# checkCIF/PLATON report

Structure factors have been supplied for datablock(s) ACu5

THIS REPORT IS FOR GUIDANCE ONLY. IF USED AS PART OF A REVIEW PROCEDURE FOR PUBLICATION, IT SHOULD NOT REPLACE THE EXPERTISE OF AN EXPERIENCED CRYSTALLOGRAPHIC REFEREE.

## **Datablock: ACu5**

```
Bond precision: C-C = 0.0042 A
                                        Wavelength=0.71073
Cell:
               a=8.3204(17)
                                b=10.874(2)
                                                 c=12.726(3)
               alpha=76.29(3)
                                beta=75.33(3)
                                                 qamma = 83.76(3)
Temperature:
               298 K
               Calculated
                                         Reported
Volume
               1080.7(4)
                                         1080.7(4)
Space group
              P -1
                                         P -1
Hall group
               -P 1
                                         -P 1
Moiety formula C44 H38 Cl4 Cu2 N2 O8
Sum formula
              C44 H38 C14 Cu2 N2 O8
                                         C44 H38 C14 Cu2 N2 O8
Mr
               991.66
                                         991.64
               1.524
                                         1.524
Dx,g cm-3
Ζ
                                         1
Mu (mm-1)
               1.286
                                         1.286
F000
               506.0
                                         506.0
F000′
               507.41
h,k,lmax
               11,14,16
                                         11,14,16
Nref
               5360
                                         5327
               0.689,0.720
                                         0.780,0.911
Tmin,Tmax
Tmin'
               0.676
Correction method= # Reported T Limits: Tmin=0.780 Tmax=0.911
AbsCorr = NUMERICAL
Data completeness= 0.994
                                 Theta(max) = 28.281
R(reflections) = 0.0351( 4045) wR2(reflections) = 0.0888( 5327)
S = 1.008
                          Npar= 272
```

The following ALERTS were generated. Each ALERT has the format test-name\_ALERT\_alert-type\_alert-level.

Click on the hyperlinks for more details of the test.

```
Alert level C
PLAT242_ALERT_2_C Low
                        'MainMol' Ueg as Compared to Neighbors of
                                                                       C9 Check
PLAT334 ALERT 2 C Small Aver. Benzene C-C Dist C1 -C6
                                                                       1.37 Ang.
PLAT910_ALERT_3_C Missing # of FCF Reflection(s) Below Theta(Min).
                                                                         8 Note
PLAT911_ALERT_3_C Missing FCF Refl Between Thmin & STh/L= 0.600
                                                                         16 Report
  Alert level G
PLAT154_ALERT_1_G The s.u.'s on the Cell Angles are Equal ..(Note)
                                                                      0.03 Degree
PLAT794_ALERT_5_G Tentative Bond Valency for Cul
                                                                       2.10 Info
PLAT883_ALERT_1_G No Info/Value for _atom_sites_solution_primary .
                                                                   Please Do !
PLAT912_ALERT_4_G Missing # of FCF Reflections Above STh/L= 0.600
                                                                        10 Note
PLAT941_ALERT_3_G Average HKL Measurement Multiplicity .....
                                                                        2.0 Low
PLAT978 ALERT 2 G Number C-C Bonds with Positive Residual Density.
                                                                          2 Info
  0 ALERT level A = Most likely a serious problem - resolve or explain
  0 ALERT level B = A potentially serious problem, consider carefully
   4 ALERT level C = Check. Ensure it is not caused by an omission or oversight
  6 ALERT level G = General information/check it is not something unexpected
  2 ALERT type 1 CIF construction/syntax error, inconsistent or missing data
  3 ALERT type 2 Indicator that the structure model may be wrong or deficient
  3 ALERT type 3 Indicator that the structure quality may be low
  1 ALERT type 4 Improvement, methodology, query or suggestion
  1 ALERT type 5 Informative message, check
```

It is advisable to attempt to resolve as many as possible of the alerts in all categories. Often the minor alerts point to easily fixed oversights, errors and omissions in your CIF or refinement strategy, so attention to these fine details can be worthwhile. In order to resolve some of the more serious problems it may be necessary to carry out additional measurements or structure refinements. However, the purpose of your study may justify the reported deviations and the more serious of these should normally be commented upon in the discussion or experimental section of a paper or in the "special\_details" fields of the CIF. checkCIF was carefully designed to identify outliers and unusual parameters, but every test has its limitations and alerts that are not important in a particular case may appear. Conversely, the absence of alerts does not guarantee there are no aspects of the results needing attention. It is up to the individual to critically assess their own results and, if necessary, seek expert advice.

### Publication of your CIF in IUCr journals

A basic structural check has been run on your CIF. These basic checks will be run on all CIFs submitted for publication in IUCr journals (*Acta Crystallographica*, *Journal of Applied Crystallography*, *Journal of Synchrotron Radiation*); however, if you intend to submit to *Acta Crystallographica Section C* or *E* or *IUCrData*, you should make sure that full publication checks are run on the final version of your CIF prior to submission.

#### **Publication of your CIF in other journals**

Please refer to the *Notes for Authors* of the relevant journal for any special instructions relating to CIF submission.

### PLATON version of 10/08/2020; check.def file version of 06/08/2020

Datablock ACu5 - ellipsoid plot

