17)	171 (2)
O2W—H21W···O21^{iii}	0.77 (3)	2.49 (3)	3.050 (2)	131 (3)
O2W—H22W···O32^{i}	0.91 (4)	2.04 (5)	2.882 (2)	153 (4)

Symmetry codes: (i) $x, -y+\frac{1}{2}, z-\frac{1}{2}$; (ii) $-x, -y, -z$; (iii) $x, -y+\frac{3}{2}, z-\frac{1}{2}$.

Data collection: *CrysAlis Pro* (Oxford Diffraction, 2009); cell refinement: *CrysAlis Pro*; data reduction: *CrysAlis Pro*; program(s) used to solve structure: *SIR92* (Altomare *et al.*, 1994); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *PLATON* (Spek, 2009); software used to prepare material for publication: *PLATON*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: SU2131).

References

Altomare, A., Cascarano, G., Giacovazzo, C., Guagliardi, A., Burla, M. C., Polidori, G. & Camalli, M. (1994). *J. Appl. Cryst.* **27**, 435.
Crowther, D., Chowdhury, M. & Kariuki, B. M. (2008). *J. Mol. Struct.* **872**, 64–71.
Kariuki, B. M., Valim, J. B., Jones, W. & King, J. (1995). *Acta Cryst.* C**51**, 1051–1053.
Kula, A., Mazur, L. & Rzaczynska, Z. (2007). *J. Coord. Chem.* **60**, 843–850.
Oxford Diffraction (2009). *CrysAlis Pro*. Oxford Diffraction Ltd, Yarnton, England.
Schmidt, G. M. J. (1964). *J. Chem. Soc.* pp. 2014–2021.
Sheldrick, G. M. (1996). *SADABS*, University of Göttingen, Germany.
Sheldrick, G. M. (2008). *Acta Cryst.* A**64**, 112–122.
Smith, G., Wermuth, U. D., Young, D. J. & White, J. M. (2006). *Acta Cryst.* E**62**, o2024–o2026.
Spek, A. L. (2009). *Acta Cryst.* D**65**, 148–155.
Trividi, D. R., Ballabh, A. & Dastidar, P. (2005). *J. Mater. Chem.* **15**, 2606–2614.

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Sodium 2-nitrocinnamate dihydrate: a one-dimensional hydrogen-bonded coordination polymer

Graham Smith and Urs D. Wermuth

S1. Comment

Although the structures of two polymorphs of *trans*-cinnamic acid have been determined (Schmidt, 1964; Smith *et al.*, 2006), the structures of neither *trans*-2-nitrocinnamic acid [(*E*)-3-(2-nitrophenyl)propenoic acid] nor any of its alkali metal salts are known, although the dicyclohexylaminium salt has been reported (Trividi *et al.*, 2005). The only structures of alkali metal compounds of analogous ring-substituted *trans*-cinnamic acids are the sodium complexes with 2-chloro-cinnamic acid (Kariuki *et al.*, 1995), 3-chlorocinnamic acid (Crowther *et al.*, 2008), and 4-hydroxy-2-methoxycinnamic acid (Kula *et al.*, 2007). We have now prepared the sodium salt of *trans*-2-nitrocinnamic acid, a dihydrate $[Na(C_9H_6NO_4)(H_2O)_2]_n$ and its structure is reported here.

The molecular structure of the title compound is illustrated in Fig. 1. The polymeric structure is based on octahadral six-coordinate NaO_6 centres comprising three facially related monodentate carboxylate O-donors from separate ligands (all bridging) [Na–O, 2.4370 (13)– 2.5046 (13) Å] and three water molecules (two bridging, one monodentate) [Na–O, 2.3782 (13)–2.4404 (17) Å]. These units are linked into one-dimensional coordination polymer chains which extend along direction [010] (Fig. 1). The structure is similar to that of the sodium 2-chlorocinnamate complex (Kariuki *et al.*, 1995). The polymer chains are stabilized by intra-chain water *O–H*···$O_{carboxylate}$ and *O–H*···O_{nitro}hydrogen bonds (Table 1).

In the substituted cinnamate ligand molecule, the nitro group is rotated out of the plane of the benzene ring [torsion angle C1–C2–N21–O22, 144.65 (17)°], while the carboxylate group is similarly non-coplanar [C11–C21–C31–O31, -169.51 (17)°].

S2. Experimental

The title compound was synthesized by heating together for 10 minutes under reflux 1 mmol quantities of *trans*-cinnamic acid [(*E*-3-(2-nitrophenyl)propenoic acid] and sodium carbonate in 50 ml of 50% ethanol-water. After concentration to *ca* 30 ml, partial rt evaporation of the hot-filtered solution gave thin colourless plate-like crystals, suitable for X-ray analysis.

S3. Refinement

The H-atoms of the water molecules were located in difference electron-density maps and were freely refined: O-H = 0.77 (3) - 0.91 (4) Å. The C-bound H-atoms were included in calculated positions and treated as riding atoms: C–H = 0.93 Å with U_{iso}(H) = 1.2U_{eq}(C).

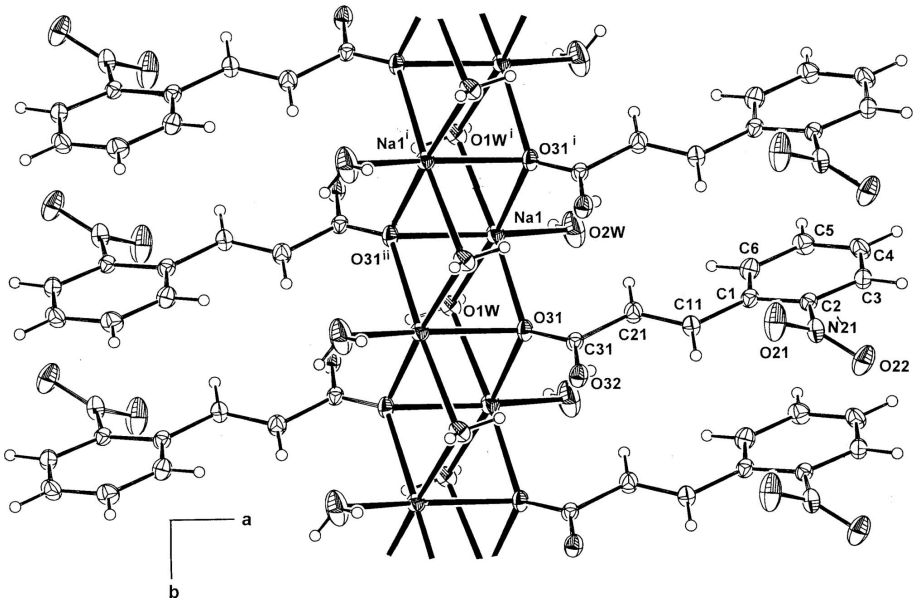

Figure 1

Molecular configuration and atom naming scheme for the title compound, showing the one-dimensional chain polymer structure extending along direction [010]. Displacement ellipsoids are drawn at the 50% probability level [Symmetry codes: (i) *x*, *y* + 1, *z*; (ii) -*x*, -*y* + 1, -*z*].

catena-poly[aquasodium-μ_2-aqua-μ_3-2-nitrocinnamato]

Crystal data

$[Na(C_9H_6NO_4)(H_2O)_2]$
$M_r = 251.17$
Monoclinic, $P2_1/c$
Hall symbol: -P 2ybc
$a = 19.4179\ (7)$ Å
$b = 3.6899\ (2)$ Å
$c = 14.8738\ (7)$ Å
$\beta = 92.239\ (4)°$
$V = 1064.90\ (9)$ Å^3
$Z = 4$
$F(000) = 520$
$D_x = 1.567$ Mg m^{-3}
Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å
Cell parameters from 2943 reflections
$\theta = 3.0–28.7°$
$\mu = 0.17$ mm^{-1}
$T = 297$ K
Plate, colourless
0.40 × 0.30 × 0.13 mm

Data collection

Oxford Diffraction Gemini-S CCD-detector diffractometer
Radiation source: Enhance (Mo) X-ray source
Graphite monochromator
ω scans
Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996)
$T_{min} = 0.93$, $T_{max} = 0.98$
6531 measured reflections
2100 independent reflections
1626 reflections with $I > 2\sigma(I)$
$R_{int} = 0.019$
$\theta_{max} = 26.0°$, $\theta_{min} = 3.0°$
$h = -23\rightarrow21$
$k = -4\rightarrow4$
$l = -18\rightarrow17$

Refinement

Refinement on F^2
Least-squares matrix: full
$R[F^2 > 2\sigma(F^2)] = 0.038$
$wR(F^2) = 0.113$
$S = 1.09$
2100 reflections
170 parameters
0 restraints
Primary atom site location: structure-invariant direct methods
Secondary atom site location: difference Fourier map
Hydrogen site location: inferred from neighbouring sites
H atoms treated by a mixture of independent and constrained refinement
$w = 1/[\sigma^2(F_o^2) + (0.0708P)^2]$ where $P = (F_o^2 + 2F_c^2)/3$
$(\Delta/\sigma)_{max} < 0.001$
$\Delta\rho_{max} = 0.30$ e Å^{-3}
$\Delta\rho_{min} = -0.19$ e Å^{-3}

Special details

Geometry. Bond distances, angles *etc*. have been calculated using the rounded fractional coordinates. All su's are estimated from the variances of the (full) variance-covariance matrix. The cell e.s.d.'s are taken into account in the estimation of distances, angles and torsion angles

Refinement. Refinement of F^2 against ALL reflections. The weighted *R*-factor *wR* and goodness of fit *S* are based on F^2, conventional *R*-factors *R* are based on *F*, with *F* set to zero for negative F^2. The threshold expression of $F^2 > \sigma(F^2)$ is used only for calculating *R*-factors(gt) *etc*. and is not relevant to the choice of reflections for refinement. *R*-factors based on F^2 are statistically about twice as large as those based on *F*, and *R*- factors based on ALL data will be even larger.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (Å²)

	x	*y*	*z*	U_{iso}*/U_{eq}
Na1	0.05494 (3)	0.73072 (17)	−0.06704 (4)	0.0272 (2)
O1W	0.02635 (6)	0.2374 (3)	−0.16440 (8)	0.0319 (4)
O2W	0.16105 (8)	0.6812 (5)	−0.14886 (12)	0.0691 (7)
O21	0.29871 (7)	0.4625 (6)	0.31366 (9)	0.0637 (6)
O22	0.39979 (7)	0.2280 (5)	0.32066 (10)	0.0541 (6)
O31	0.06940 (6)	0.2430 (3)	0.04255 (8)	0.0282 (4)
O32	0.10417 (6)	0.0743 (4)	0.18086 (8)	0.0336 (4)
N21	0.35246 (7)	0.3769 (4)	0.27927 (10)	0.0339 (5)
C1	0.30837 (8)	0.4562 (5)	0.12070 (11)	0.0267 (5)
C2	0.36252 (8)	0.4742 (4)	0.18535 (11)	0.0261 (5)
C3	0.42813 (9)	0.5880 (5)	0.16617 (12)	0.0327 (6)
C4	0.44154 (10)	0.6987 (5)	0.08063 (14)	0.0384 (6)
C5	0.38972 (10)	0.6860 (5)	0.01494 (13)	0.0368 (6)
C6	0.32514 (9)	0.5634 (5)	0.03431 (12)	0.0346 (6)
C11	0.23935 (9)	0.3141 (5)	0.13955 (12)	0.0297 (5)
C21	0.18383 (9)	0.3638 (5)	0.08772 (13)	0.0345 (6)
C31	0.11421 (8)	0.2155 (4)	0.10637 (11)	0.0256 (5)
H3	0.46290	0.58950	0.21090	0.0390*
H4	0.48520	0.78150	0.06720	0.0460*
H5	0.39850	0.76120	−0.04320	0.0440*
H6	0.29140	0.55150	−0.01180	0.0420*
H11	0.23490	0.17970	0.19190	0.0360*
H11W	0.0460 (11)	0.241 (6)	−0.209 (2)	0.055 (8)*
H12W	−0.0168 (12)	0.157 (7)	−0.1733 (17)	0.044 (8)*
H21	0.18810	0.50100	0.03580	0.0410*

H21W	0.1828 (16)	0.851 (9)	−0.138 (2)	0.093 (13)*
H22W	0.1500 (18)	0.666 (12)	−0.209 (2)	0.101 (14)*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Na1	0.0279 (4)	0.0275 (4)	0.0260 (4)	−0.0019 (3)	−0.0005 (3)	0.0007 (3)
O1W	0.0363 (7)	0.0350 (7)	0.0244 (7)	−0.0082 (5)	0.0026 (5)	−0.0002 (5)
O2W	0.0399 (9)	0.1143 (15)	0.0532 (10)	−0.0214 (9)	0.0046 (7)	0.0098 (9)
O21	0.0357 (8)	0.1222 (15)	0.0337 (8)	0.0081 (9)	0.0067 (6)	−0.0065 (9)
O22	0.0475 (9)	0.0775 (12)	0.0364 (9)	0.0164 (7)	−0.0096 (7)	0.0112 (7)
O31	0.0220 (6)	0.0370 (7)	0.0253 (6)	−0.0013 (5)	−0.0039 (5)	0.0011 (5)
O32	0.0290 (7)	0.0461 (8)	0.0255 (7)	0.0007 (5)	0.0005 (5)	0.0056 (6)
N21	0.0266 (8)	0.0471 (9)	0.0276 (8)	−0.0020 (7)	−0.0027 (6)	−0.0022 (7)
C1	0.0235 (8)	0.0281 (9)	0.0283 (9)	0.0035 (7)	0.0004 (7)	−0.0017 (7)
C2	0.0264 (8)	0.0257 (8)	0.0260 (9)	0.0040 (7)	0.0003 (7)	−0.0020 (7)
C3	0.0255 (9)	0.0331 (10)	0.0393 (11)	−0.0010 (8)	−0.0021 (7)	−0.0033 (8)
C4	0.0299 (10)	0.0362 (10)	0.0499 (13)	−0.0071 (8)	0.0106 (9)	−0.0017 (9)
C5	0.0405 (11)	0.0374 (10)	0.0331 (11)	−0.0045 (8)	0.0097 (8)	0.0035 (8)
C6	0.0340 (10)	0.0415 (11)	0.0281 (9)	0.0024 (8)	−0.0026 (7)	0.0020 (8)
C11	0.0261 (9)	0.0353 (10)	0.0276 (9)	−0.0015 (7)	0.0001 (7)	−0.0008 (7)
C21	0.0266 (9)	0.0435 (11)	0.0334 (10)	−0.0035 (8)	−0.0004 (7)	0.0093 (8)
C31	0.0231 (8)	0.0288 (9)	0.0248 (9)	0.0039 (7)	0.0008 (7)	−0.0028 (7)

Geometric parameters (Å, º)

Na1—O1W	2.3782 (13)	C1—C2	1.399 (2)
Na1—O2W	2.4404 (17)	C1—C6	1.395 (2)
Na1—O31	2.4370 (13)	C1—C11	1.476 (2)
Na1—O1W^{i}	2.4162 (13)	C2—C3	1.382 (2)
Na1—O31^{i}	2.5046 (13)	C3—C4	1.371 (3)
Na1—O31^{ii}	2.4577 (13)	C4—C5	1.376 (3)
O21—N21	1.222 (2)	C5—C6	1.374 (3)
O22—N21	1.217 (2)	C11—C21	1.314 (3)
O31—C31	1.267 (2)	C21—C31	1.494 (2)
O32—C31	1.247 (2)	C3—H3	0.9300
O1W—H11W	0.78 (3)	C4—H4	0.9300
O1W—H12W	0.89 (2)	C5—H5	0.9300
O2W—H21W	0.77 (3)	C6—H6	0.9300
O2W—H22W	0.91 (4)	C11—H11	0.9300
N21—C2	1.463 (2)	C21—H21	0.9300
O1W—Na1—O2W	79.67 (5)	C2—C1—C6	115.06 (15)
O1W—Na1—O31	81.96 (4)	C2—C1—C11	123.37 (15)
O1W—Na1—O1W^{i}	100.64 (4)	C6—C1—C11	121.46 (15)
O1W—Na1—O31^{i}	172.53 (5)	N21—C2—C1	121.34 (14)
O1W—Na1—O31^{ii}	85.02 (4)	N21—C2—C3	115.49 (14)
O2W—Na1—O31	101.58 (6)	C1—C2—C3	123.16 (15)

O1W^{i}—Na1—O2W	86.43 (5)	C2—C3—C4	119.49 (17)
O2W—Na1—O31^{i}	107.80 (5)	C3—C4—C5	119.26 (18)
O2W—Na1—O31^{ii}	158.48 (6)	C4—C5—C6	120.69 (18)
O1W^{i}—Na1—O31	171.93 (5)	C1—C6—C5	122.29 (16)
O31—Na1—O31^{i}	96.60 (4)	C1—C11—C21	124.71 (17)
O31—Na1—O31^{ii}	91.04 (4)	C11—C21—C31	124.59 (17)
O1W^{i}—Na1—O31^{i}	79.83 (4)	O31—C31—C21	115.57 (14)
O1W^{i}—Na1—O31^{ii}	81.62 (4)	O32—C31—C21	119.48 (15)
O31^{i}—Na1—O31^{ii}	87.68 (4)	O31—C31—O32	124.96 (15)
Na1—O1W—Na1^{iii}	100.64 (5)	C2—C3—H3	120.00
Na1—O31—C31	128.20 (10)	C4—C3—H3	120.00
Na1—O31—Na1^{iii}	96.60 (5)	C3—C4—H4	120.00
Na1—O31—Na1^{ii}	88.96 (4)	C5—C4—H4	120.00
Na1^{iii}—O31—C31	118.92 (10)	C4—C5—H5	120.00
Na1^{ii}—O31—C31	122.84 (10)	C6—C5—H5	120.00
Na1^{iii}—O31—Na1^{ii}	92.32 (4)	C1—C6—H6	119.00
H11W—O1W—H12W	112 (2)	C5—C6—H6	119.00
H21W—O2W—H22W	111 (4)	C1—C11—H11	118.00
O21—N21—C2	118.98 (14)	C21—C11—H11	118.00
O22—N21—C2	117.87 (14)	C11—C21—H21	118.00
O21—N21—O22	123.06 (16)	C31—C21—H21	118.00

O2W—Na1—O1W—Na1^{iii}	95.66 (6)	O31—Na1—O31^{ii}—C31^{ii}	−136.41 (11)
O31—Na1—O1W—Na1^{iii}	−7.75 (5)	Na1—O31—C31—O32	145.52 (13)
O1W^{i}—Na1—O1W—Na1^{iii}	180.00 (6)	Na1—O31—C31—C21	−35.11 (19)
O31^{ii}—Na1—O1W—Na1^{iii}	−99.51 (5)	Na1^{iii}—O31—C31—O32	−87.48 (18)
O1W—Na1—O31—C31	142.67 (13)	Na1^{iii}—O31—C31—C21	91.89 (14)
O1W—Na1—O31—Na1^{iii}	7.39 (4)	Na1^{ii}—O31—C31—O32	26.8 (2)
O1W—Na1—O31—Na1^{ii}	−84.82 (4)	Na1^{ii}—O31—C31—C21	−153.81 (11)
O2W—Na1—O31—C31	65.02 (14)	O21—N21—C2—C1	−38.6 (2)
O2W—Na1—O31—Na1^{iii}	−70.26 (6)	O21—N21—C2—C3	140.47 (18)
O2W—Na1—O31—Na1^{ii}	−162.47 (5)	O22—N21—C2—C1	144.65 (17)
O31^{i}—Na1—O31—C31	−44.72 (13)	O22—N21—C2—C3	−36.2 (2)
O31^{i}—Na1—O31—Na1^{iii}	180.00 (4)	C6—C1—C2—N21	178.85 (15)
O31^{i}—Na1—O31—Na1^{ii}	87.79 (4)	C6—C1—C2—C3	−0.2 (3)
O31^{ii}—Na1—O31—C31	−132.51 (13)	C11—C1—C2—N21	−4.8 (3)
O31^{ii}—Na1—O31—Na1^{iii}	92.21 (5)	C11—C1—C2—C3	176.18 (17)
O31^{ii}—Na1—O31—Na1^{ii}	0.00 (3)	C2—C1—C6—C5	−1.5 (3)
O1W—Na1—O1W^{i}—Na1^{i}	−180.00 (6)	C11—C1—C6—C5	−177.99 (17)
O2W—Na1—O1W^{i}—Na1^{i}	−101.22 (6)	C2—C1—C11—C21	164.51 (18)
O2W—Na1—O31^{i}—Na1^{i}	75.56 (6)	C6—C1—C11—C21	−19.4 (3)
O2W—Na1—O31^{i}—C31^{i}	−65.26 (12)	N21—C2—C3—C4	−177.36 (16)
O31—Na1—O31^{i}—Na1^{i}	180.00 (3)	C1—C2—C3—C4	1.7 (3)
O31—Na1—O31^{i}—C31^{i}	39.19 (12)	C2—C3—C4—C5	−1.6 (3)
O1W—Na1—O31^{ii}—Na1^{ii}	81.83 (4)	C3—C4—C5—C6	−0.1 (3)
O1W—Na1—O31^{ii}—C31^{ii}	−54.58 (11)	C4—C5—C6—C1	1.7 (3)
O2W—Na1—O31^{ii}—Na1^{ii}	126.42 (15)	C1—C11—C21—C31	179.16 (16)

O2W—Na1—O31^{ii}—C31^{ii}	−10.0 (2)	C11—C21—C31—O31	−169.51 (17)
O31—Na1—O31^{ii}—Na1^{ii}	0.00 (5)	C11—C21—C31—O32	9.9 (3)

Symmetry codes: (i) *x*, *y*+1, *z*; (ii) −*x*, −*y*+1, −*z*; (iii) *x*, *y*−1, *z*.

Hydrogen-bond geometry (Å, º)

D—H···*A*	*D*—H	H···*A*	*D*···*A*	*D*—H···*A*
O1*W*—H11*W*···O32^{iv}	0.78 (3)	2.14 (3)	2.8871 (17)	162 (2)
O1*W*—H12*W*···O32^{v}	0.89 (2)	1.90 (2)	2.7852 (17)	171 (2)
O2*W*—H21*W*···O21^{vi}	0.77 (3)	2.49 (3)	3.050 (2)	131 (3)
O2*W*—H22*W*···O32^{iv}	0.91 (4)	2.04 (5)	2.882 (2)	153 (4)
C11—H11···O21	0.93	2.39	2.846 (2)	110

Symmetry codes: (iv) *x*, −*y*+1/2, *z*−1/2; (v) −*x*, −*y*, −*z*; (vi) *x*, −*y*+3/2, *z*−1/2.