

Tetrakis(trimethylsilyl) isocyanatomethylene-  
bisphosphonatePetar Y. Petrov,<sup>a</sup> Robert  
McDonald,<sup>b\*</sup> Robert D.  
Lukowski,<sup>b</sup> Ronald G. Cavell<sup>b</sup>  
and Christo M. Angelov<sup>b</sup><sup>a</sup>Department of Organic Chemistry, Faculty of  
Chemistry, University of Sofia, 1126 Sofia,  
Bulgaria, and <sup>b</sup>Chemistry Department,  
University of Alberta, Edmonton, Alberta  
T6G 2G2, Canada

\* X-ray Crystallography Laboratory.

Correspondence e-mail:  
bob.mcdonald@ualberta.ca

## Key indicators

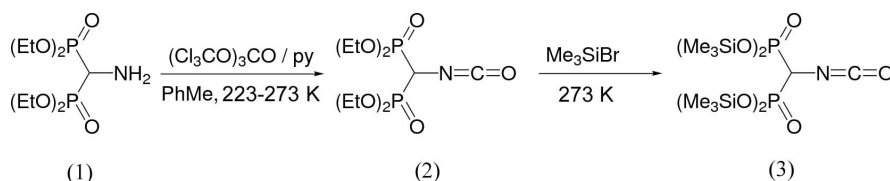
Single-crystal X-ray study  
T = 193 K  
Mean  $\sigma(\text{N}-\text{C}) = 0.003 \text{ \AA}$   
Disorder in main residue  
R factor = 0.035  
wR factor = 0.102  
Data-to-parameter ratio = 21.7For details of how these key indicators were  
automatically derived from the article, see  
<http://journals.iucr.org/e>.Syntheses of tetraethyl isocyanatomethylenediphosphonate  
and the title compound,  $\text{C}_{14}\text{H}_{37}\text{NO}_7\text{P}_2\text{Si}_4$ , are reported. The  
structure of the latter shows disorder of the isocyanate group,  
but bond lengths and angles within this group are normal.

Received 29 March 2005

Accepted 29 April 2005

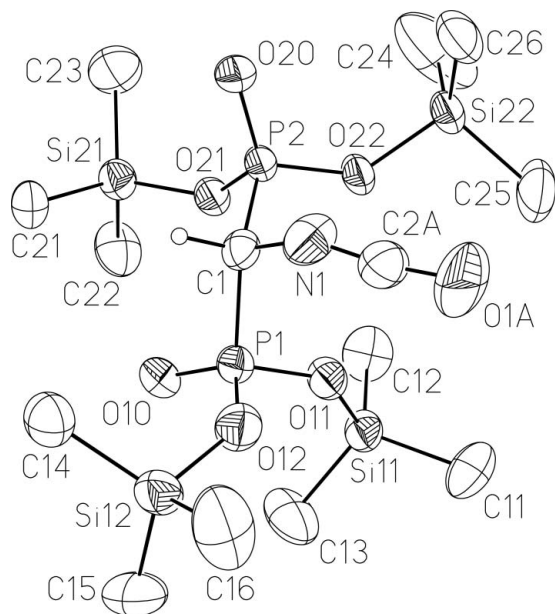
Online 7 May 2005

## Comment

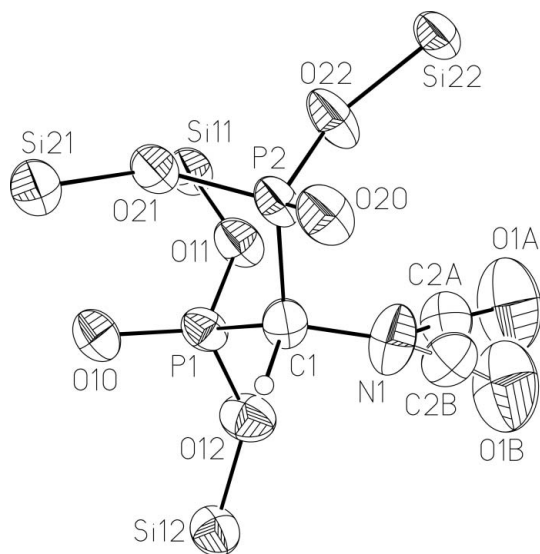
Bisphosphonates are a valuable class of compounds due to  
their ability to participate in calcium metabolism in bones,  
thus opening up possibilities for effective treatment of various  
bone diseases (Body, 1998; Sparidans *et al.*, 1998). Recently,  
bisphosphonates have been used in the treatment of bone  
metastasis diseases (Mundy & Yoneda, 1998; Smith, 2003). A  
relatively new use for bisphosphonic acids has been in the  
development of targeted anticancer drugs (El-Mabhough *et al.*,  
2004), leading us to view compounds bearing the desired  
bisphosphonate moiety and a reactive pre-linker as our  
synthetic targets. We report here an efficient synthetic  
approach, leading to the title compound, (3), as a precursor to  
new drugs.Reaction of tetraethyl aminomethylenediphosphonate, (1),  
with triphosgene affords tetraethyl isocyanatomethylenedi-  
phosphonate, (2), which may be treated with trimethylsilyl  
bromide to produce the title compound, tetrakis(trimethyl-  
silyl) isocyanatomethylenebisphosphonate, (3). The structure  
of (3) is shown in Fig. 1. Although the isocyanate group is  
disordered (Fig. 2), the distances and angles involving the  
disordered components are in satisfactory mutual agreement.  
The isocyanate  $\text{N}=\text{C}$  bond distance values are slightly less  
than the shortest previously observed (1.131 Å) for an  
isocyanate group bound to carbon (Chini *et al.*, 1988), but the  
difference is statistically insignificant (and the aforementioned  
disorder makes the present values less reliable).

## Experimental

For the preparation of tetraethyl isocyanatomethylenediphospho-  
nate, (2), a solution of tetraethyl aminomethylenediphospho-  
nate, (1) (3.0 g, 9.9 mmol), and pyridine (1.57 g, 1.60 ml, 19.8 mmol) in toluene  
(50 ml) was slowly (15 min) added dropwise to a stirred cooled  
(223 K) solution of triphosgene [ $(\text{Cl}_3\text{CO})_2\text{C}=\text{O}$ : 1.2 g, 3.7 mmol] in  
toluene (100 ml). After the addition was complete, the reaction  
mixture was allowed to warm to 273 K and stirred at this temperature



**Figure 1**  
View of (3), with displacement ellipsoids drawn at the 50% probability level. The methine H atom is drawn as a sphere of arbitrary radius, while methyl H atoms are not shown. Atoms O1A and C2A were refined at 0.65 occupancy (O1B and C2B of 0.35 occupancy are not shown).



**Figure 2**  
Illustration of the disorder of the isocyanate group. Methyl C atoms bound to silicon have been omitted.

for an additional 2 h. The precipitate of pyridinium chloride was filtered off under argon and washed with toluene (2 × 15 ml), after which the combined toluene solutions were evaporated under vacuum and the residue distilled (yield: 2.7 g, 8.1 mmol, 82%; b.p. 408–413 K/0.15 mm Hg). For the preparation of tetrakis(trimethylsilyl) isocyanatomethylenebisphosphonate, (3), neat (2) (1.67 g, 5.1 mmol) was reacted with an eightfold excess of Me<sub>3</sub>SiBr (6.23 g, 5.35 ml, 4.07 mmol) at 273 K for 48 h. The excess Me<sub>3</sub>SiBr was evaporated under vacuum and the residue distilled (yield: 2.4 g, 4.7 mmol, 93%; b.p. 393 K/0.05 mm Hg). The slightly yellow liquid solidifies on standing or cooling (m.p. 317–318 K). Single crystals suitable for diffraction measurements were obtained by slowly cooling molten (3).

**Crystal data**

C<sub>14</sub>H<sub>37</sub>NO<sub>7</sub>P<sub>2</sub>Si<sub>4</sub>  
M<sub>r</sub> = 505.75  
Monoclinic, P2<sub>1</sub>/n  
a = 10.2012 (6) Å  
b = 26.7355 (15) Å  
c = 11.0576 (6) Å  
β = 107.6500 (9)°  
V = 2873.8 (3) Å<sup>3</sup>  
Z = 4

D<sub>x</sub> = 1.169 Mg m<sup>-3</sup>  
Mo Kα radiation  
Cell parameters from 6971 reflections  
θ = 2.4–26.4°  
μ = 0.35 mm<sup>-1</sup>  
T = 193 (2) K  
Prism, pale yellow  
0.47 × 0.43 × 0.43 mm

**Data collection**

Bruker SMART 1000 CCD area-detector/PLATFORM diffractometer  
ω scans  
Absorption correction: multi-scan (SADABS; Sheldrick, 2003)  
T<sub>min</sub> = 0.854, T<sub>max</sub> = 0.865  
21 491 measured reflections

5905 independent reflections  
5076 reflections with I > 2σ(I)  
R<sub>int</sub> = 0.022  
θ<sub>max</sub> = 26.4°  
h = -12 → 12  
k = -33 → 33  
l = -13 → 13

**Refinement**

Refinement on F<sup>2</sup>  
R[F<sup>2</sup> > 2σ(F<sup>2</sup>)] = 0.035  
wR(F<sup>2</sup>) = 0.102  
S = 1.03  
5905 reflections  
272 parameters  
H-atom parameters constrained

w = 1/[σ<sup>2</sup>(F<sub>o</sub><sup>2</sup>) + (0.0569P)<sup>2</sup> + 0.9513P]  
where P = (F<sub>o</sub><sup>2</sup> + 2F<sub>c</sub><sup>2</sup>)/3  
(Δ/σ)<sub>max</sub> = 0.001  
Δρ<sub>max</sub> = 0.41 e Å<sup>-3</sup>  
Δρ<sub>min</sub> = -0.24 e Å<sup>-3</sup>

**Table 1**

Selected geometric parameters (Å, °).

P1–O10	1.4596 (14)	Si12–O12	1.6713 (14)
P1–O11	1.5502 (12)	Si21–O21	1.6826 (13)
P1–O12	1.5617 (14)	Si22–O22	1.6730 (12)
P1–C1	1.8214 (17)	O1A–C2A	1.164 (9)
P2–O20	1.4653 (12)	O1B–C2B	1.162 (17)
P2–O21	1.5514 (13)	N1–C1	1.447 (2)
P2–O22	1.5539 (12)	N1–C2A	1.129 (9)
P2–C1	1.8253 (18)	N1–C2B	1.111 (17)
Si11–O11	1.6813 (13)		
O10–P1–O11	116.85 (8)	P1–O11–Si11	136.56 (9)
O10–P1–O12	115.46 (8)	P1–O12–Si12	135.64 (10)
O11–P1–O12	103.12 (8)	P2–O21–Si21	134.84 (8)
O10–P1–C1	113.06 (8)	P2–O22–Si22	136.75 (8)
O11–P1–C1	104.03 (7)	C1–N1–C2A	141.6 (4)
O12–P1–C1	102.63 (8)	C1–N1–C2B	151.4 (9)
O20–P2–O21	115.85 (7)	P1–C1–P2	117.44 (9)
O20–P2–O22	115.37 (7)	P1–C1–N1	111.79 (12)
O21–P2–O22	103.97 (7)	P2–C1–N1	107.95 (13)
O20–P2–C1	111.43 (8)	N1–C2A–O1A	170.3 (9)
O21–P2–C1	104.54 (7)	N1–C2B–O1B	174.8 (12)
O22–P2–C1	104.46 (8)		

Disorder within the isocyanate (–N=C=O) group was handled by splitting the C and O atoms into two sets of positions (C2A/O1A and C2B/O1B), which were given respective occupancy factors of 0.65 and 0.35. Occupancy factors were chosen that allowed the U<sub>eq</sub> values for corresponding atoms (O1A/O1B and C2A/C2B) to refine to comparable values, and produced roughly equivalent N–C and C–O distances in the disordered fragments. H atoms were placed in idealized positions according to the sp<sup>3</sup> geometries about their parent C atoms, at C–H distances of 0.98 (methine) or 1.00 Å (methyl), and with U<sub>iso</sub>(H) = 1.2U<sub>eq</sub>(C).

Data collection: SMART (Bruker, 2001); cell refinement: SAINT (Bruker, 2003); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics:

*SHELXTL* (Bruker, 2003); software used to prepare material for publication: *SHELXTL*.

We are indebted to the University of Alberta for support of this work. The Bruker SMART 1000 CCD area detector/PLATFORM diffractometer was purchased with the generous support of the Natural Sciences and Engineering Research Council (NSERC) of Canada.

## References

- Body, J.-J. (1998). *Eur. J. Cancer*, **34**, 263–269.
- Bruker (2001). *SMART*. Version 5.054. Bruker AXS Inc., Madison, Wisconsin, USA.
- Bruker (2003). *SAINT* (Version 6.45A) and *SHELXTL* (Version 6.14). Bruker AXS Inc., Madison, Wisconsin, USA.
- Chini, M., Crotti, P., Macchia, F., Domiano, P. & Monti, L. (1988). *Gazz. Chim. Ital.* **118**, 123–125.
- El-Mabhouh, A., Angelov, C., McEwan, A., Jia, G. F. & Mercer, J. (2004). *Cancer Biother. Radiopharm.* **19**, 627–640.
- Mundy, G. R. & Yoneda, T. (1998). *New England J. Med.* **339**, 398–400.
- Sheldrick, G. M. (1997). *SHELXS97* and *SHELXL97*. University of Göttingen, Germany.
- Sheldrick, G. M. (2003). *SADABS*. Version 2.10. University of Göttingen, Germany.
- Smith, M. R. (2003). *Clin. Cancer Res.* **9**, 5433–5434.
- Sparidans, R. W., Twiss, I. M. & Talbot, S. (1998). *Pharm. World Sci.* **20**, 206–213.