

Acta Crystallographica Section C

Crystal Structure
Communications

ISSN 0108-2701

Pseudopolymorphism in brucine:
brucine–water (1/2), the third crystal
hydrate of brucineGraham Smith,^{a*} Urs D. Wermuth^b and Jonathan M.
White^c^aSchool of Physical and Chemical Sciences, Queensland University of Technology,
GPO Box 2434, Brisbane, Queensland 4001, Australia, ^bSchool of Biomolecular and
Physical Sciences, Griffith University, Nathan, Queensland 4111, Australia, and
^cBIO-21 Molecular Science and Biotechnology, University of Melbourne, Parkville,
Victoria 3052, Australia

Correspondence e-mail: g.smith@qut.edu.au

Received 20 May 2007

Accepted 2 July 2007

Online 26 July 2007

The structure of a third pseudopolymorphic hydrate of
brucine, brucine–water (1/2) [systematic name: 2,3-dimethoxy-
strychnidin-10-one–water (1/2)], C₂₃H₂₆N₂O₄·2H₂O, has beendetermined at 130 K. The asymmetric unit comprises two
independent brucine molecules and four water molecules of
solvation. The four water molecules form uncommon cyclic
hydrogen-bonded homomolecular R_s⁴(8) tetramer rings, which
then form primary hydrogen-bonded chain substructures
extending down the 2₁ screw axis in the unit cell. The two
brucine molecules are linked peripherally to these substructures
by either single O–H···N_{brucine} or asymmetric three-
centre O–H···O_{brucine} hydrogen bonds.

Comment

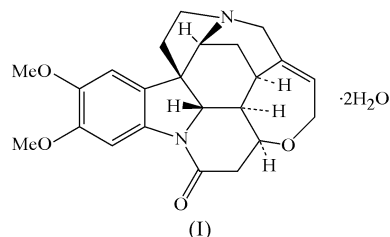
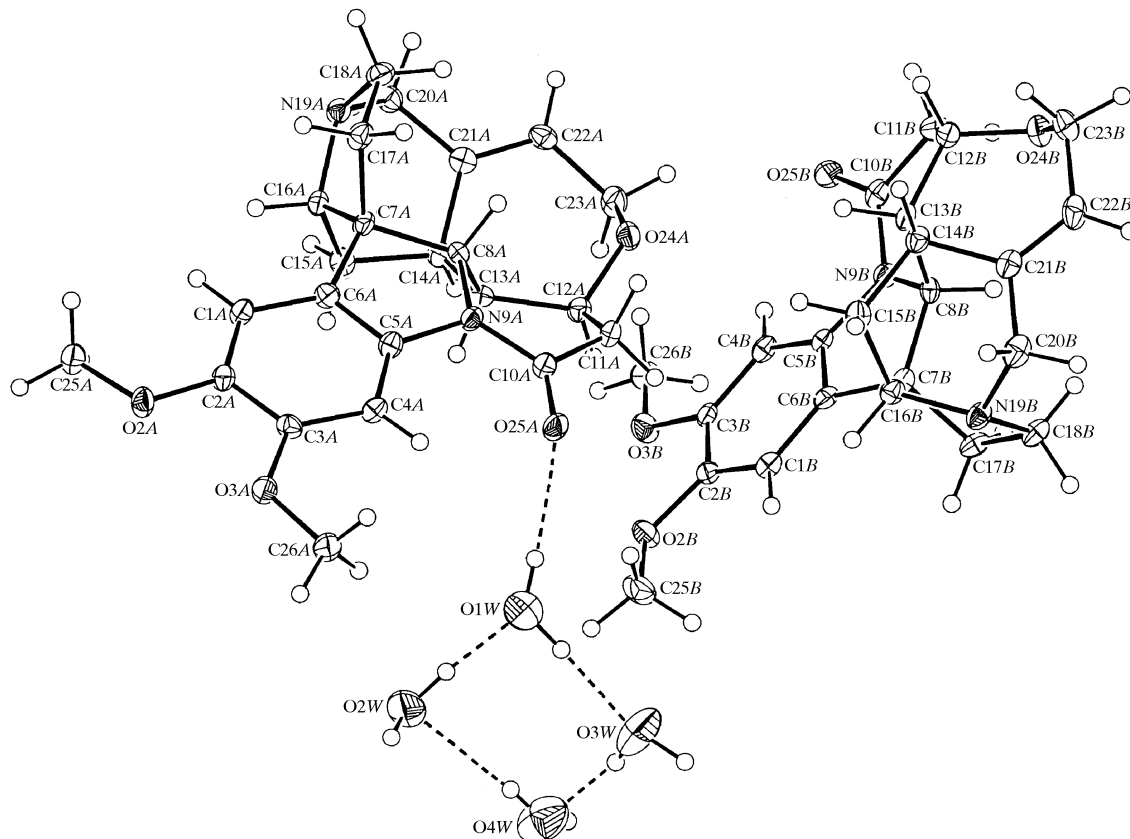
The common crystalline form of the alkaloid brucine is a
tetrahydrate [brucine–water (1/4)], and we have completedthe structure determination of this and another pseudopoly-
morphic hydrate, brucine–water (1/5.25), obtained from the
attempted preparation of a brucine adduct with urea (Smith *et al.*,
2006a). Other crystallographically characterized brucine

Figure 1

The molecular configuration and atom-numbering scheme for the two brucine molecules (A and B) and the four water molecules of solvation in the
asymmetric unit in (I). The hydrogen bonding in the cyclic R_s⁴(8) tetrameric water units and the intra-unit hydrogen bonds are shown as dashed lines.
Non-H atoms are shown as 40% probability displacement ellipsoids.

pseudopolymorphs include anhydrous brucine, brucine–acetone (1/1) and brucine–2-propanol–water (1/1/2) (Białońska & Ciunik, 2004a), and brucine–ethanol–water (1/1/2) (Glover *et al.*, 1985). The 2-propanol–water and ethanol–water solvates are isomorphous.

In a number of these solvate structures, as well as in the proton-transfer compounds of brucine, it has been recognized (Gould & Walkinshaw, 1984; Białońska & Ciunik, 2004b; Smith *et al.*, 2006a,b) that the brucine species form regular undulating parallel or antiparallel host sheet substructures, accommodating the guest molecules in the interstitial cavities. Thus, molecules of solvation are similarly incorporated and associated through hydrogen bonding. In the orthorhombic $P2_12_12_1$ or monoclinic $P2_1$ examples, the presence of a *ca* 12.5 Å unit-cell repeat along a crystallographic 2_1 screw axis was reasonably indicative of this characteristic substructure, *e.g.* in brucine–2-propanol–water (1/1/2) (12.37 Å; Białońska & Ciunik, 2004a) and brucine–ethanol–water (1/1/2) (12.34 Å; Glover *et al.*, 1985) (both $P2_12_12_1$), or in the proton-transfer example brucinium D-glucuronate (12.7 Å, $P2_1$; Dijkstra *et al.*, 1998). In examples where the *ca* 12.5 Å/ $P2_1$ cell parameter/space group combination is not found, the characteristic structuring is usually absent, *e.g.* brucine–acetone (1/1) (7.14 Å, $P2_1$) in the present set of brucine solvates (see Table 2); brucine–water (1/4) (11.53 Å, $P2_12_12_1$), in which the structure is present, is the exception. On the basis of this generalization, the structure of the monoclinic dihydrate, *viz.* brucine–water (1/2), (I), obtained from the attempted

preparation of a brucine–diethanolamine adduct in 95% ethanol, was not expected to have the structuring (with a *b* cell length of *ca* 7.45 Å; $P2_1$). This type of structuring was in fact not found, distinguishing the structure from those of brucine–water (1/4) and brucine–water (1/5.25) (Smith *et al.*, 2006a).

In (I), the asymmetric unit comprises two independent brucine molecules and four water molecules of solvation. The two brucine molecules (*A* and *B*; Fig. 1) have the overall Cahn–Ingold–Prelog absolute configuration [C7(*R*), C8(*S*), C12(*S*), C13(*R*), C14(*R*), C16(*S*)], as found in strychnine (Peerdeman, 1956). Because of the rigid nature of the brucine molecular cage, both molecules, as expected, are conformationally identical, including the methoxy substituents at C2 and C3, which in all brucine structures lie essentially in the plane of the benzene ring. In the efflorescent brucine–water (1/5.25) structure, the asymmetric unit also comprises two independent brucine molecules together with 10.5 molecules of solvent water, some of these having split-occupancy sites. However, unlike this structure, in which one set of brucine molecules forms the common undulating sheet substructure, in (I) there is no such structuring. Instead, the water molecules dominate the structure assembly, forming uncommon cyclic hydrogen-bonded tetramer units (graph set $R_4^4(8)$; Etter *et al.*, 1990; Bernstein *et al.*, 1995) (Figs. 2 and 3). These are analogous to the cyclic water pentamer units found in the structure of brucinium L-glycerate 4.75-hydrate (Białońska *et al.*, 2005). In (I), O–H...O–H...O associations (O4W–H42W...O3W^{vii}; symmetry code as in Table 1) link the tetramers into chain structures which form down the 2_1 screw axis in the unit cell (Fig. 2).

The brucine molecules are linked peripherally to these water structures by hydrogen bonds; there is a single linear interaction with a nitrogen acceptor of a *B* molecule (O2W–H22W...N19Bⁱ), as well as an asymmetric three-centred interaction with the carbonyl O-atom acceptor of an *A* molecule [O2W–H12W...O25A and O3W^{vii}–H32W...O25A; symmetry code: (vii) $-x + 1, y + \frac{1}{2}, -z$]. The second carbonyl O

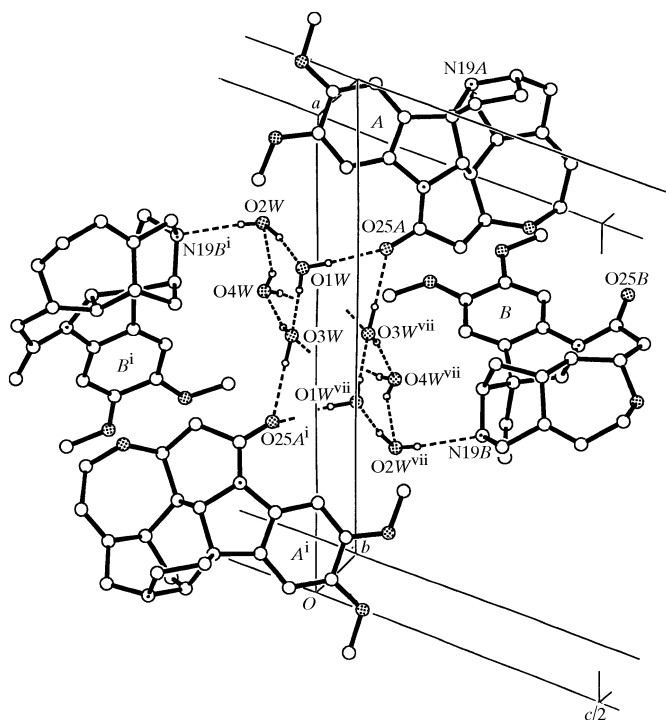


Figure 2

A perspective view of the intermolecular hydrogen bonding in the water chain structure of (I), extending approximately down the *b* axial direction. Non-associative H atoms have been omitted. [Symmetry code: (vii) $-x + 1, y + \frac{1}{2}, -z$; for other symmetry codes, see Table 1.]

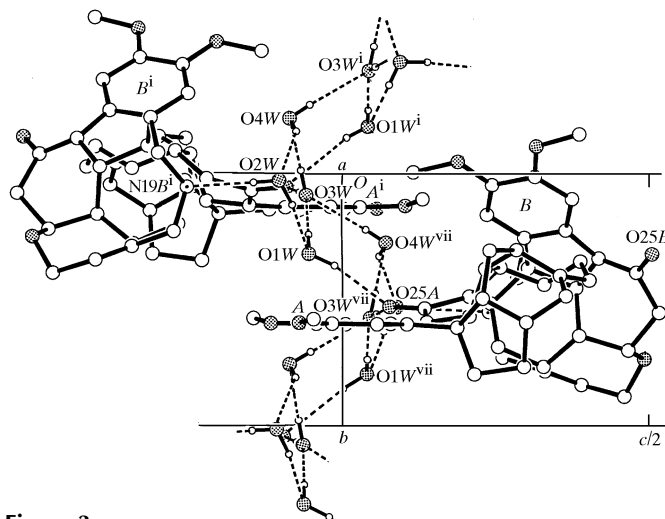


Figure 3

The infinite hydrogen-bonded water chain extension, viewed perpendicular to the (vertical) 2_1 screw axial direction.

atom of molecule *B* (O25*B*) and atom N19*A* of molecule *A* are unassociated except for some weak C—H···O and C—H···N interactions. Unlike the structures of both of the other brucine hydrates (Smith *et al.*, 2006*a*), there are no water–methoxy O—H···O interactions in (I).

It can be assumed that formation of different pseudopolymorphic solvates of brucine depends not only upon solvent composition (Bernstein, 1987; Kumar *et al.*, 1999) but, in the case of the pure hydrates, also upon the presence of additional non-incorporated but structure-influencing solute components, *e.g.* diethanolamine in the case of (I) and urea in brucine–water (1/5.25).

Experimental

Brucine dihydrate, (I), was obtained from the attempted preparation of a 1:1 brucine–diethanolamine adduct by refluxing 0.1 mmol quantities of brucine tetrahydrate and diethanolamine in 40 ml of 50% ethanol–water for 10 min. Colourless prismatic crystals were obtained after total room-temperature evaporation of the solvent.

Crystal data

| | |
|--|--------------------------------------|
| C ₂₃ H ₂₆ N ₂ O ₄ ·2H ₂ O | <i>V</i> = 2064.8 (5) Å ³ |
| <i>M_r</i> = 430.49 | <i>Z</i> = 4 |
| Monoclinic, <i>P</i> 2 ₁ | Mo <i>K</i> α radiation |
| <i>a</i> = 15.178 (2) Å | <i>μ</i> = 0.10 mm ⁻¹ |
| <i>b</i> = 7.4496 (12) Å | <i>T</i> = 130 (2) K |
| <i>c</i> = 19.751 (3) Å | 0.50 × 0.15 × 0.10 mm |
| <i>β</i> = 112.397 (3)° | |

Data collection

| | |
|---------------------------|---|
| Bruker SMART CCD detector | 3936 independent reflections |
| diffractometer | 3072 reflections with <i>I</i> > 2σ(<i>I</i>) |
| 9011 measured reflections | <i>R</i> _{int} = 0.064 |

Refinement

| | |
|---|---|
| <i>R</i> [<i>F</i> ² > 2σ(<i>F</i> ²)] = 0.048 | 1 restraint |
| <i>wR</i> (<i>F</i> ²) = 0.090 | H atoms parameters constrained |
| <i>S</i> = 0.87 | Δ <i>ρ</i> _{max} = 0.20 e Å ⁻³ |
| 3936 reflections | Δ <i>ρ</i> _{min} = -0.21 e Å ⁻³ |
| 559 parameters | |

H atoms potentially involved in hydrogen-bonding interactions were generally located by difference Fourier methods. Some of the H atoms of the water molecules could not be located and were included in the refinement at calculated sites dictated by the assumed hydrogen-bonding geometry, and their positional and isotropic displacement parameters were refined. However, because of the poor reflection–parameter ratio, these were fixed in the final refinement cycles. Brucine H atoms were included at calculated positions (aromatic C—H = 0.95 Å and aliphatic C—H = 0.98–1.00 Å) and treated as riding, with *U*_{iso}(H) values of 1.2*U*_{eq}(C). The absolute configuration determined for the parent strychnine (Peerdeman, 1956) was invoked. Friedel pairs were averaged for the data used in the refinement.

Data collection: *SMART* (Bruker, 2000); cell refinement: *SAINTE* (Bruker, 1999); data reduction: *SAINTE*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics:

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

| <i>D</i> —H··· <i>A</i> | <i>D</i> —H | H··· <i>A</i> | <i>D</i> ··· <i>A</i> | <i>D</i> —H··· <i>A</i> |
|---|-------------|---------------|-----------------------|-------------------------|
| O1 <i>W</i> —H11 <i>W</i> ···O3 <i>W</i> | 0.93 | 1.86 | 2.785 (6) | 170 |
| O1 <i>W</i> —H12 <i>W</i> ···O25 <i>A</i> | 0.87 | 2.11 | 2.960 (4) | 168 |
| O2 <i>W</i> —H21 <i>W</i> ···O1 <i>W</i> | 0.91 | 1.90 | 2.815 (5) | 175 |
| O2 <i>W</i> —H22 <i>W</i> ···N19 <i>B</i> ⁱ | 0.77 | 2.02 | 2.781 (4) | 170 |
| O3 <i>W</i> —H31 <i>W</i> ···O4 <i>W</i> | 0.81 | 2.00 | 2.785 (7) | 164 |
| O3 <i>W</i> —H32 <i>W</i> ···O25 <i>A</i> ⁱ | 0.89 | 1.90 | 2.794 (4) | 179 |
| O4 <i>W</i> —H41 <i>W</i> ···O2 <i>W</i> | 0.73 | 2.13 | 2.843 (7) | 164 |
| O4 <i>W</i> —H42 <i>W</i> ···O3 <i>W</i> ⁱ | 0.73 | 2.55 | 3.210 (6) | 151 |
| C4 <i>A</i> —H4 <i>A</i> ···O25 <i>A</i> | 0.95 | 2.39 | 2.930 (4) | 115 |
| C4 <i>B</i> —H4 <i>B</i> ···O25 <i>B</i> | 0.95 | 2.43 | 2.923 (4) | 112 |
| C22 <i>A</i> —H22 <i>A</i> ···O25 <i>B</i> ⁱⁱ | 0.95 | 2.57 | 3.327 (4) | 137 |
| C26 <i>B</i> —H30 <i>B</i> ···O24 <i>A</i> ⁱⁱⁱ | 0.98 | 2.56 | 3.107 (4) | 115 |
| C17 <i>A</i> —H31 <i>A</i> ···O3 <i>A</i> ^{iv} | 0.99 | 2.44 | 3.324 (4) | 149 |
| C20 <i>B</i> —H35 <i>B</i> ···N19 <i>A</i> ^v | 0.99 | 2.58 | 3.525 (4) | 159 |
| C15 <i>A</i> —H38 <i>A</i> ···O3 <i>A</i> ^{vi} | 0.99 | 2.44 | 3.164 (4) | 129 |

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

Table 2
Comparative unit-cell data for the pseudopolymorphic brucine solvates.

| Cell parameters | Brucine | Brucine·2H ₂ O | Brucine·4H ₂ O | Brucine·5.25H ₂ O | Brucine·EtOH·2H ₂ O | Brucine· ⁱ PrOH·2H ₂ O | Brucine–acetone |
|----------------------------|-------------------------|---------------------------|---|------------------------------|---|---|-------------------------|
| <i>a</i> (Å) | 7.992 (2) | 15.178 (2) | 7.555 (2) | 23.351 (5) | 7.723 (1) | 7.9297 (3) | 12.765 (3) |
| <i>b</i> (Å) | 12.704 (3) | 7.4496 (12) | 11.531 (3) | 12.200 (3) | 12.337 (1) | 12.3289 (7) | 7.1360 (14) |
| <i>c</i> (Å) | 9.471 (2) | 19.751 (3) | 26.492 (8) | 16.972 (4) | 25.212 (2) | 25.1631 (10) | 13.686 (3) |
| α (°) | 90 | 90 | 90 | 90 | 90 | 90 | 90 |
| β (°) | 99.68 (3) | 112.397 (3) | 90 | 96.202 (4) | 90 | 90 | 114.35 (3) |
| γ (°) | 90 | 90 | 90 | 90 | 90 | 90 | 90 |
| <i>V</i> (Å ³) | 947.9 (4) | 2064.7 (6) | 2307.9 (11) | 4806.7 (19) | 2403 (1) | 2460.06 (19) | 1135.8 (4) |
| <i>Z</i> | 2 | 4 | 4 | 8 | 4 | 4 | 2 |
| Space group | <i>P</i> 2 ₁ | <i>P</i> 2 ₁ | <i>P</i> 2 ₁ 2 ₁ 2 ₁ | <i>C</i> 2 | <i>P</i> 2 ₁ 2 ₁ 2 ₁ | <i>P</i> 2 ₁ 2 ₁ 2 ₁ | <i>P</i> 2 ₁ |
| Reference | <i>a</i> | <i>b</i> | <i>c</i> | <i>c</i> | <i>d</i> | <i>a</i> | <i>a</i> |

Notes: (*a*) Białońska & Ciunik (2004*a*); (*b*) this work; (*c*) Smith *et al.* (2006*a*); (*d*) Glover *et al.* (1985).

PLATON (Spek, 2003); software used to prepare material for publication: PLATON.

The authors acknowledge financial support from the School of Physical and Chemical Sciences (Queensland University of Technology), the School of Biomolecular and Physical Sciences (Griffith University) and the School of Chemistry (University of Melbourne).

Supplementary data for this paper are available from the IUCr electronic archives (Reference: GA3057). Services for accessing these data are described at the back of the journal.

References

- Bernstein, J. (1987). *Organic Solid State Chemistry*, Vol. 32, edited by G. R. Desiraju, pp. 471–518. Amsterdam: Elsevier.
- Bernstein, J., Davis, R. E., Shimoni, L. & Chang, N.-L. (1995). *Angew. Chem. Int. Ed. Engl.* **34**, 1555–1573.
- Bialońska, A. & Ciunik, Z. (2004a). *Acta Cryst.* **C60**, o853–o855.
- Bialońska, A. & Ciunik, Z. (2004b). *CrystEngComm*, **6**, 276–279.
- Bialońska, A., Ciunik, Z., Popek, T. & Lis, T. (2005). *Acta Cryst.* **C61**, o88–o91.
- Bruker (1999). *SAINT*. Version 6.02. Bruker AXS Inc., Madison, Wisconsin, USA.
- Bruker (2000). *SMART*. Version 5.55. Bruker AXS Inc., Madison, Wisconsin, USA.
- Dijksma, F. J. J., Gould, R. O., Parsons, S., Taylor, J. & Walkinshaw, M. D. (1998). *Chem. Commun.* pp. 745–746.
- Etter, M. C., MacDonald, J. C. & Bernstein, J. (1990). *Acta Cryst.* **B46**, 256–262.
- Glover, S. S. B., Gould, R. O. & Walkinshaw, M. D. (1985). *Acta Cryst.* **C41**, 990–994.
- Gould, R. O. & Walkinshaw, M. D. (1984). *J. Am. Chem. Soc.* **106**, 7840–7842.
- Kumar, V. S. S., Kuduva, S. S. & Desiraju, G. R. (1999). *J. Chem. Soc. Perkin Trans. 2*, pp. 1069–1073.
- Peerdeman, A. F. (1956). *Acta Cryst.* **9**, 824.
- Sheldrick, G. M. (1997). *SHELXL97* and *SHELXS97*. University of Göttingen, Germany.
- Smith, G., Wermuth, U. D., Healy, P. C. & White, J. M. (2006a). *Acta Cryst.* **C62**, o203–o207.
- Smith, G., Wermuth, U. D., Healy, P. C. & White, J. M. (2006b). *Aust. J. Chem.* **59**, 320–328.
- Spek, A. L. (2003). *J. Appl. Cryst.* **36**, 7–13.