data\_k05kcm23 \_audit\_creation\_method SHELXL-97 \_chemical\_name\_systematic ? ; ? \_chemical\_name\_common ? \_chemical\_melting\_point ? \_chemical\_formula\_moiety \_chemical\_formula\_sum 'C16 H20 Cl6 N2 O2 Sn' \_chemical\_formula\_weight 603.73 loop\_ \_atom\_type\_symbol \_atom\_type\_description \_atom\_type\_scat\_dispersion\_real \_atom\_type\_scat\_dispersion\_imag \_atom\_type\_scat\_source 'C' 'C' 0.0033 0.0016 'International Tables Vol C Tables 4.2.6.8 and 6.1.1.4' 'H' 'H' 0.0000 0.0000 'International Tables Vol C Tables 4.2.6.8 and 6.1.1.4' 'Cl' 'Cl' 0.1484 0.1585 'International Tables Vol C Tables 4.2.6.8 and 6.1.1.4' 'N' 'N' 0.0061 0.0033 'International Tables Vol C Tables 4.2.6.8 and 6.1.1.4' 'O' 'O' 0.0106 0.0060 'International Tables Vol C Tables 4.2.6.8 and 6.1.1.4' 'Sn' 'Sn' -0.6537 1.4246 'International Tables Vol C Tables 4.2.6.8 and 6.1.1.4' \_symmetry\_cell\_setting ? \_symmetry\_space\_group\_name\_H-M ? loop\_ \_symmetry\_equiv\_pos\_as\_xyz 'x, y, z' '-x, -y, -z' \_cell\_length\_a 7.37800(10) \_cell\_length\_b 7.44800(10) \_cell\_length\_c 11.5170(2) \_cell\_angle\_alpha 97.6240(10) \_cell\_angle\_beta 91.4810(10) \_cell\_angle\_gamma 117.3890(10) \_cell\_volume 554.243(14) \_cell\_formula\_units\_Z 1

```
_cell_measurement_temperature
                                  150(2)
                                 ?
_cell_measurement_reflns_used
cell measurement theta min
                                 ?
                                 ?
_cell_measurement_theta_max
_exptl_crystal_description
                              ?
                            ?
_exptl_crystal_colour
                              ?
_exptl_crystal_size_max
                             ?
_exptl_crystal_size_mid
                             ?
_exptl_crystal_size_min
                               ?
_exptl_crystal_density_meas
_exptl_crystal_density_diffrn
                               1.809
_exptl_crystal_density_method
                                'not measured'
_exptl_crystal_F_000
                             298
exptl absorpt coefficient mu
                                1.891
_exptl_absorpt_correction_type
                                ?
_exptl_absorpt_correction_T_min ?
exptl absorpt correction T max ?
_exptl_absorpt_process_details ?
_exptl_special_details
?
;
_diffrn_ambient_temperature
                                150(2)
_diffrn_radiation_wavelength
                               0.71073
_diffrn_radiation_type
                            MoK\a
_diffrn_radiation_source
                             'fine-focus sealed tube'
_diffrn_radiation_monochromator graphite
_diffrn_measurement_device_type ?
                                 ?
_diffrn_measurement_method
_diffrn_detector_area_resol_mean ?
_diffrn_standards_number
diffrn standards interval count ?
_diffrn_standards_interval_time
                                ?
_diffrn_standards_decay_%
                               ?
diffrn reflns number
                             8539
_diffrn_reflns_av_R_equivalents 0.0270
_diffrn_reflns_av_sigmaI/netI
                               0.0261
_diffrn_reflns_limit_h_min
                              -8
                               8
diffrn reflns limit h max
_diffrn_reflns_limit_k_min
                              -8
_diffrn_reflns_limit_k_max
                               8
diffrn reflns limit 1 min
                              -13
_diffrn_reflns_limit_l_max
                              13
_diffrn_reflns_theta_min
                             3.85
_diffrn_reflns_theta_max
                              24.97
_reflns_number_total
                            1909
reflns number gt
                            1890
```

reflns threshold expression >2sigma(I) ? computing data collection ? \_computing\_cell\_refinement ? \_computing\_data\_reduction \_computing\_structure\_solution ? \_computing\_structure\_refinement 'SHELXL-97 (Sheldrick, 1997)' computing molecular graphics \_computing\_publication\_material ? \_refine\_special\_details Refinement of F^2<sup>^</sup> 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  $F^{2^{}} > 2sigma(F^{2^{}})$  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 Rfactors based on ALL data will be even larger. : \_refine\_ls\_structure\_factor\_coef Fsqd refine ls matrix type full \_refine\_ls\_weighting\_scheme calc \_refine\_ls\_weighting details 'calc w=1/[ $s^2(Fo^2)$ +(0.0120P)<sup>2</sup>+0.2090P] where P=(Fo<sup>2</sup>+2Fc<sup>2</sup>)/3' \_atom\_sites\_solution\_primary direct \_atom\_sites\_solution\_secondary difmap \_atom\_sites\_solution\_hydrogens geom refine ls hydrogen treatment mixed \_refine\_ls\_extinction\_method none ? \_refine\_ls\_extinction\_coef \_refine\_ls\_number\_reflns 1909 \_refine\_ls\_number\_parameters 136 refine ls number restraints 0 \_refine\_ls\_R\_factor\_all 0.0156 \_refine\_ls\_R\_factor\_gt 0.0154 refine ls wR factor ref 0.0390 \_refine\_ls\_wR\_factor\_gt 0.0388 \_refine\_ls\_goodness\_of\_fit\_ref 1.104 \_refine\_ls\_restrained\_S\_all 1.104 refine ls shift/su max 0.001 \_refine\_ls\_shift/su\_mean 0.000 loop\_ \_atom\_site\_label \_atom\_site\_type\_symbol \_atom\_site\_fract\_x \_atom\_site\_fract\_v \_atom\_site\_fract\_z

atom site U iso or equiv \_atom\_site\_adp\_type atom site occupancy \_atom\_site\_symmetry\_multiplicity \_atom\_site\_calc\_flag \_atom\_site\_refinement\_flags \_atom\_site\_disorder\_assembly atom site disorder group Sn Sn 0.5000 0.0000 0.0000 0.01455(7) Uani 1 2 d S . . Cl1 Cl 0.16150(6) -0.22514(6) 0.05433(3) 0.02076(10) Uani 1 1 d ... Cl2 Cl 0.64998(6) -0.18451(6) 0.09292(3) 0.02024(10) Uani 1 1 d ... Cl3 Cl 0.56731(6) 0.22768(6) 0.18357(3) 0.02169(10) Uani 1 1 d . . . O1 O 1.36508(17) 0.52884(18) 0.36931(10) 0.0281(3) Uani 1 1 d . . . N1 N 1.2489(2) 0.4143(3) 0.14543(14) 0.0232(3) Uani 1 1 d ... H10A H 1.216(4) 0.497(4) 0.120(2) 0.050(7) Uiso 1 1 d . . . H10B H 1.262(4) 0.338(4) 0.087(2) 0.055(7) Uiso 1 1 d . . . H10C H 1.366(4) 0.494(4) 0.186(2) 0.052(7) Uiso 1 1 d . . . C1 C 1.1055(2) 0.2873(3) 0.22441(14) 0.0224(3) Uani 1 1 d . . . H1A H 1.0878 0.1459 0.2086 0.027 Uiso 1 1 calc R . . H1B H 0.9697 0.2812 0.2104 0.027 Uiso 1 1 calc R . . C2 C 1.1941(2) 0.3834(2) 0.35142(14) 0.0199(3) Uani 1 1 d . . . C3 C 1.0664(2) 0.2936(2) 0.44582(14) 0.0189(3) Uani 1 1 d . . . C4 C 1.1441(3) 0.3808(2) 0.56302(14) 0.0220(3) Uani 1 1 d . . . H4 H 1.2768 0.4952 0.5808 0.026 Uiso 1 1 calc R . . C5 C 1.0272(3) 0.2998(3) 0.65272(14) 0.0255(4) Uani 1 1 d . . . H5 H 1.0795 0.3592 0.7323 0.031 Uiso 1 1 calc R ... C6 C 0.8340(3) 0.1323(3) 0.62699(15) 0.0257(4) Uani 1 1 d . . . H6 H 0.7537 0.0786 0.6891 0.031 Uiso 1 1 calc R . . C7 C 0.7570(3) 0.0425(3) 0.51110(15) 0.0240(4) Uani 1 1 d . . . H7 H 0.6257 -0.0741 0.4939 0.029 Uiso 1 1 calc R . . C8 C 0.8728(2) 0.1238(2) 0.42053(14) 0.0216(3) Uani 1 1 d . . . H8 H 0.8200 0.0636 0.3410 0.026 Uiso 1 1 calc R . . loop \_atom\_site\_aniso\_label \_atom\_site\_aniso\_U\_11 atom site aniso U 22 atom site aniso U 33 \_atom\_site\_aniso\_U\_23 \_atom\_site\_aniso\_U\_13 atom site aniso U 12 Sn 0.01356(10) 0.01427(10) 0.01314(10) -0.00040(6) -0.00002(6) 0.00500(7) Cl1 0.0150(2) 0.0224(2) 0.0208(2) 0.00499(16) 0.00225(15) 0.00495(17) C12 0.0202(2) 0.0200(2) 0.0205(2) 0.00314(15) -0.00052(15) 0.00952(17) C13 0.0228(2) 0.0233(2) 0.01627(19) -0.00566(15) -0.00246(15) 0.01108(17) 01 0.0199(6) 0.0277(6) 0.0269(6) -0.0001(5) -0.0011(5) 0.0043(5) N1 0.0233(8) 0.0247(8) 0.0223(8) 0.0046(7) 0.0031(6) 0.0116(7) C1 0.0190(8) 0.0241(8) 0.0211(8) 0.0038(7) 0.0021(6) 0.0075(7) C2 0.0193(8) 0.0202(8) 0.0227(8) 0.0005(6) -0.0015(6) 0.0123(7) C3 0.0203(8) 0.0198(8) 0.0211(8) 0.0015(6) 0.0009(6) 0.0136(7)

C4 0.0228(9) 0.0189(8) 0.0233(8) -0.0017(6) -0.0024(7) 0.0105(7) C5 0.0346(10) 0.0264(9) 0.0182(8) -0.0015(7) 0.0002(7) 0.0180(8) C6 0.0305(9) 0.0266(9) 0.0247(9) 0.0058(7) 0.0077(7) 0.0167(8) C7 0.0218(9) 0.0213(8) 0.0301(9) 0.0038(7) 0.0026(7) 0.0111(7) C8 0.0220(8) 0.0232(8) 0.0209(8) -0.0007(7) -0.0023(7) 0.0129(7)

\_geom\_special\_details

All esds (except the esd in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell esds are taken into account individually in the estimation of esds in distances, angles and torsion angles; correlations between esds in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell esds is used for estimating esds involving l.s. planes. ;

loop\_

geom bond atom site label 1 \_geom\_bond\_atom\_site\_label\_2 \_geom\_bond\_distance \_geom\_bond\_site\_symmetry\_2 \_geom\_bond\_publ\_flag Sn Cl3 2.4107(4) 2 655 ? Sn Cl3 2.4107(4) . ? Sn Cl1 2.4386(4) 2 655 ? Sn Cl1 2.4386(4).? Sn Cl2 2.4467(4) . ? Sn Cl2 2.4467(4) 2\_655 ? O1 C2 1.215(2) . ? N1 C1 1.483(2).? N1 H10A 0.83(3).? N1 H10B 0.86(3).? N1 H10C 0.87(3).? C1 C2 1.524(2).? C1 H1A 0.9900 . ? C1 H1B 0.9900.? C2 C3 1.478(2) . ? C3 C8 1.394(2).? C3 C4 1.399(2).? C4 C5 1.381(2).? C4 H4 0.9500 . ? C5 C6 1.386(2).? C5 H5 0.9500 . ? C6 C7 1.388(2).? C6 H6 0.9500 . ? C7 C8 1.386(2).? C7 H7 0.9500.? C8 H8 0.9500.?

loop\_

\_geom\_angle\_atom\_site\_label\_1 \_geom\_angle\_atom\_site\_label 2 \_geom\_angle\_atom\_site\_label\_3 \_geom\_angle \_geom\_angle\_site\_symmetry\_1 \_geom\_angle\_site\_symmetry\_3 \_geom\_angle\_publ\_flag Cl3 Sn Cl3 180.000(11) 2\_655 . ? Cl3 Sn Cl1 89.722(13) 2\_655 2\_655 ? Cl3 Sn Cl1 90.278(13) . 2\_655 ? Cl3 Sn Cl1 90.278(13) 2 655.? Cl3 Sn Cl1 89.722(13) . . ? Cl1 Sn Cl1 180.00(3) 2\_655 . ? Cl3 Sn Cl2 90.030(13) 2\_655 . ? Cl3 Sn Cl2 89.970(13) . . ? Cl1 Sn Cl2 88.929(13) 2\_655 . ? Cl1 Sn Cl2 91.071(12) . . ? Cl3 Sn Cl2 89.970(13) 2 655 2 655 ? Cl3 Sn Cl2 90.030(13) . 2\_655 ? Cl1 Sn Cl2 91.071(13) 2\_655 2\_655 ? Cl1 Sn Cl2 88.929(13) . 2\_655 ? Cl2 Sn Cl2 180.000(19) . 2\_655 ? C1 N1 H10A 114.2(17) . . ? C1 N1 H10B 110.8(17) . . ? H10A N1 H10B 109(2) . . ? C1 N1 H10C 108.3(16) . . ? H10A N1 H10C 103(2) . . ? H10B N1 H10C 111(2)..? N1 C1 C2 108.49(13) . . ? N1 C1 H1A 110.0 . . ? C2 C1 H1A 110.0 . . ? N1 C1 H1B 110.0 . . ? C2 C1 H1B 110.0 . . ? H1A C1 H1B 108.4 . . ? O1 C2 C3 123.83(15) . . ? O1 C2 C1 118.35(14) . . ? C3 C2 C1 117.82(14) . . ? C8 C3 C4 119.73(15) . . ? C8 C3 C2 121.53(15) . . ? C4 C3 C2 118.73(15) . . ? C5 C4 C3 119.75(15) . . ? C5 C4 H4 120.1 . . ? C3 C4 H4 120.1 . . ? C4 C5 C6 120.25(16) . . ? C4 C5 H5 119.9 . . ? C6 C5 H5 119.9 . . ? C5 C6 C7 120.43(16) . . ? C5 C6 H6 119.8 . . ? C7 C6 H6 119.8 . . ? C8 C7 C6 119.66(16) . . ?

C8 C7 H7 120.2 . . ? C6 C7 H7 120.2 . . ? C7 C8 C3 120.15(15) . . ? C7 C8 H8 119.9 . . ? C3 C8 H8 119.9 . . ?

\_diffrn\_measured\_fraction\_theta\_max 0.983 \_diffrn\_reflns\_theta\_full 24.97 \_diffrn\_measured\_fraction\_theta\_full 0.983 \_refine\_diff\_density\_max 0.285 \_refine\_diff\_density\_min -0.702 \_refine\_diff\_density\_rms 0.071