- O'GORMAN, J. M., SHAND, W. & SCHOMAKER, V. (1950). J. Amer. Chem. Soc. 73, 4222.
- PAULING, L. (1952). J. Phys. Chem. 56, 341.
- PAULING, L. & COREY, R. B. (1953). Proc. Nat. Acad. Sci., Wash. 39, 84.
- ROLLETT, J. S. & DAVIES, D. R. (1955). Acta Cryst. 8, 125.
- ROMERS, C., KETELAAR, J. A. A. & MACGILLAVRY, C. M. (1949). Nature, Lond. 164, 960.
- SMITH, J. P., BROWN, W. E. & LEHR, J. R. (1955). J. Amer. Chem. Soc. 77, 2728.
- SPARKS, R. A., PROSEN, R. J., KRUSE, F. H. & TRUE-BLOOD, K. N. (1956). Acta Cryst. 9, 350.
- WEST, J. (1930). Z. Kristallogr. 74, 306.

Acta Cryst. (1956). 9, 334

# A Refinement of the Crystal Structure of Succinamide\*

By DAVID R. DAVIEST AND R. A. PASTERNAK

Gates and Crellin Laboratories of Chemistry, California Institute of Technology, Pasadena, California, U.S.A.

(Received 6 September 1955)

The structure of succinamide has been refined to a high degree of precision by three-dimensional Fourier and least-squares methods. In this structure the molecules are planar and are tied into layers by a net of hydrogen bonds. The bond lengths in the amide group are C-N = 1.333 and C-O = 1.238 Å; the bond angles are  $C-C-N = 115^{\circ} 36'$ ,  $C-C-O = 122^{\circ} 23'$ , and  $N-C-O = 122^{\circ} 2'$ . The two non-equivalent C-C bonds have lengths of 1.512 and 1.501 Å, the central bond being slightly but significantly longer.

#### Introduction

When the crystal-structure determination of succinamide was initiated, the dimensions of the terminal amide group had been determined in only three compounds: acetamide (Senti & Harker, 1940), glutamine (Cochran & Penfold, 1952) and glycylasparagine (Pasternak, Katz & Corey, 1954). Because the bond lengths and angles reported for this group in these three investigations differed significantly, a precise determination of the succinamide structure, which is simpler than any of the other three, was considered to be of value. Recently, the crystal structures of formamide (Ladell & Post, 1954) and oxamide (Ayerst & Duke, 1954) have also been published.

### Experimental

### Unit cell and space group

Crystals of succinamide grown from water solutions were monoclinic plates bounded by  $\{001\}$  faces and showing perfect cleavage along  $\{100\}$ . Unit-cell dimensions were derived from single-crystal rotation photographs obtained with a Straumanis-type camera; they are, with their limits of error,

$$a_0 = 6.932 \pm 0.003, \ b = 7.994 \pm 0.003, \ c = 9.878 \pm 0.004 \text{ Å}, \ \beta = 102^{\circ} 28' + 5'.$$

The density observed by flotation is  $1.442\pm0.005$  g.cm.<sup>-3</sup>, and the number of molecules per unit cell is 4 (calculated 3.998).

The observed systematic absences are hkl when h+k is odd and h0l when h or l are odd; they are characteristic of the space groups Cc (4 general positions) and C2/c (centrosymmetric, 8 general positions).

Trial structure

A trial structure for succinamide



has been reported previously (Pasternak, 1953). The observations of (a) perfect cleavage of the crystals along {100}, (b) outstandingly strong (h00) reflections showing normal decline, (c) extremely weak (hk0) reflections when h and k are odd, (d) a very similar intensity distribution on layer lines with h even and with h odd, respectively, pointed to an almost perfect layer structure parallel to (100) with the layers of molecules at  $x = \frac{1}{4}$  and  $\frac{3}{4}$ . By taking into account known bond lengths and bond angles, van der Waals distances, and the expectation of hydrogen bond formation, the configuration of the molecules and their

<sup>\*</sup> Contribution No. 2034 from the Gates and Crellin Laboratories. This investigation was supported by a research grant RG 1987, from the National Institutes of Health, Public Health Service.

<sup>†</sup> Present address: National Institute of Mental Health, Bethesda 14, Maryland, U.S.A.

arrangement within a layer was easily derived. The arrangement was consistent with the space group C2/c with four centrosymmetric molecules in the unit cell.

## Intensity data

Because of the shortness of the a axis and the apparent arrangement of the molecules in layers parallel to (100), Weissenberg photographs around a were most desirable. The crystals, however, were plates along (001) and could not be shaped into cylinders, as they cleaved almost on touch. An attempt was therefore made to modify the crystal habit by the addition of methylene blue to a hot concentrated aqueous solution of succinamide. This method proved to be successful; small laths or needles with a as the needle axis were obtained. A comparison between Weissenberg photographs taken with the modified and unmodified crystals showed no detectable differences in intensities, and a needle was therefore used for the intensity work; the specimen chosen had an almost square cross-section of about 0.2 mm. on edge. Equi-inclination Weissenberg photographs of the zero through fifth layer lines in respect to h were taken with Cu  $K\alpha$  radiation, and of the zero through eighth layer lines with Mo  $K\alpha$  radiation; the eighth layer line was very weak. In addition, the (h0l), (h1l), and (h2l) reflections were recorded with Cu radiation from a crystal oscillating around the b axis; they were used for the determination of an average anisotropic temperature factor along a (see below).

Intensities were estimated visually from multiple films, the range of intensities being 1 to 10,000. They were corrected in the usual way for Lorentz and polarization factors; no absorption or extinction corrections were applied. Factors correlating Cu to Mo data were derived by comparing the corrected intensities of a number of medium strong reflections common to both. Since the quality of the intensity data obtained from Cu radiation was considered superior to that obtained from Mo radiation, the Mo data were used only for those reflections lying outside the Cu sphere. A few reflections with noticeable intensities appeared even at spacings smaller than 0.5 Å, but this value was taken as limit of observation.

The data were brought to an absolute scale at a later stage by comparing, separately for each layer, the observed and calculated structure factors. The introduction of these additional parameters, the scale factors, was considered legitimate in view of the great number—about 1100—of observed structure factors.

### The refinement of the structure

On the basis of the y and z parameters of the trial structure, and taking  $x = \frac{1}{4}$  for all atoms, structure factors were calculated for all (0kl) and (1kl) reflections lying within the sphere of reflection for Cu radiation; no temperature factor was applied at this

stage. The agreement between calculated and observed structure factors was encouraging, the discrepancy factor R being 0.27.

The refinement of the y and z parameters progressed rapidly by the use of Fourier and difference Fourier projections and by least-squares treatment of the combined (0kl) and (1kl) data. When the reliability factor had been reduced to 0.17, after the contribution of the hydrogen atoms to the scattering of the structure and a temperature factor of B = 0.85 Å<sup>2</sup> had been introduced, three-dimensional refinement of the structure was begun. Structure factors for all reflections of the zero through fifth layer lines with  $\sin \theta_{MO}$ smaller than 0.6 were calculated, disregarding the obviously large temperature anisotropy along a, but the approximate scale factors which were obtained for each layer individually included a temperature-factor contribution. Three least-squares refinements of all the positional parameters were carried out; in addition to shifts in y and z, small, but significant, displacements of the atoms from the plane  $x = \frac{1}{4}$  were obtained.

Although R had dropped to below 0.15, systematic discrepancies were apparent; in general, the (hkl), structure factors calculated stronger and the (hkl)structure factors weaker than the observed ones. These discrepancies clearly originated from a large temperature anisotropy along a. The anisotropy had been taken into account only in a crude way, namely, as a constant for each layer line; this is not quite correct for a monoclinic crystal. For a monoclinic space group with an average anisotropic temperature factor T the structure factor has the form  $F_c = F_g T$ , where

$$T = \exp\left[-\frac{1}{4}(\alpha a^{*2}h^2 + \beta b^{*2}k^2 + \gamma c^{*2}l^2 + \delta a^*c^*hl)\right].$$
 (1)

Here,  $F_g$  is the geometric part of the structure factor,  $\alpha$ ,  $\beta$  and  $\gamma$  are the average anisotropic temperaturefactor coefficients along the reciprocal axes, and  $\delta$  is a coefficient related to the orientation of the ellipsoid of anisotropy with respect to  $a^*$  and  $c^*$ . These coefficients are given in the same units as the isotropic temperature factor *B*. For symmetry reasons one axis of the ellipsoid must coincide with  $b^*$ .

The temperature factors of a pair of reflections the indexes of which differ only by the sign of l are not identical, because of the term  $\exp\left[-\frac{1}{4}\delta a^*c^*hl\right]$ . However, by taking ratios one obtains the simple relation

$$\frac{F(hkl)_o}{F(hk\bar{l})_o} \approx \frac{F(hkl)_c}{F(hk\bar{l})_c} = \frac{F(hkl)_g}{F(hk\bar{l})_g} \exp\left[-\frac{1}{2}\delta a^* c^* hl\right], \quad (2)$$

which contains  $\delta$  as single variable, assuming that the positional parameters have converged and that  $\delta$  is independent of the scale factor. This expression is therefore very convenient for the determination of  $\delta$  by least-squares methods. Such a least-squares calculation was carried out for the layer lines h = 1 to h = 6 individually and gave an average value of

 $\delta = 2.1$ , the average deviations of the individual values from the mean being 8%.

In the present structure the coefficient  $\alpha$  can be obtained directly from  $\delta$  by making the reasonable assumption that one principal axis of temperature anisotropy is perpendicular to the molecular layer, i.e. along  $a^*$ ; the other axes of anisotropy are then parallel to  $b^*$  and c. On transforming the temperature-factor expression (1) to orthogonal axes parallel to these principal axes of temperature anisotropy, the temperature factor becomes (see also Rollett & Davies, 1955)

$$T = \exp\left[-\frac{1}{4}\left\{\alpha' a^{*2}(h + (c^{*}/a^{*})\cos\beta^{*}l)^{2} + \beta' b^{*2}k^{2} + \gamma' \frac{c^{*2}}{\sin^{2}\beta^{*}}l^{2}\right\}\right].$$
 (3)

By comparing equations (1) and (3) the relationships between the coefficients  $\alpha$ ,  $\beta$ ,  $\gamma$ ,  $\delta$  and  $\alpha'$ ,  $\beta'$ ,  $\gamma'$ , respectively, are easily derived; thus,

$$\alpha = \alpha', \ \beta = \beta', \ \gamma = \frac{\gamma'}{\sin^2 \beta^*} + \alpha' \cos^2 \beta^*, \ \delta = 2 \cos \beta^* \alpha'.$$

Inserting the known values of  $\delta$  and  $\beta^*$  into the last equation, the value  $\alpha' = 4.9$  was obtained. A direct determination of  $\alpha$  from the (hk0), (hk1), and (hk2)data, taking  $\beta$  equal to  $\gamma$  and identical with the average in-plane temperature factor B = 0.85 Å<sup>2</sup>, produced very nearly the same value,  $\alpha' = 5.2$ ; it confirmed the assumption that the main axis of temperature anisotropy is perpendicular to the layer of the molecule.

All structure factors were now recalculated using the complete temperature-factor expression, and the scale factors were redetermined; the agreement between observed and calculated structure factors improved significantly. A three-dimensional difference Fourier, in which the contribution of hydrogen atoms to the calculated F's was omitted, was then calculated in order to determine the individual atomic anisotropic temperature factors and to locate experimentally the hydrogen atoms. Peaks of heights approximately 1 e.Å<sup>-3</sup> were indeed found near the positions where hydrogen atoms were expected, namely, two in the plane of the molecule at a distance of 0.9 Å from the nitrogen atom and two symmetrically above and below the plane at a distance of 1.0 Å from carbon atom C<sub>1</sub>.

Sizeable differences in the magnitudes of the temperature anisotropies for the various atoms were apparent on the difference map; fortunately, however, the directions of the axes of the ellipsoid of anisotropy were for all atoms approximately parallel to  $a^*$ ,  $b^*$ and c. The temperature factor for each atom was therefore represented by equation (3),  $\alpha'$ ,  $\beta'$  and  $\gamma'$ being now the individual atomic temperature-factor coefficients. The magnitudes of these coefficients were determined by least-squares treatment of the complete data. Three refinements similar to those described by Davies & Blum (1955) were carried out, interspersed with adjustment of the scale factors of the layer lines. In a final step, the positional parameters were also adjusted by least squares. The shifts in the parameters were barely significant, the maximum being 0.004 Å.

It is interesting to note that in the refinement of the structure of succinamide the large atomic temperature anisotropies virtually did not interfere with the determination of the positional parameters. This may be due to a special feature of the structure; namely, that the large temperature vibration is along a direction in which atoms are very widely spaced, and therefore there is little interaction between vibrational and positional parameters.

The final positional and vibrational parameters of the atoms are listed in Table 1. The average standard

### Table 1. Atomic coordinates

	$\boldsymbol{x}$	$\boldsymbol{y}$	z	$\alpha'$ (Å <sup>2</sup> )	$eta^\prime$ (Ų)	γ' (Ų)
$C_1$	0.2496	-0.1943	0.0611	4.50	0.95	1.28
$\overline{C_2}$	0.2537	-0.0097	0.0285	3.85	0.82	1.22
N	0.2532	0.0933	0.1346	5.86	1.04	1.30
0	0.2559	0.0429	-0.0890	6.37	1.01	1.10
$\mathbf{H}_{1}$	0.125	-0.225	0.110			
$\mathbf{H}_{2}$	0.375	-0.225	0.110			
$H_3$	0.250	0.503	0.117			
$\mathbf{H}_{\mathbf{A}}$	0.250	0.055	0.220			

error in the positional parameters was calculated as  $2 \cdot 2 \times 10^{-3}$  Å in x and  $1 \cdot 4 \times 10^{-3}$  Å in y and z; the standard error in the bond lengths is then  $2 \times 10^{-3}$  Å except for the C<sub>1</sub>-C<sub>1</sub> bond, its standard error being larger ( $3 \times 10^{-3}$  Å) because the two carbon atoms are related by a center of symmetry. The standard error in the bond angles is 10'.

The final structure-factor agreement is very satisfactory. In Table 2 the observed and calculated (0kl)and (1kl) structure factors are compared.\* The discrepancy factors are given in Table 3 as a function of h; they are virtually independent of h. The over-all R factor is 0.089. Structure factors too weak to be observed were omitted; with very few exceptions they calculated weaker than the minimum observable values. The few very strong reflections show pronounced extinction and are also omitted in the calculation of the R factor.

As a final check of the results a Fourier section and a difference Fourier section at  $x = \frac{1}{4}$  were calculated. The Fourier section (Fig. 1(*a*)) is remarkable for the resolution and the heights of the electron density peaks, their heights being about 14, 16 and 18 e.Å<sup>-3</sup> for the carbon, nitrogen and oxygen atoms, respec-

<sup>\*</sup> A list of about 1100 values of  $F_o$  and  $F_c$  has been deposited as Document No. 4719 with the ADI Auxiliary Publications Project, Photoduplication Service, Library of Congress, Washington 25, D.C., U.S.A. A copy may be secured by citing the Document number and by remitting \$2.50 for photoprints, or \$1.75 for 35 mm. microfilm. Advance payment is required. Make checks or money orders payable to: Chief, Photoduplication Service, Library of Congress.

## DAVID R. DAVIES AND R. A. PASTERNAK

## Table 2. Observed and calculated structure factors

The calculations are based on McWeeny scattering curves (1951). The F's are multiplied by ten.

k	e	F.,	Fc	k	٤	Р,	Fc	k	ł	Fo	F.	k	e	Fo	P c
N N N N N N N N N N N N N N N N N N N	د 246802468 1234567890123. 11111	Fo 0k 3113076847 3551682998791230 544214758847 5 2945682998791230 5 228 3 498791230	F 945328100 58334696087682471 942 512326 304467239785155 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	k 1001100 1000100 1001100 1001100 1001100 11222 1122 112 1122 1122 1122 1122 112 1122 1122 1122 1122 112 1122 1122 1122 1122 1122 1122 1122 112 112 1122 1122 1122 1122 1122 1122 1122 112 1122 1122 1122 1122 1122 1122 1122 1122 1122 1122 1122 1122 1122 112 1122 1122 1122 112 1122 112 1111 1111 1111 1111 11111 11111 111111	2 2345678901123 1234567890112 1111	F 0 40999012045556 2222255569555665 50 6222655564 10 60 355569555665 50 10 60 555665 50	F <sub>0</sub> 4772577537729099 473334111377484 6 3	¥ 5555 777777777777777999999	4 3456 1234567890123456 1234	Fo 64556 434719500270141777 1699491	F <sub>c</sub> 3844 24 	* 333333333333333 555555555555555555555	د 345678901234567 1234567890122	Fo 75330602442893556 7829552464420 22311112943556 78295524664202	$\begin{array}{c} P_{c} \\ 292614\\ 202514\\ 2017213\\ 2017213\\ 14334\\ -1123534\\ 126535\\ 2017213\\ -2017213\\ 2017212\\ 2017212\\ 2017212\\ 2017212\\ 2017212\\ 2017212\\ 2017212\\ 2017212\\ 2017212\\ 2017212\\ 2017212\\ 2017212\\ 2017212\\ 2017212\\ 2017212\\ 2017212\\ 2017212\\ 2017212\\ $
~~~~~	14 15 16 17 18	59 <33 91 <35 72 295 121	- 51 - 95 - 25 - 79 287 - 82	14 14 14 14 14 14	1234567	66 <35 75 67 <36 44 45	- 64 9 - 80 66 15 50 44	9 9 9 9 9 9 9 9	567890 1112	74 <29 148 <31 40 48 <35	$ \begin{array}{r} 76 \\ -10 \\ 162 \\ -21 \\ -51 \\ -57 \\ -28 \\ -28 \\ \end{array} $	5 5 5 7 7 7 7	13 14 15 16 1 2 3	59 <33 <34 <36 59 79 <17	- 52 13 - 29 - 75 - 62 9
4444444444444	234567890112345 1112345	2 2 4 8 3 7 5 8 3 7 5 8 3 7 5 8 3 7 5 8 3 7 5 8 3 9 5 9 7 1 4 5 3 1 4 6 2 9 7 1 5 3	$\begin{array}{r} 313\\43\\15\\7\\-96\\4\\99\\-119\\33\\5\\15\\5\\-61\end{array}$	111111111111111111111111111111111111111	1234567890 10	Lkt 47 157 127 125 125 125 135 135 135 157	$\begin{array}{r} - 38\\ - 1669\\ 2833\\ - 74\\ - 1364\\ - 2801\\ - 183\end{array}$	9 9 9 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	13 14 15 10 12 34 56 78	<pre>&lt;36 &lt;37 &lt;37 &lt;30 161 103 33 4 </pre>	- 372 - 326 - 25 - 1899 - 1129 - 1129 - 1129 - 1129 - 117	77777777777777777777777777777777777777	45678901123456 111211111	48 999 1730 3477 4588 83 83 83	$\begin{array}{r} 44\\ 770\\ 9992\\ -\\ 4615\\ -\\ 11155\\ -\\ 869\\ -\\ 63\\ -\\ 869\\ -\\ -\\ 869\\ -\\ -\\ -\\ -\\ -\\ -\\ -\\ -\\ -\\ -\\ -\\ -\\ -\\$
44 66666	16 17 12 3	61 <36 15 206 186 337 217	- 63 - 31 18 215 187 360	111111111	11 12 13 14 15 16 17	<pre>&lt;2 0 19 &lt;31 60 &lt;34 &lt;35 &lt;36</pre>	- 15 29 59 - 12 36	11 13 13 13 13 13	9 1 2 3 4	<pre>&lt;35 &lt;34 &lt;34 34 65 &lt;35</pre>	- 40 16 - 37 71 - 12	9 9 9 9 9 9 9 9 9 9 9 9	12345670	51 162 60 112 92 <28 171	- 43 153 - 48 97 85 181
00000000000	45678901123 1123	1267 275 1137 333 330 331 332	133 - 32 - 108 - 55 - 35 - 35 3	3 3 3 3 3 3 3 3 3 3 3 3 3 3	12345678	16 4996 2399 185 298 30 146	- 17 - 575 378 241 194 278 - 97 27 - 167	15 15 15 15 15 15	1 2 3 4 5 1	<37 <37 <37 <37 <37 78 <39 93	1 28 - 26 37 - 78 6 96	9999999999999	89011231151	<pre>     59     51     71     34     44     44     &lt;37     &lt;37 </pre>	- 396 - 396 - 396 - 32 - 32
666 88888	14 15 16 12 34	59 35 25 102 130 16 143 104	$ \begin{array}{r} - 52 \\ 30 \\ - 31 \\ - 84 \\ 146 \\ - 15 \\ 161 \\ 106 \\ \end{array} $	3 3 3 3 3 3 3 3 3 3 3 3 3	901234567 11234567	<pre>&lt;35 122 21 86 38 &lt;33 54 &lt;36 &lt;36 &lt;36</pre>	$ \begin{array}{r} - & 2 \\ - & 147 \\ 20 \\ - & 91 \\ - & 39 \\ - & 29 \\ - & 44 \\ 12 \\ \end{array} $	1 1 1 1 1	123456	1kł 93 148 311 141 87 92	81 157 340 -138 87 93	11 11 11 11 11 11 11 11	123456789	80 181 75 121 <32 <32 <33 <34 42	$ \begin{array}{r} 78\\ 198\\ 75\\ 29\\ -\\ 30\\ -\\ 30\\ -\\ 46\\ \end{array} $
588888888888	567890112345	549 673 52 319 52 339 534 436 44	- 48 147 - 60 88 44 3 - 49 21 - 53 44	5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5	123456789	V11 224 84 211 1780 990 10	$ \begin{array}{r} - & 1 \\ - & 2 & 3 \\ 4 & 9 \\ 7 & 2 & 1 \\ - & 2 & 1 & 9 \\ - & 2 & 8 & 2 \\ - & 7 & 0 \\ - & 7 & 0 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 \\ - & 1 & 9 $	1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	7 8 9012345 67	154 3354 1214 550634 336 336	-162 349 -128 -128 -128 -45 -84 18 200 49	1333 113 1555 1555	1234 12345	<pre>&lt;34 34 69 &lt;35 &lt;37 &lt;37 &lt;37 &lt;37 &lt;37 &lt;37 &lt;37 &lt;37 &lt;37 &lt;37</pre>	148370 771 276808 888
10 10	1	152 53	44 141 46	5 5 5	10 11 12	29 59 <31	- 18 19 - 58 - 23	1 3 3	12	537 408	- 595 - 434	15 15 17	5 6 1	79 103	67 98

tively. An elongation of the peaks in the z direction is noticeable, an indication of temperature anisotropy. The most striking feature of the section, however, is the high electron density between bonded atoms, especially in the bonds  $\rm C_1-\rm C_2$  and  $\rm C_2-\rm N.$ 

This feature is also apparent in the difference-



Fig. 1. (a) Fourier section at  $x = \frac{1}{4}$ . Broken contour at 1 e.Å<sup>-3</sup>, other contours at 2, 4, 6, ... e.Å<sup>-3</sup>. (b) Difference Fourier section at  $x = \frac{1}{4}$ . Contours at 0.25, 0.5, 0.75... e.Å<sup>-3</sup>. Negative contours are broken.

Fourier section (Fig. 1(b)). The general background of the difference Fourier is very low, almost everywhere less than  $\pm 0.25$  e.Å<sup>-3</sup>. The hydrogen atoms, which are attached to the nitrogen atom, appear as well-defined, fairly round peaks of heights about 1 e.Å-3; the N-H bonds are very probably coplanar with the molecule and form angles of about 120° with the C-N bond. Hydrogen atoms  $H_1$  and  $H_2$ , which are 0.8 Å above and below the plane, also seem to show up weakly. All the heavy atoms fall into holes, the oxygen atom into a fairly deep one. By contrast, a positive electron density is found between bonded atoms, especially between C<sub>2</sub> and N. As for the physical meaning of this observed electron-density variation, it is difficult to escape the conclusion that a partial displacement of electrons from the atoms into the bonds is being observed. However, it should be pointed out that all refinements were carried out by least-squares methods, and the weights given to individual reflections were different from those in the difference Fourier synthesis; refinement of the temperature factors based on the difference Fourier and, in addition, introduction of the anisotropic f curve for the oxygen atom (McWeeny,

Table 3. Discrepancy factor R as a function of h

	Number of observed	
h	reflections	R
0	89	0.082
1	141	0.084
2	165	0.085
3	154	0.092
4	128	0.087
5	135	0.090
6	101	0.099
7	96	0.112
8	69	0.096
hkl	1078	0.089

1951) might have smoothed the difference Fourier somewhat.

It is felt that the refinement has approached the limit set by the errors in the intensity measurements; more convincing evidence for the displacement of the electrons would have to be based on more accurate experimental data.

#### **Discussion of the structure**

#### Molecular configuration

One layer of the structure of succinamide is shown in Fig. 2; bond lengths and angles are also listed in Table 4.

## Table 4. Bond lengths and angles

C1-C1	1·501 Å	$C_1 - C_1 - C_2$	113° 54′
$C_1 - C_2$	1.512	$C_1 - C_2 - N$	115° 36′
$C_2 - N$	1.333	$C_1 - C_2 - O$	122° 23′
$C_2 - O$	1.238	$\tilde{N-C_2-O}$	122° 2'
NT O	∫ 2·942		( 119° 23'
N 0	2.939	$C_2 - N \cdots O$	120° 5′

A comparison of the molecular configuration of succinamide with those of succinic acid (Morrison & Robertson, 1949) and oxamide (Ayerst & Duke, 1954) is most fruitful. Both of these structures, and especially the latter, have been determined with great care.

The succinamide molecule is perfectly planar; no atom deviates from the median plane by more than the standard error associated with it. This contrasts with the configuration of succinic acid, in which the planes of the carboxyl groups are rotated by  $9^{\circ}$  in respect to the plane of the carbon chain, the four carbon atoms being coplanar. However, perfect planarity was observed for the molecule of oxamide. Of the two non-equivalent C–C bonds the central  $C_1-C_1$  bond is slightly, but significantly, shorter (1.501 Å compared to 1.512 Å); both bonds are shorter than the value of 1.54 Å usually assigned to



Fig. 2. A schematic drawing showing the arrangement of the molecules in the layer  $x = \frac{1}{4}$ .

a C-C single bond. Furthermore, the  $C_1-C_1-C_2$  bond angle (113° 54') is appreciably larger than the tetrahedral angle. These observations are in accord with the tentative results of Morrison & Robertson (1949) in their studies of succinic acid and other dicarboxylic acids containing an even number of carbon atoms. In particular, they report an alternation between long and short C-C bonds along the chain, the bond adjacent to the carboxyl group always being the longer. This systematic variation in the bond lengths might be explained by hyperconjugation.

In the amide group of succinamide the C–O bond has the same length as in oxamide, but the C–N bond

Table 5. Bond lengths in the amide group

	C-O (Å)	C–N (Å)	Reported standard error (Å)
Acetamide	1.28	1.38	?
Succinamide	1.238	1.333	0.002
Oxamide	1.243	1.315	0.004
Formamide	1.225	1.30	0.017
Glycyl asparagine*	1.22	1.36	0.014
Glutamine	1.27	1.28	0.024
Peptide (average)†	1.24	1.32	—

\* A value of 1.39 Å for the C-N bond had been reported in the original paper (Pasternak *et al.*, 1954). On checking the calculations, an error in the *x* coordinate of nitrogen atom  $N_1$  has been discovered (correct x = -0.064); the corrected bond length is reported here. Other molecular dimensions involving this atom do not change significantly.

† Corey & Pauling, 1953.

is somewhat longer (1.333 Å compared with 1.315 Å). It is interesting to note that, except for this small though significant difference, the lengths of the C–O and of the C–N bonds in the amide groups of all terminal amides and peptides studied to date have been found to be the same within the limits of error of the investigations (Table 5).

The amide group of succinamide, however, differs pronouncedly from other amides in the relative values of the three bond angles around the carbon atom (Table 6). In succinamide angle O-C-C is slightly

Table 6. Bond angles in the amide group

N-C-C (°)	O-C-C (°)	0-C-N (°)
109	129	122
115.6	$122 \cdot 4$	122.0
114.8	119.5	125.7
115.2	$124 \cdot 9$	119.9
118	118	123
114	121	125
	N-C-C (°) 109 115·6 114·8 115·2 118 114	$\begin{array}{ccc} \text{N-C-C} (°) & \text{O-C-C} (°) \\ 109 & 129 \\ 115\cdot6 & 122\cdot4 \\ 114\cdot8 & 119\cdot5 \\ 115\cdot2 & 124\cdot9 \\ 118 & 118 \\ 114 & 121 \end{array}$

\* In terminal amide.

† Corey & Pauling, 1953.

larger than angle O–C–N, but in oxamide and in peptide amide links the former angle is significantly smaller than the latter. The variation in bond angles as found in peptide amide links agrees with what one might expect on the basis of resonance considerations. It is interesting, however, that in succinamide, and even more pronouncedly in the terminal amide group of its derivative glycylasparagine, the angles deviate from the predicted ones. Similar deviations in the relative magnitudes of the angles were reported also for the carboxyl groups of N-acetylglycine (Carpenter & Donohue, 1950), succinic acid, and glutaric acid (Morrison & Robertson, 1949).

Finally, the three bonds formed by the nitrogen atom in succinamide are coplanar and trigonally arranged within the accuracy of the determination of the hydrogen atoms.

#### Arrangement of the molecules

The plane of the succinamide molecules is slightly but significantly tilted in respect to the (100) plane; the two planes intersect approximately along the C1-C1 bond, and the dihedral angle between them is  $1.0^{\circ}$ . The molecules in the layer are connected by a net of hydrogen bonds. There are two sets of non-equivalent hydrogen bonds, which have, however, the same length and form angles of very nearly 120° with the C-N bond. The hydrogen bonds are arranged not quite so symmetrically around the oxygen atoms, the angles being 118° 35' and 138° 31'. One set of hydrogen bonds joins the amide groups of adjacent molecules into an eight-membered ring around a center of symmetry. The same ring formation is found in oxamide, which displays a very similar arrangement of the molecules in the layer, and in formamide (Ladell & Post, 1954). It is interesting that these hydrogen

bonds, which are of identical length in the three structures, are fairly long, 2.94 Å.

The spacing between molecular layers is 3.385 Å, compared with 3.007 Å in oxamide. The increase in spacing is reasonable considering the presence of the bulky methylene groups in succinamide; in both structures, however, the packing is very loose, assuming conventional van der Waals radii for the individual atoms.

The superposition of two molecular layers in the unit cell is shown in Fig. 3. In this view along a, the



Fig. 3. A schematic drawing of the structure viewed along the a axis of the crystal. The molecular layers are at  $x = \frac{1}{4}$  and  $x = \frac{3}{4}$ , and the twofold axes are at  $x = 0, \frac{1}{2}$ .

structure might be described as consisting of sheets of molecules parallel to the (001) plane; these sheets are held together by hydrogen bonds. Finally, the anisotropic temperature factors of the structure (Table 1) are well explained by the packing of the molecules. The largest temperature vibrations are, as expected, perpendicular to the molecular layers. The oxygen and nitrogen atoms, which form only one covalent bond (not counting those with hydrogen atoms), have the largest temperature factors, and carbon atom  $C_2$ , which forms three bonds, has the smallest. Within the layer the temperature factors are small, vibrations along c being somewhat larger than along b.

The authors wish to express their gratitude to Dr Richard E. Marsh for critical review of the manuscript.

### References

- AYERST, E. M. & DUKE, J. R. C. (1954). Acta Cryst. 7, 588.
- CARPENTER, G. B. & DONOHUE, J. (1950). J. Amer. Chem. Soc. 72, 2315.
- COCHRAN, W. & PENFOLD, B. R. (1952). Acta Cryst. 5, 644.
- COREY, R. B. & PAULING, L. (1953). Proc. Roy. Soc. B, 141, 10.
- DAVIES, D. R. & BLUM, J. J. (1955). Acta Cryst. 8, 129.
- LADELL, J. & POST, B. (1954). Acta Cryst. 7, 559.
- MCWEENY, R. (1951). Acta Cryst. 4, 513.
- MORRISON, J. D. & ROBERTSON, J. M. (1949). J. Chem. Soc. pp. 980 et seq.
- PASTERNAK, R. A. (1953). Acta Cryst. 6, 808.
- PASTERNAK, R.A., KATZ, L. & COREY, R. B. (1954). Acta Cryst. 7, 225.
- ROLLETT, J. S. & DAVIES, D. R. (1955). Acta Cryst. 8, 125.
- SENTI, F. & HARKER, D. (1940). J. Amer. Chem. Soc. 62, 2008.