Electronic Supplementary Material (ESI) for New Journal of Chemistry.

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New Journal of Chemistry

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Supplementary material for

Tricarbonylrhenium(I) complexes with 2-acetylpyridine-derived hydrazones are cytotoxic to NCI-H460 human large cell lung cancer

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<sup>&</sup>lt;sup>b</sup> Departamento de Imunologia, Instituto de Ciências Biomédicas, Universidade de São Paulo, 05508-900, São Paulo, SP, Brazil

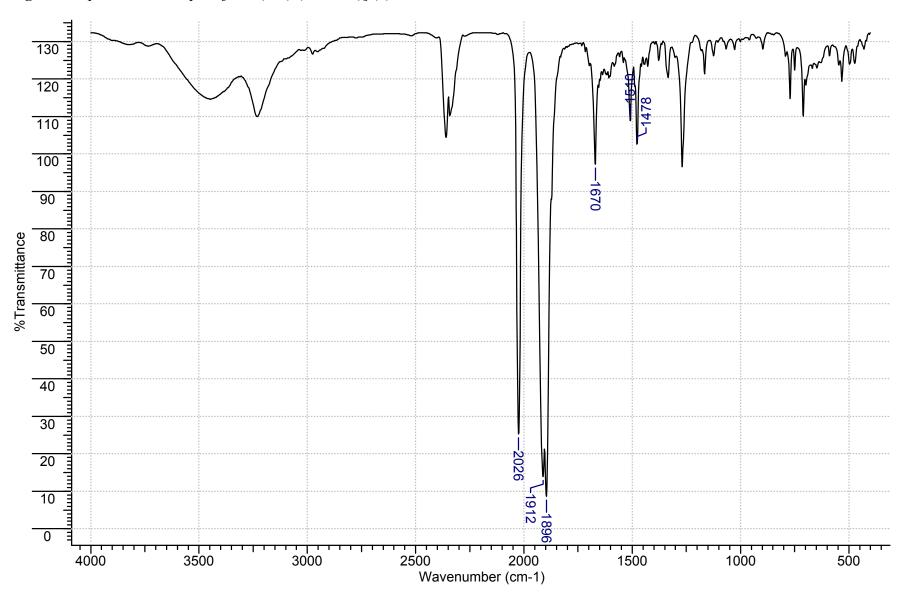
<sup>\*</sup> hberaldo@ufmg.br

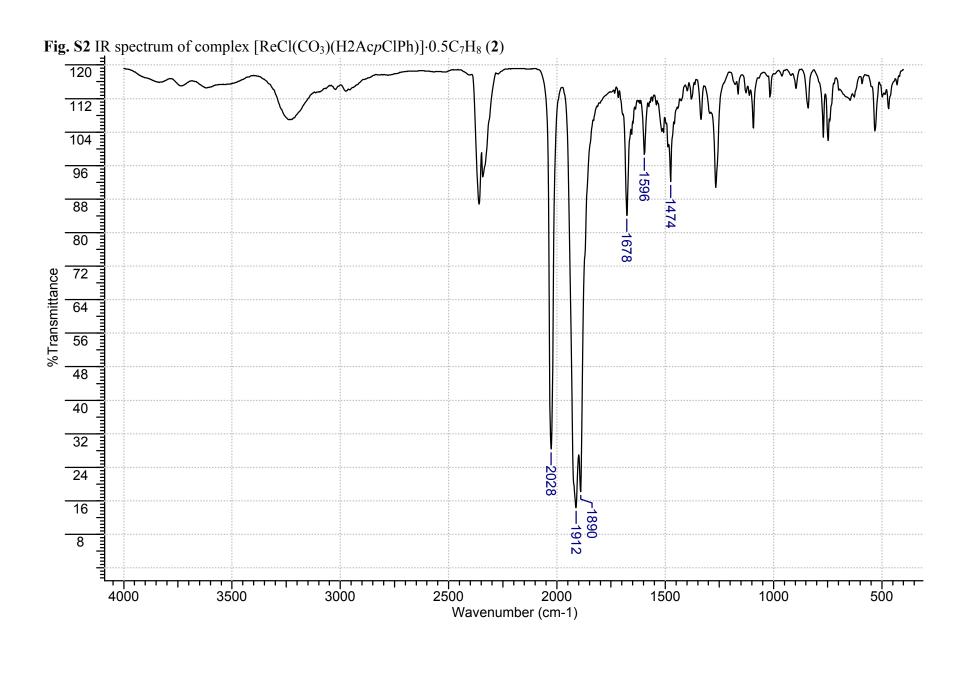
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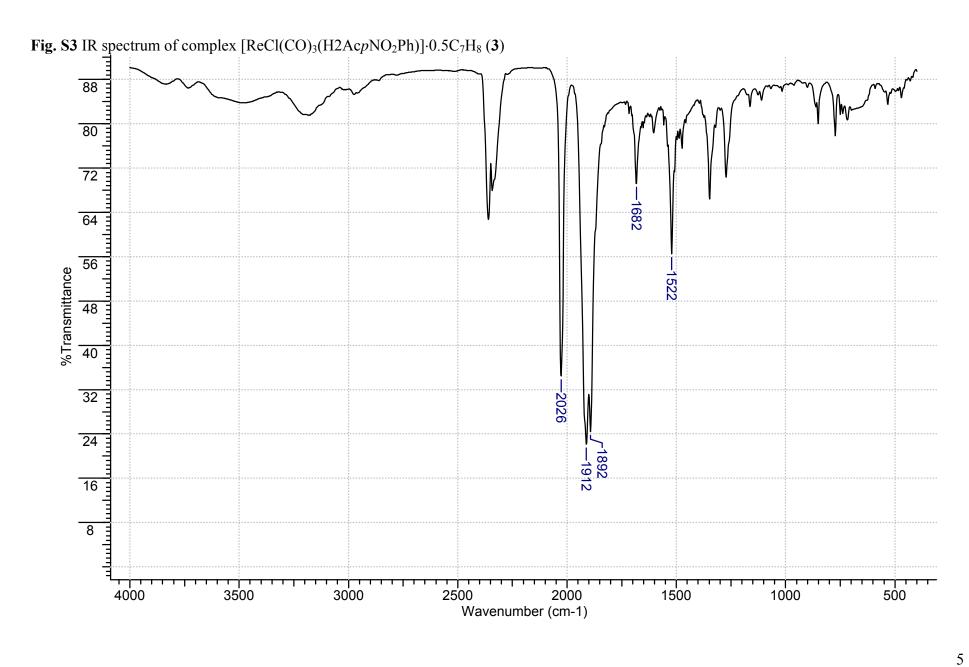
Infra	red spec	etra of	complexe	s [ReCl(CC	(H2A)	AcPh)] (1), [	$[ReCl(CO_3)(H2AcpClPh)]\cdot 0.5$	$5C_7H_8$
( <b>2</b> ) a	nd [ReC	Cl(CO)3	(H2AcpN	$[O_2Ph)]\cdot 0.5$	C <sub>7</sub> H <sub>8</sub> (3	3)	•••••••••••••••••••••••••••••••••••••••	2
<sup>1</sup> H	and	<sup>13</sup> C	NMR	spectra	of	complexes	[ReCl(CO) <sub>3</sub> (H2AcPh)]	(1),
[ReC	Cl(CO <sub>3</sub> )(	Н2Аср	oClPh)]·0.	5C <sub>7</sub> H <sub>8</sub> ( <b>2</b> ) a	and [Re	Cl(CO) <sub>3</sub> (H2	$AcpNO_2Ph)]\cdot 0.5C_7H_8$ (3)	6
CHECKCIFs		S	files	of	comp	lexes	[ReCl(CO) <sub>3</sub> (H2AcPh)]	(1),
[ReC	Cl(CO3)(	H2Acr	ClPh)]·D	MSO ( <b>2</b> a) a	and [Re	Cl(CO)3(H2	$(2AcpNO_2Ph)]\cdot DMSO(3a)$	23

Infrared spectra of complexes [ReCl(CO)<sub>3</sub>(H2AcPh)] (1), [ReCl(CO<sub>3</sub>)(H2AcpClPh)]·0.5C<sub>7</sub>H<sub>8</sub> (2) and [ReCl(CO)<sub>3</sub>(H2AcpNO<sub>2</sub>Ph)]·0.5C<sub>7</sub>H<sub>8</sub> (3)

Fig. S1 IR spectrum of complex [ReCl(CO)<sub>3</sub>(H2AcPh)] (1)







 $^{1}$ H and  $^{13}$ C NMR spectra of complexes [ReCl(CO)<sub>3</sub>(H2AcPh)] (1), [ReCl(CO<sub>3</sub>)(H2AcpClPh)]·0.5C<sub>7</sub>H<sub>8</sub> (2) and [ReCl(CO)<sub>3</sub>(H2AcpNO<sub>2</sub>Ph)]·0.5C<sub>7</sub>H<sub>8</sub> (3)

Fig. S4  $^{1}$ H NMR spectrum of complex [ReCl(CO)<sub>3</sub>(H2AcPh)] (1) in DMSO- $d_6$  (200 MHz) at room temperature 12.53 ight] 3.00

5.(

10.0

ppm (t1)

0.0

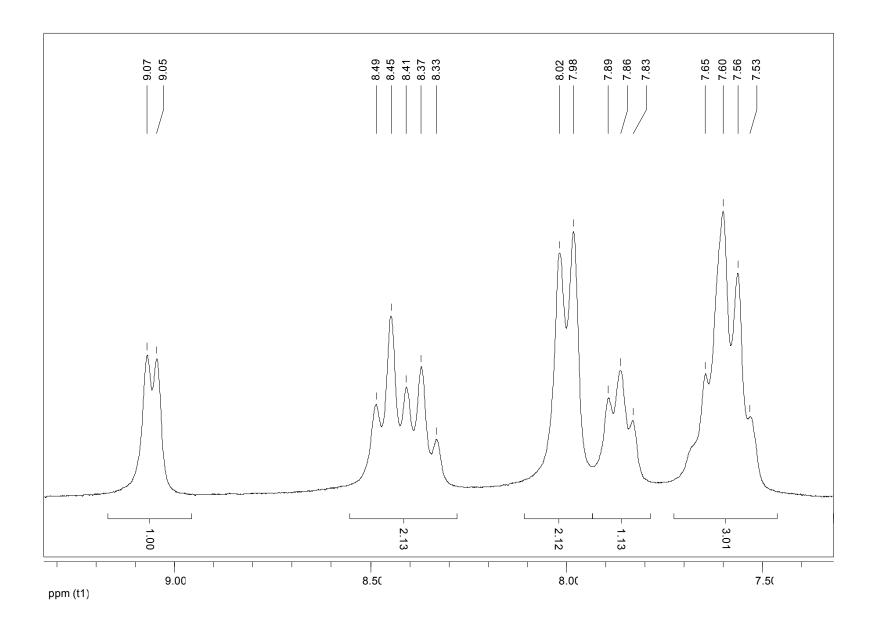
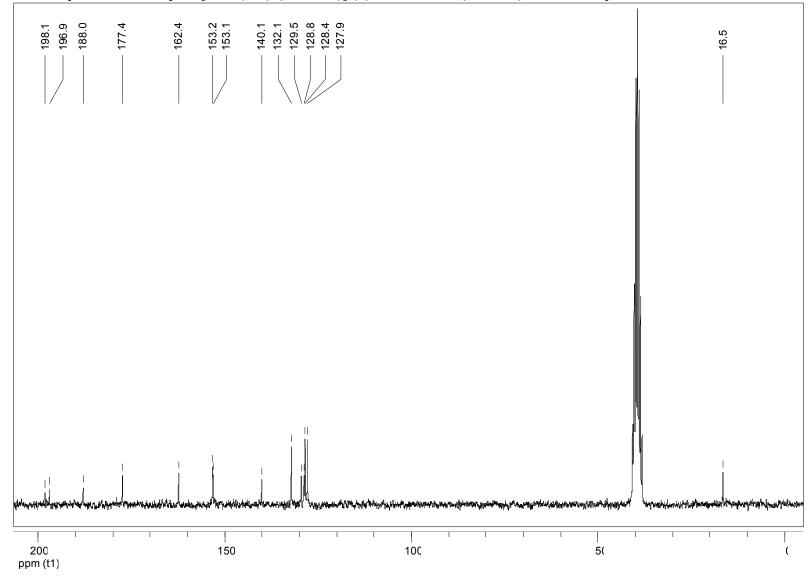
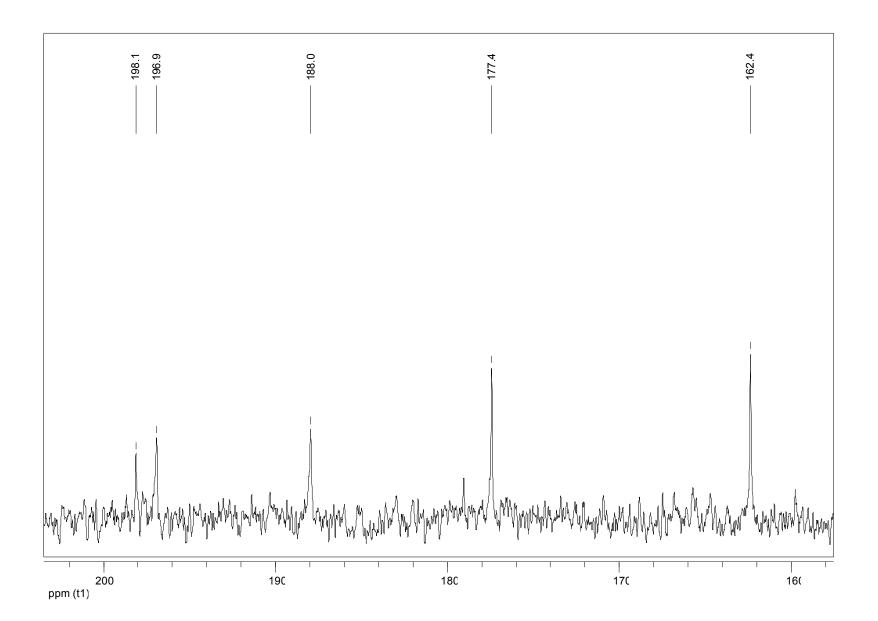
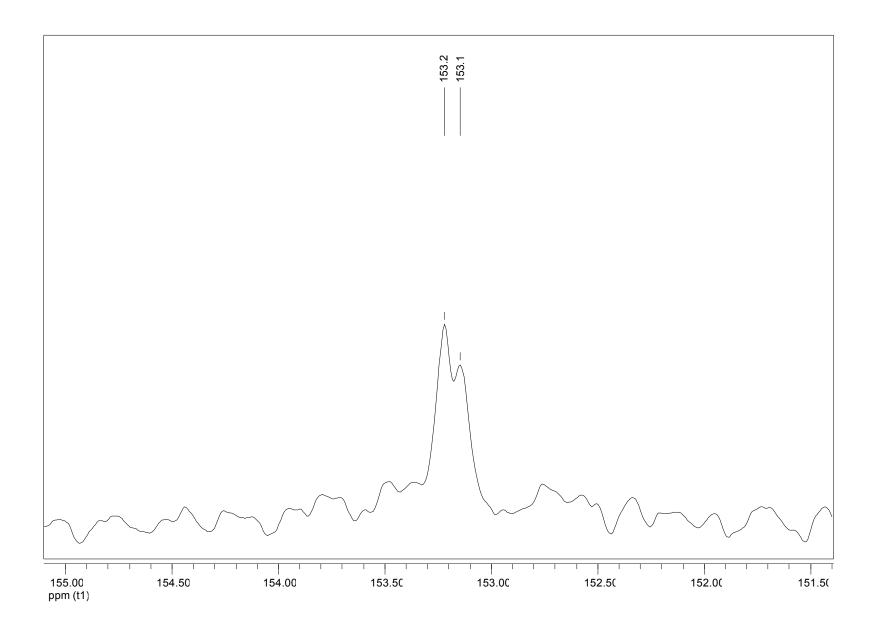
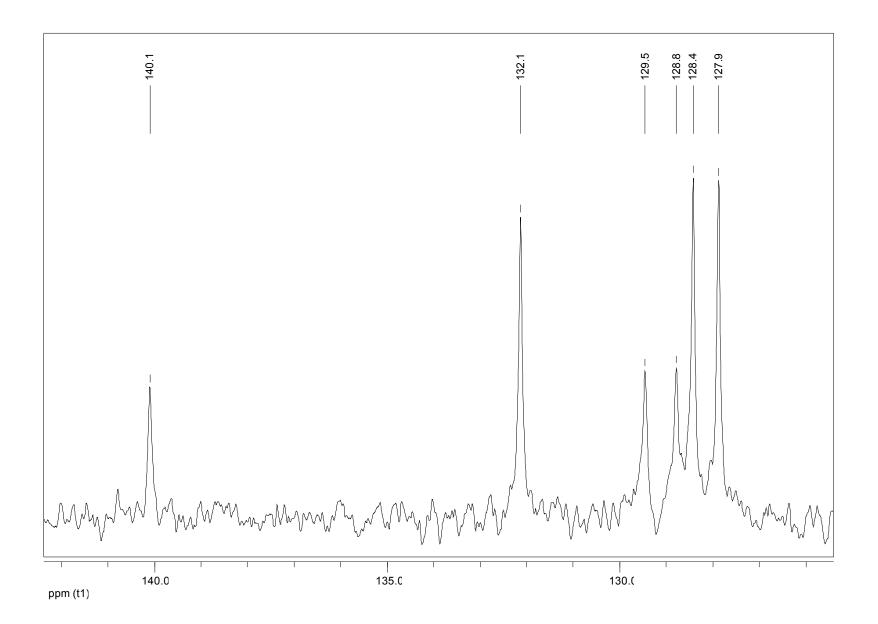


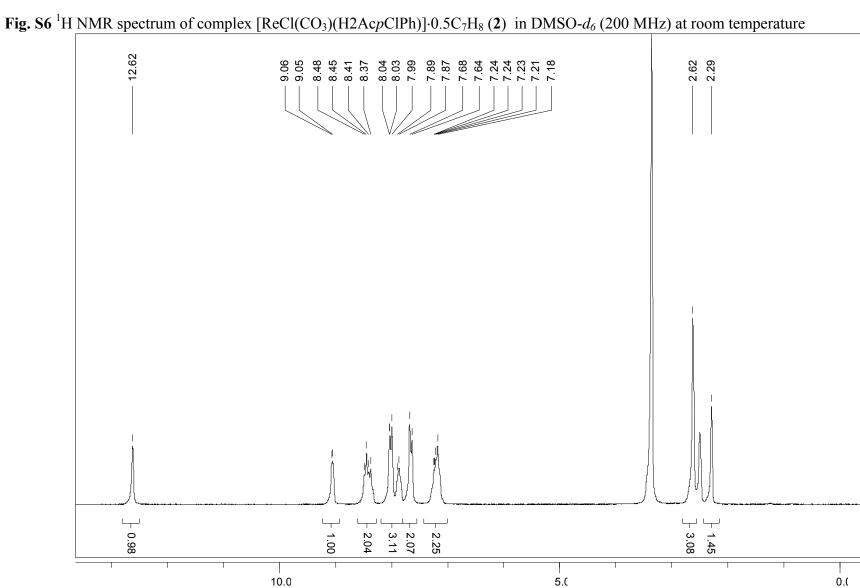
Fig. S5  $^{13}$ C NMR spectrum of complex [ReCl(CO)<sub>3</sub>(H2AcPh)] (1) in DMSO- $d_6$  (50 MHz) at room temperature



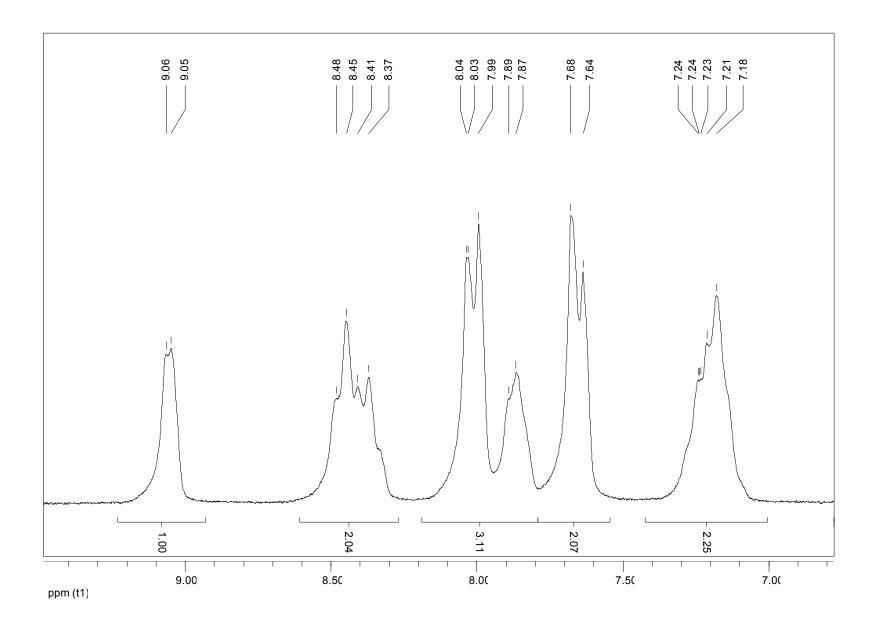


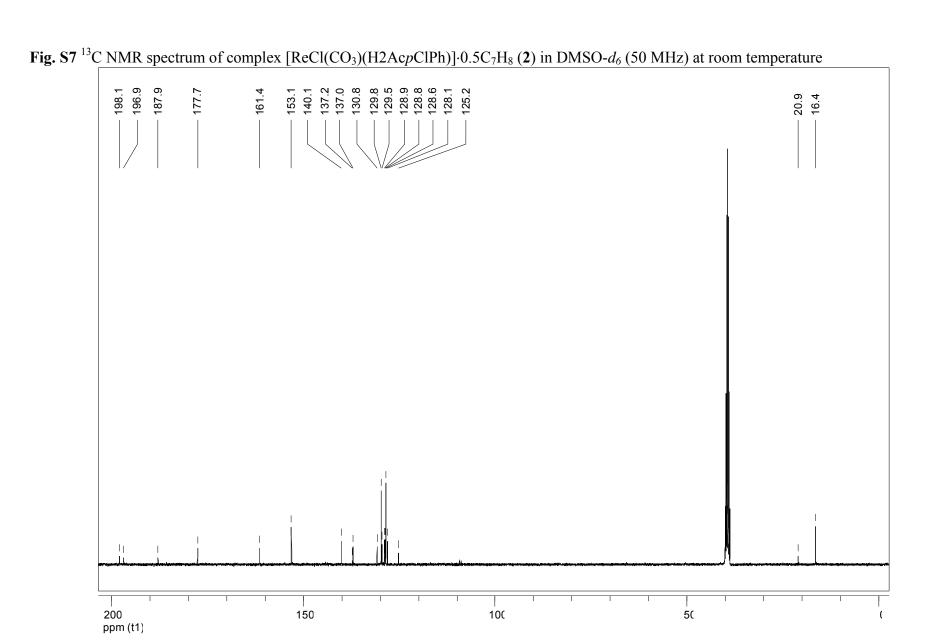


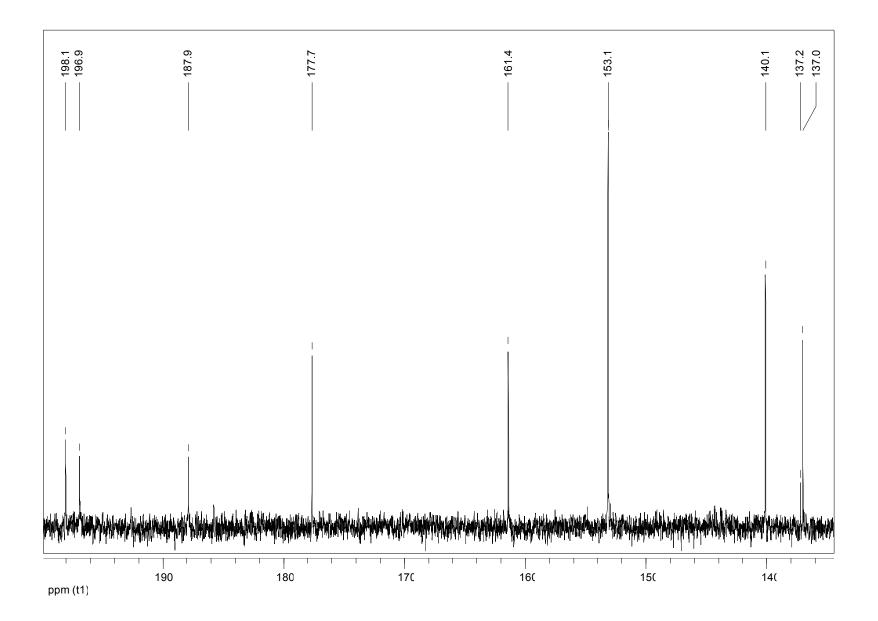




ppm (t1)







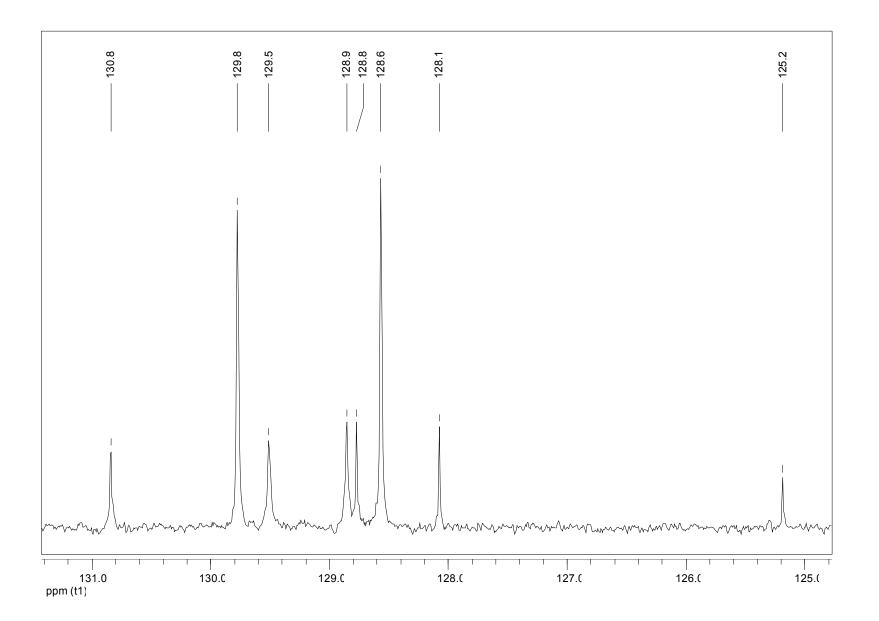


Fig. S8  $^{1}$ H NMR spectrum of complex [ReCl(CO)<sub>3</sub>(H2AcpNO<sub>2</sub>Ph)] $\cdot$ 0.5C<sub>7</sub>H<sub>8</sub> (3) in DMSO- $d_6$  (200 MHz) at room temperature 12.93 2.67 → 1.46→ 3.03 

 → 1.10

 → 2.09

 → 4.11

 <u>}</u> 0.95 10.0 5.0 0.0 ppm (t1)

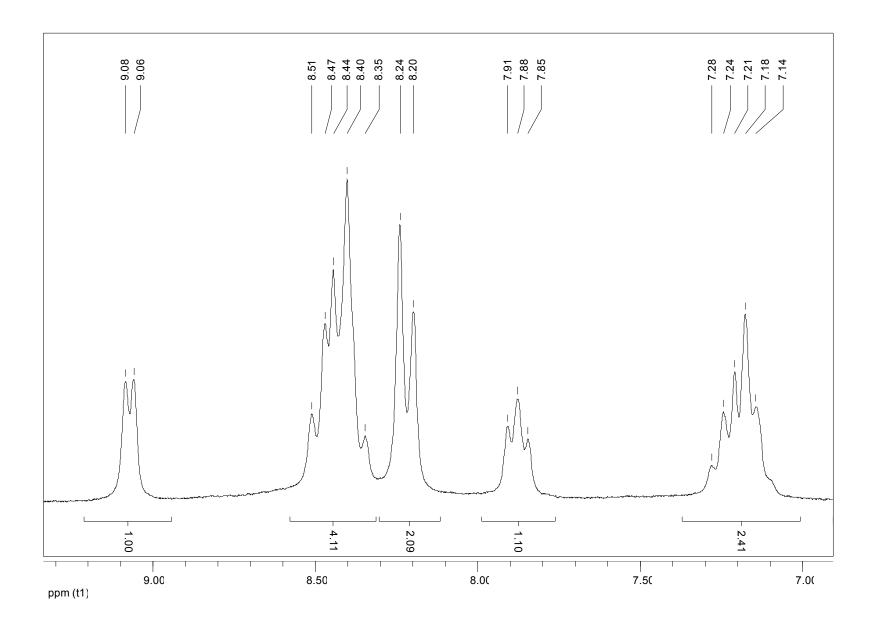
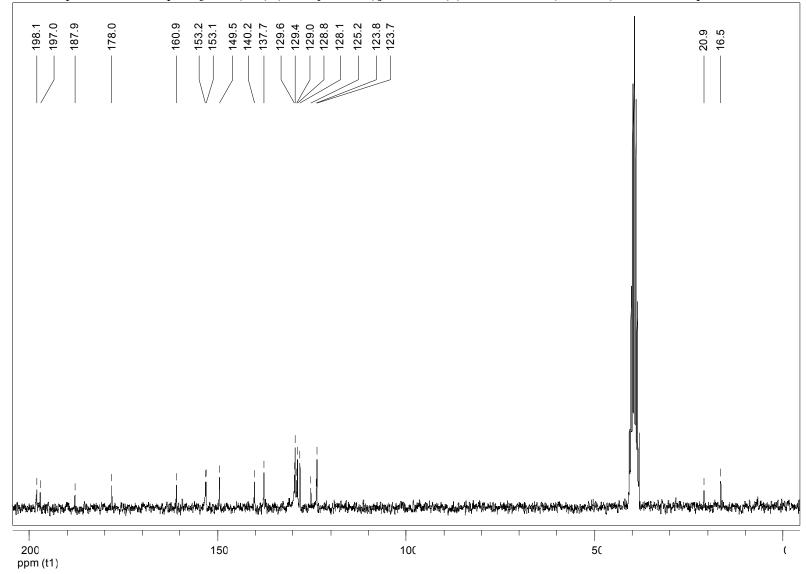
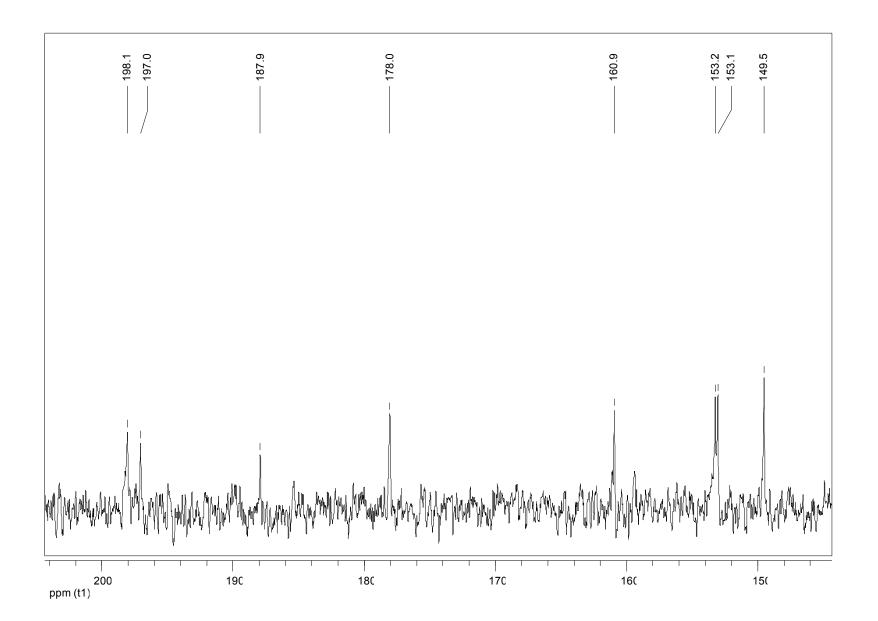
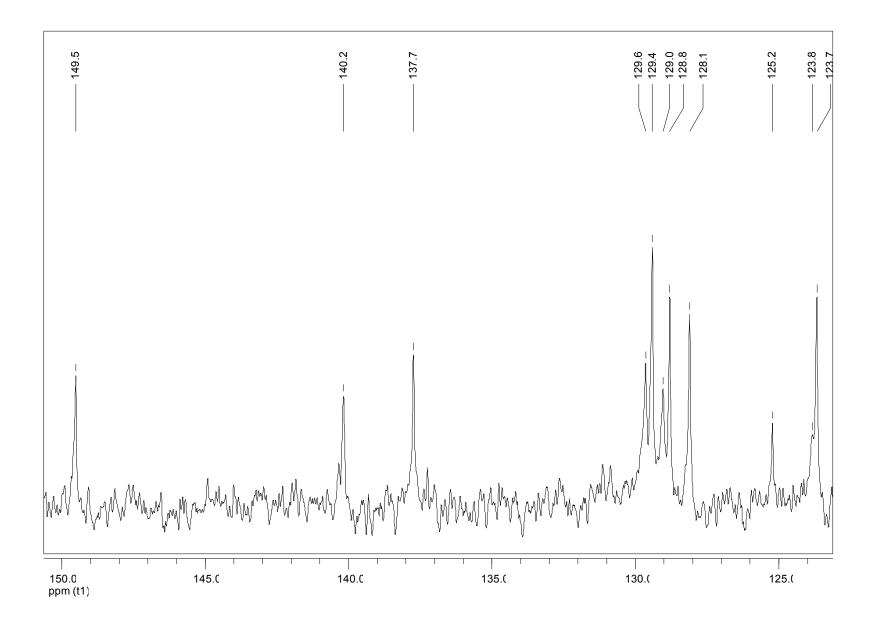


Fig. S9 <sup>13</sup>C NMR spectrum of complex [ReCl(CO)<sub>3</sub>(H2AcpNO<sub>2</sub>Ph)]·0.5C<sub>7</sub>H<sub>8</sub> (3) in DMSO-d<sub>6</sub> (50 MHz) at room temperature







CHECKCIFs files of complexes [ReCl(CO)<sub>3</sub>(H2AcPh)] (1), [ReCl(CO<sub>3</sub>)(H2AcpClPh)]·DMSO (2a) and [ReCl(CO)<sub>3</sub>(H2AcpNO<sub>2</sub>Ph)]·DMSO (3a)

# checkCIF/PLATON report

You have not supplied any structure factors. As a result the full set of tests cannot be run.

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### **Datablock: RePh** Complex (1)

```
Bond precision: C-C = 0.0048 A
                                        Wavelength=0.71073
Cell:
                a=12.6872(2)
                                b=9.5720(2)
                                                  c=14.9830(2)
                alpha=90
                                beta=98.300(2)
                                                   gamma=90
Temperature:
                293 K
               Calculated
                                         Reported
Volume
               1800.51(5)
                                         1800.51(5)
Space group
              P 21/n
                                         P_21/n
Hall group
               -P 2yn
                                         -P_2yn
Moiety formula C17 H13 Cl N3 O4 Re
Sum formula
               C17 H13 Cl N3 O4 Re
                                         C17 H13 Cl N3 O4 Re
Mr
               544.96
                                         544.95
               2.010
                                         2.010
Dx,g cm-3
Ζ
               4
Mu (mm-1)
               6.925
                                         6.925
F000
               1040.0
                                         1040.0
F000′
               1037.09
h,k,lmax
               19,14,22
                                         19,14,22
Nref
               6806
                                         6375
               0.667,0.871
                                         0.534,1.000
Tmin,Tmax
Tmin'
               0.495
Correction method= # Reported T Limits: Tmin=0.534 Tmax=1.000
AbsCorr = MULTI-SCAN
Data completeness= 0.937
                                 Theta(max) = 33.010
R(reflections) = 0.0222(5218) wR2(reflections) = 0.0650(6375)
S = 1.189
                          Npar= 235
```

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 G
PLAT005_ALERT_5_G No Embedded Refinement Details found in the CIF
                                                                                                      Please Do !
PLAT193_ALERT_1_G No s.u.'s on H-positions, Refinement Reported as mixed Check PLAT153_ALERT_1_G The s.u.'s on the Cell Axes are Equal ..(Note) 0.0002 Ang. PLAT180_ALERT_4_G Check Cell Rounding: # of Values Ending with 0 = 3 Note PLAT199_ALERT_1_G Reported _cell_measurement_temperature .... (K) 293 Check PLAT200_ALERT_1_G Reported _diffrn_ambient_temperature .... (K) 293 Check PLAT232_ALERT_2_G Hirshfeld Test Diff (M-X) Re1 -- Cl1
PLAT007_ALERT_5_G Number of Unrefined Donor-H Atoms .....
                                                                                                          1 Report
                                                                                                       mixed Check
                                                                                                          293 Check
                                                                                                        293 Check
PLAT232_ALERT_2_G Hirshfeld Test Diff (M-X) Re1 -- Cl1 ..
                                                                                                         6.5 s.u.
                                                                          -- C16
PLAT232_ALERT_2_G Hirshfeld Test Diff (M-X) Re1
                                                                                                         6.8 s.u.
PLAT232_ALERT_2_G Hirshfeld Test Diff (M-X) Re1 -- C17 ..
                                                                                                         8.1 s.u.
PLAT720_ALERT_4_G Number of Unusual/Non-Standard Labels .....
                                                                                                            1 Note
     0 ALERT level A = Most likely a serious problem - resolve or explain
```

```
O ALERT level A = Most likely a serious problem - resolve or explain
O ALERT level B = A potentially serious problem, consider carefully
O ALERT level C = Check. Ensure it is not caused by an omission or oversight
11 ALERT level G = General information/check it is not something unexpected

4 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
O ALERT type 3 Indicator that the structure quality may be low
2 ALERT type 4 Improvement, methodology, query or suggestion
2 ALERT type 5 Informative message, check
```

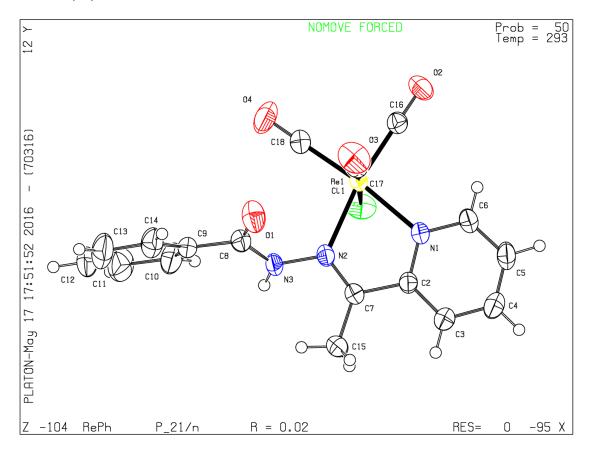
It is advisable to attempt to resolve as many as possible of the alerts in all categories. Often the minor alerts point to easily fixed oversights, errors and omissions in your CIF or refinement strategy, so attention to these fine details can be worthwhile. In order to resolve some of the more serious 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.

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### Datablock: Re\_pCl complex (2a)

```
Bond precision: C-C = 0.0039 A
                                        Wavelength=0.71073
Cell:
              a=8.7687(4)
                                b=11.1002(5)
                                               c=12.9282(5)
                                beta=97.163(4)
              alpha=113.002(4)
                                                 qamma = 93.603(4)
Temperature:
              200 K
               Calculated
                                         Reported
Volume
               1140.66(9)
                                         1140.66(9)
                                         P -1
Space group
              P -1
Hall group
               -P 1
                                         -P 1
               C17 H12 Cl2 N3 O4 Re, C2
                                         C19 H18 Cl2 N3 O5 Re S
Moiety formula
               H6 O S
Sum formula
               C19 H18 Cl2 N3 O5 Re S
                                         C19 H18 Cl2 N3 O5 Re S
Mr
               657.53
                                         657.52
Dx,g cm-3
               1.914
                                         1.914
               2
                                         2
Mu (mm-1)
               5.688
                                         5.688
F000
               636.0
                                         636.0
F000′
               635.10
h,k,lmax
               13,16,19
                                         13,16,19
Nref
               8528
                                         7953
                                         0.555,1.000
Tmin,Tmax
               0.626,0.892
Tmin'
               0.422
Correction method= # Reported T Limits: Tmin=0.555 Tmax=1.000
AbsCorr = MULTI-SCAN
Data completeness= 0.933
                                 Theta(max) = 32.880
R(reflections) = 0.0255(7050) wR2(reflections) = 0.0508(7953)
S = 1.061
                          Npar= 283
```

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 G

```
PLAT005_ALERT_5_G No Embedded Refinement Details found in the CIF Please Do !
PLAT007_ALERT_5_G Number of Unrefined Donor-H Atoms .....
                                                                 1 Report
                                                          1 Report
Please Check
PLAT042_ALERT_1_G Calc. and Reported MoietyFormula Strings Differ
                                                         mixed Check
PLAT093_ALERT_1_G No s.u.'s on H-positions, Refinement Reported as
PLAT154_ALERT_1_G The s.u.'s on the Cell Angles are Equal ..(Note)
                                                            0.004 Degree
PLAT232_ALERT_2_G Hirshfeld Test Diff (M-X) Re1 -- C16 ..
                                                             6.9 s.u.
PLAT232_ALERT_2_G Hirshfeld Test Diff (M-X) Re1
                                            -- C17
                                                              6.0 s.u.
                                                      . .
PLAT232_ALERT_2_G Hirshfeld Test Diff (M-X) Re1 -- C18 ..
                                                              7.3 s.u.
PLAT720_ALERT_4_G Number of Unusual/Non-Standard Labels .....
                                                               1 Note
                                                             4.09 Ang.
PLAT721_ALERT_1_G Bond Calc 4.95000, Rep 0.86000 Dev...
                          1.555 1.555 ..... Bond #
          N3 -H3
                                                           20 Check
PLAT722_ALERT_1_G Angle Calc 144.00, Rep 120.60 Dev...
                                                            23.40 Degree
           C8 -N3 -H3
                            1.555 1.555 1.555 #
                                                           26
PLAT722_ALERT_1_G Angle Calc
                            38.00, Rep 120.60 Dev...
                                                            82.60 Degree
           N2 -N3 -H3
                           1.555 1.555 1.555 #
                                                           27
```

- 0 ALERT level A = Most likely a serious problem resolve or explain
- 0 ALERT level B = A potentially serious problem, consider carefully
- 0 ALERT level C = Check. Ensure it is not caused by an omission or oversight
- 12 ALERT level G = General information/check it is not something unexpected
- 6 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
- $\ensuremath{\text{0}}$  ALERT type  $\ensuremath{\text{3}}$  Indicator that the structure quality may be low
- 1 ALERT type 4 Improvement, methodology, query or suggestion
- 2 ALERT type 5 Informative message, check

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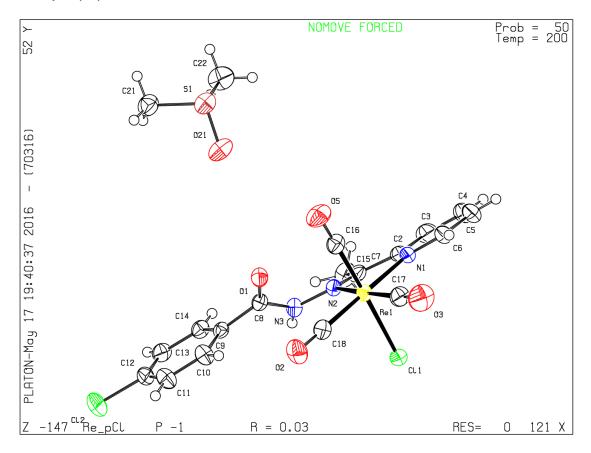
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# Datablock: RepNO2 complex (3a)

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                                 b=11.1200(6)
                                               c=12.9995(6)
              alpha=112.409(4)
                                 beta=96.989(4)
                                                 qamma = 93.490(4)
Temperature: 293 K
               Calculated
                                         Reported
Volume
               1172.08(11)
                                         1172.08(11)
                                         P - 1
Space group
              P -1
Hall group
               -P 1
                                         -P1
               C17 H12 Cl N4 O6 Re, C2 H6 _{3}
Moiety formula
               0 S
Sum formula
               C19 H18 Cl N4 O7 Re S
                                         C19 H18 Cl N4 O7 Re S
Mr
               668.09
                                         668.08
Dx,g cm-3
               1.893
                                         1.893
               2
                                         2
               5.434
Mu (mm-1)
                                         5.434
               648.0
F000
                                         648.0
F000′
               646.83
               12,15,18
                                         11,15,17
h,k,lmax
Nref
               6568
                                         5446
                                         0.669,1.000
Tmin,Tmax
               0.729,0.897
Tmin'
               0.575
Correction method= # Reported T Limits: Tmin=0.669 Tmax=1.000
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Data completeness= 0.829
                                 Theta(max) = 29.550
R(reflections) = 0.0297( 4833) wR2(reflections) = 0.0715( 5446)
S = 1.082
                          Npar= 300
```

# 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' Ueq as Compared to Neighbors of	N4	Check								
PLAT751_ALERT_4_C Bond Calc 0.93000, Rep 0.927(4) N3 -HN2 1.555 1.555 Bond #		s.u.								
PLAT752_ALERT_4_C Angle Calc 117.00, Rep 117.3(3) C8 -N3 -HN2 1.555 1.555 #	Senseless 26	s.u.								
PLAT752_ALERT_4_C Angle Calc 123.00, Rep 122.5(3) N2 -N3 -HN2 1.555 1.555 #	Senseless 27	s.u.								
Alert level G										
PLAT005_ALERT_5_G No Embedded Refinement Details found in the CIF	Please	Do !								
PLAT007_ALERT_5_G Number of Unrefined Donor-H Atoms	1	Report								
PLAT093_ALERT_1_G No s.u.'s on H-positions, Refinement Reported as	mixed	Check								
PLAT154_ALERT_1_G The s.u.'s on the Cell Angles are Equal(Note)	0.004	Degree								
PLAT199_ALERT_1_G Reported _cell_measurement_temperature (K)	293	Check								
PLAT200_ALERT_1_G Reporteddiffrn_ambient_temperature (K)	293	Check								
PLAT380_ALERT_4_G Incorrectly? Oriented X(sp2)-Methyl Moiety	C15	Check								
PLAT720_ALERT_4_G Number of Unusual/Non-Standard Labels										
0 ALERT level A = Most likely a serious problem - resolve or exp	olain									
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4 ALERT level C = Check. Ensure it is not caused by an omission or oversight										
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1 ALERT type 2 Indicator that the structure model may be wrong or deficient										
O ALERT type 3 Indicator that the structure quality may be low										

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