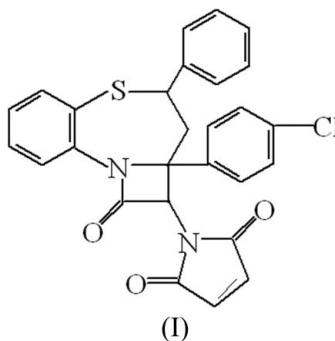


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Key indicators

Single-crystal X-ray study
T = 293 K
Mean $\sigma(\text{C}-\text{C}) = 0.004 \text{ \AA}$
R factor = 0.041
wR factor = 0.108
Data-to-parameter ratio = 13.5For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.2a-(4-Chlorophenyl)-2-(2,5-dioxo-3-pyrazolin-1-yl)-
4-phenyl-2,2a,3,4-tetrahydro-1*H*-azeto[2,1-*d*][1,5]-
benzothiazepin-1-oneThe conformation of the seven-membered ring in the title
molecule, $\text{C}_{27}\text{H}_{19}\text{ClN}_2\text{O}_3\text{S}$, is that of a chair. The 2,5-dioxo-3-
pyrazolin-1-yl and 4-chlorophenyl groups attached to the β -
lactam ring are in *cis* positions.Received 25 July 2006
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Comment

1,5-Benzothiazepines are useful and important compounds
utilized in drug research owing to their well known bioactiv-
ities (Ansari *et al.*, 2005; Dandia *et al.*, 1998). Cycloaddition
reactions of their $\text{C}=\text{N}$ double bond with ketene may produce
2-azetidinone which is the essential component of some anti-
biotics (Gerard *et al.*, 2004). We are interested in the synthesis
of 2-azetidinone derivatives of 1,5-benzothiazepines and we
report here the synthesis and the structure of the title
compound, (I).Compound (I) was obtained by the cycloaddition of 2-*tert*-
butoxy-*N*-glyciny succinimide with 4-(4-chlorophenyl)-2-
phenyl-1,5-benzothiazepine. The C—O bond of the *tert*-
butoxy group is also cleaved and subsequent elimination of H_2O
yielded the double bond, in agreement with previous work
(Ping *et al.*, 2006).A view of the molecule is shown in Fig. 1. The seven-
membered ring in the molecule has a chair conformation,
while in the 1,5-benzothiazepine it has a boat form (Lu *et al.*,
1987). Thus, the conformation of the seven-membered ring
changes in the cycloaddition reaction. The four-membered
ring in the molecule results from the cycloaddition reaction.
The 2,5-dioxo-3-pyrazolin-1-yl and chlorophenyl groups
attached to C22 and C15 are oriented away from the plane of
the four-membered ring and are in *cis* positions.

Experimental

A mixture of 2-*tert*-butoxy-*N*-glyciny succinimide (20 mmol) and
 SOCl_2 (30 ml) was refluxed for 4 h. When the reaction was complete,

SOCl₂ was removed. 4-(4-Chlorophenyl)-2-phenyl-1,5-benzothiazepine (10 mmol), benzene (30 ml) and Et₃N (6 ml) were added to the above mixture and stirred at room temperature for 1 h. After the reaction was complete, the mixture was filtered and the filtrate was washed with 10% HCl; the benzene layer was separated and dried over anhydrous Na₂SO₄. The benzene solution was evaporated and the residue was purified by column chromatography (ethyl acetate/petroleum ether = 1:3) to give the title product. Analysis calculated for C₂₇H₁₉ClN₂O₃S: C 66.54, H 3.90, N 5.75%; found: C 66.49, H 3.97, N 5.70%.

Crystal data

C₂₇H₁₉ClN₂O₃S
M_r = 486.95
 Monoclinic, *P*2₁/*c*
a = 12.3034 (12) Å
b = 16.6438 (16) Å
c = 12.2606 (12) Å
 β = 110.041 (1)°
V = 2358.6 (4) Å³

Z = 4
D_x = 1.371 Mg m⁻³
 Mo *K*α radiation
 μ = 0.28 mm⁻¹
T = 293 (2) K
 Block, colorless
 0.26 × 0.22 × 0.18 mm

Data collection

Bruker SMART CCD area-detector diffractometer
 φ and ω scans
 Absorption correction: multi-scan (*SADABS*; Sheldrick, 1997)
T_{min} = 0.834, *T_{max}* = 0.855
 (expected range = 0.927–0.950)

12674 measured reflections
 4157 independent reflections
 2966 reflections with *I* > 2σ(*I*)
R_{int} = 0.025
 θ_{max} = 25.0°

Refinement

Refinement on *F*²
R [*F*² > 2σ(*F*²)] = 0.041
wR (*F*²) = 0.108
S = 1.03
 4157 reflections
 307 parameters
 H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0334P)^2 + 1.0352P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{max} < 0.001$
 $\Delta\rho_{max} = 0.26 \text{ e \AA}^{-3}$
 $\Delta\rho_{min} = -0.32 \text{ e \AA}^{-3}$

H atoms were placed in calculated positions and constrained to ride on their parent atoms, with C–H = 0.93 or 0.97 Å (aryl H) and *U_{iso}*(H) = 1.2*U_{eq}*(C)

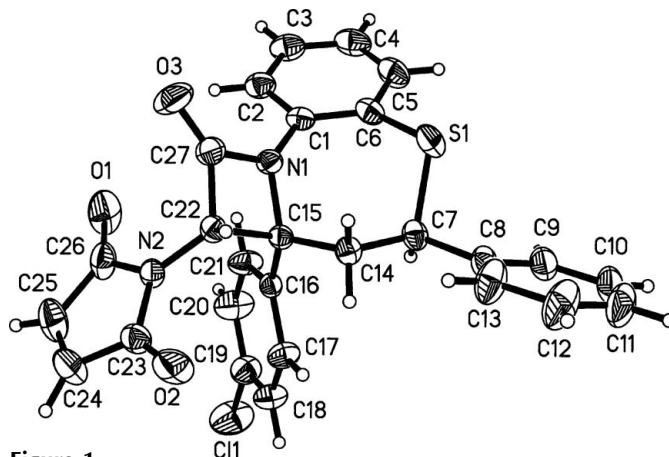


Figure 1 The molecular structure of the title compound, showing 30% probability displacement ellipsoids and the atom-numbering scheme.

Data collection: *SMART* (Bruker, 1997); cell refinement: *S SAINT* (Bruker, 1997); data reduction: *S SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 1997); software used to prepare material for publication: *SHELXTL*.

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