

Acidity Quantification and Structure Analysis of Amide-AlCl₃ Liquid Coordination Complexes for C₄ Alkylation Catalysis

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1. Gutmann-Beckett values and computed FIA values.

Table S1. Collection of literature known and measured Gutmann-Beckett values ($\Delta\delta_{31\text{P}}^{\text{exp}}$) [23].

Lewis Acid	$\Delta\delta_{31\text{P}}^{\text{exp}}$ [ppm]
BH ₃	33.1
BF ₃	29.0
BCl ₃	38.7
BBr ₃	40.3
BI ₃	42.9
BEt ₃	1.9
BPh ₃	20.6
B(C ₆ F ₅) ₃	27.0
B(C ₆ F ₅) ₂ (C ₆ Cl ₅)	25.8
B(C ₆ F ₄ - <i>o</i> -C ₆ F ₅) ₃	30.7
B(NMe ₂) ₃	-4.9
B(<i>Oi</i> Pr) ₃	0.8
B(OMe) ₃	-1.9
B(OC ₆ F ₅) ₃	30.9
B(OC ₆ H ₅) ₃	19.4
Cl-B(cat)	30.5
Cl-B(nad)	31.5
Ph-B(cat)	19.2
Ph-B(nad)	17.9
C ₆ F ₅ -B(cat)	26.6
C ₆ F ₅ -B(nad)	27.5
Me ₃ -B(cat)	-3.5
Me ₃ -B(nad)	-3.7
F ₂ -B(cat)	23.9
F ₂ -B(nad)	23.6
F ₃ -B(cat)	24.9
F ₃ -B(nad)	23.8
Cl ₂ -B(cat)	20.2
Cl ₂ -B(nad)	15.2
9-B(Trypt)	25.0
AlCl ₃	32.2
AlBr ₃	33.1 / 35.4
AlI ₃	30.8 / 37.4
AlMe ₃	16.7
AlEt ₃	15.9
Al(C ₆ F ₅) ₃	26.0

AlN(OC ₇ H ₄) ₃ Cl	21.2
AlN(OC ₇ H ₄) ₃ Me	19.1
[AlCalixMe]-	18.5
Ga(C ₆ F ₅) ₃	23.1
GaN(OC ₇ H ₄) ₃ Cl	22.1
GaN(OC ₇ H ₄) ₃ Me	19.6
GaCl ₃	31.1
GaI ₃	28.3 / 30.2
InCl ₃	1.5 / 13.7
InBr ₃	3.2 / 20.3
In(C ₆ F ₅) ₃	19.4
SiCl ₄	25.1 / 25.0
SiBr ₄	37.3 / 37.4
SiI ₄	58.1 / 58.2
1-Si(catH) ₂	33.2
2-Si(catH) ₂	17.7
1-Si(catF) ₂	36.6
2-Si(catF) ₂	22.2
1-Si(catCl) ₂	37.2
2-Si(catCl) ₂	23.1
2-Si(catBr) ₂	37.3
2-Si(catBr) ₂	23.1
1-Si(cattBu) ₂	31.6
2-Si(cattBu) ₂	20.5
1-Si(catCF ₃) ₂	39.4
2-Si(catCF ₃) ₂	24.7
[SiEt ₃] ⁺	38.6
[Si(C ₆ Me ₆) ₃] ⁺	35.4
[Si(Xylyl) ₃] ⁺	38.5
[Si(Tipp) ₃] ⁺	41.1
[Si(Tipp) ₂ Et] ⁺	41.3
[Si(NMe ₂) ₃] ⁺	35.2
2-GeF ₄	26.4
2-GeCl ₄	10.1
1-Ge(catCl) ₂	33.7
2-Ge(catCl) ₂	20.6
[Ge(Tipp) ₂ Me] ⁺	36.6
[NBn-BZIMPYGeCl] ⁺	41.9
SnCl ₄	17.4
SnI ₄	17.9 / 17.7
[Sn(iPr) ₃] ⁺	21.7
[Sn(nBu) ₃] ⁺	20.0
SnCalixEt	19.3
[NBn-BZIMPYSn] ₂ ⁺	26.5
[Ph-CAAN] ⁺	6.5
[P(C ₆ F ₅) ₃ F] ⁺	41.1
[O=P-CDipp] ⁺	6.3
[O=P-NDipp] ⁺	37.2
[Ph-P(bipy)] ₂ ⁺	34.0
[Ph-P(terpy)] ₂ ⁺	5.0
[(terpy)PPh] ₂ ⁺	5.8
[P(η ⁵ -Cp [*])] ₂ ⁺	64.7
AsCl ₃	3.4 / 7.6

AsI ₃	13.1 / 22.0
[As(η^5 -Cp [*])] ₂ ⁺	49.6
SbF ₃	7.2 / 13.0
SbCl ₅	36.1
SbCl ₃	18.1 / 21.2
Sb(Mesityl) ₂ OTf	31.6
Sb(Mesityl) ₂ Cl	1.0
Sb-NNN	-2.2
[Sb(η^5 -Cp [*])] ₂ ⁺	39.0
BiCl ₃	16.0 / 25.9
BiBr ₃	11.5 / 24.3
Bi(Mesityl) ₂ OTf	14.6
Bi(Mesityl) ₂ Cl	5.3
Bi-NNN	3.6
[S(OFPh ₂)] ⁺	34.5
[F-I(Ph)] ⁺	40.3
[Cl-I(Ph)] ⁺	35.6
[C ₆ F ₅ -I(Ph)] ⁺	13.1
ZnF ₂	0.6 / 0.5
ZnCl ₂	18.6 / 20.4
ZnBr ₂	18.5 / 20.2
ZnI ₂	18.3 / 18.3
Zn(C ₆ F ₅) ₂	17.7
[ZnCDipp(C ₆ F ₅)] ⁺	25.2
[ZnCDippMe] ⁺	10.0
TiCl ₄	22.7

Table S2. Computed FIA values of Lewis acids given in kJ/mol⁻¹ [23] (computed on DLPNO-CCSD(T)/aug-cc-pVQZ level of theory according to Ref. [24]).

Lewis Acid	FIA
BEt ₃	264
B(OMe) ₃	215
B(OiPr) ₃	220
B(NMe ₂) ₃	168
B(OC ₆ H ₅) ₃	323
B(C ₆ F ₅) ₂ (C ₆ Cl ₅)	422
B(C ₆ F ₄ -o-C ₆ F ₅) ₃	449
Cl-B(cat)	352
Cl-B(nad)	373
Ph-B(cat)	424
Ph-B(nad)	355
C ₆ F ₅ -B(cat)	378
C ₆ F ₅ -B(nad)	387
Me ₃ -B(cat)	310
Me ₃ -B(nad)	308
F ₂ -B(cat)	357
F ₂ -B(nad)	423
F ₃ -B(cat)	417
F ₃ -B(nad)	443
Cl ₂ -B(cat)	368
Cl ₂ -B(nad)	348
⁹ -B(Trypt)	444

AlEt ₃	379
AlN(OC ₇ H ₄) ₃	412
AlN(OC ₇ H ₄) ₃	363
[AlCalixMe]-	132
GaN(OC ₇ H ₄) ₃	351
GaN(OC ₇ H ₄) ₃	291
InCl ₃	426
InBr ₃	419
In(C ₆ F ₅) ₃	420
2-SiCl ₄	-28
2-SiBr ₄	17
2-SiI ₄	60
1-Si(catH) ₂	395
2-Si(catH) ₂	-93
1-Si(catF) ₂	483
2-Si(catF) ₂	9
1-Si(catCl) ₂	490
2-Si(catCl) ₂	42
1-Si(catBr) ₂	469
2-Si(catBr) ₂	52
1-Si(cattBu) ₂	367
2-Si(cattBu) ₂	-172
1-Si(catCF ₃) ₂	570
2-Si(catCF ₃) ₂	127
1-SiCalixEt	484
2-SiCalixEt	56
[SiEt ₃] ⁺	934
[Si(C ₆ Me ₆) ₃] ⁺	782
[Si(Xylyl) ₃] ⁺	838
[Si(Tipp) ₃] ⁺	732
[Si(Tipp) ₂ Et] ⁺	746
[Si(NMe ₂) ₃] ⁺	829
1-Si(cattBu) ₂	367
2-Si(cattBu) ₂	-172
1-Si(catCF ₃) ₂	570
2-Si(catCF ₃) ₂	127
1-SiCalixEt	484
2-SiCalixEt	56
[SiEt ₃] ⁺	934
[Si(C ₆ Me ₆) ₃] ⁺	782
[Si(Xylyl) ₃] ⁺	838
[Si(Tipp) ₃] ⁺	732
[Si(Tipp) ₂ Et] ⁺	746
[Si(NMe ₂) ₃] ⁺	829
2-GeF ₄	-142
2-GeCl ₄	-47
1-Ge(catCl) ₂	490
2-Ge(catCl) ₂	60
[Ge(Tipp) ₂ Me] ⁺	753
[NBn-BZIMPYGeCl] ⁺ 549	120
2-SnCl ₄	-17
2-SnI ₄	17
[Sn(iPr) ₃] ⁺	786

[Sn(nBu) ₃] ⁺	893
1-SnCalixEt	430
2-SnCalixEt	79
[NBn-BZIMPY ₂ Sn] ²⁺	938
PI ₃	298
[P(C ₆ F ₅) ₃ F] ⁺	723
[O=P-CDipp] ⁺	628
[O=P-NDipp] ⁺	666
P-NNN	302
[Ph-P(bipy)] ₂ ⁺	815
[Ph-P(terpy)] ₂ ⁺	872
[(terpy)PPh] ₂ ⁺	796
[P(η ⁵ -Cp*)] ₂ ⁺	1167
AsI ₃	289
As-NNN	313
[As(η ⁵ -Cp*)] ₂ ⁺	1123
Sb(Mesityl) ₂ Cl	241
Sb(Mesityl) ₂ OTf	309
Sb-NNN	386
BiCl ₃	320
BiBr ₃	338
Bi(Mesityl) ₂ Cl	257
Bi(Mesityl) ₂ OTf	282
Bi-NNN	311
[S(FOPh ₂)] ⁺	616
[S(FObiph)] ⁺	611
[F-I(Ph)]	776
[Cl-I(Ph)] ⁺	726
[C ₆ F ₅ -I(Ph)] ⁺	610
ZnF ₂	442
ZnCl ₂	357
ZnBr ₂	442
ZnI ₂	703
Zn(C ₆ F ₅) ₂	348
[ZnCDipp(C ₆ F ₅)] ⁺	638
[ZnCDippMe] ⁺	582
[ZnCDippEt] ⁺	582
TiCl ₄	384

2. GB-FIA model and correlation program

2.1. GB-FIA model and hypothesis test

To determine whether a new point (e.g., 400, 19.9) falls within the relevance curve established by GB-FIA model, a linear regression model can be used for fitting and hypothesis testing.

Step 1: Hypothesis

First, we need to clarify the hypothesis.

Null hypothesis (H₀): The new point (e.g., 400, 19.9) is on the relevance curve established by GB-FIA.

Alternative hypothesis (H₁): The new point (e.g., 400, 19.9) is not on the relevance curve established by GB-FIA.

Step 2: Regression Model

With the known data points FIA and GB, a linear regression model can be fitted to describe their relationship. This can be done using the least squares method.

Step 3: Calculate Residuals

After fitting the model, calculate the residuals for each data point on the fit curve (i.e., the difference between the actual observed values and the predicted values from the model). Residuals indicate the degree of deviation of each data point relative to the fit line.

Step 4: Hypothesis Test

Use the fitted model to calculate the predicted value of the new point (400, 19.9) and its residual. Then, perform hypothesis testing using the distribution of these residuals.

The key in hypothesis testing is calculating the confidence interval, which is a range containing the probable estimates of the true parameter. If the residual of the new point falls within this confidence interval, we cannot reject the null hypothesis.

Step 5: Determine

Compare the residual of the new point with the confidence interval. If the residual of the new point falls within the confidence interval, the null hypothesis cannot be rejected, implying that the new point could be on the relevance curve established by GB-FIA. If the residual falls outside the confidence interval, reject the null hypothesis, implying that the new point is not on the relevance curve established by GB-FIA.

2.2. Matlab program for establishing the GB-FIA model and hypothesis test

The Matlab program attached after this section is used to establish the GB-FIA regression model. The data used for modeling is from section 2.1, but there are some slight differences in the correlation between GB and FIA with these data. When the dataset is multi-compound dataset (large dataset), the correlation coefficient R^2 between GB and FIA is 0.923. In contrast, when the dataset only includes AI compounds, the correlation coefficient R^2 between GB and FIA is as high as 0.990. This indicates a strong correlation between the experimental GB of AI compounds, the calculated FIA, and the molecular model constructed for FIA calculations.

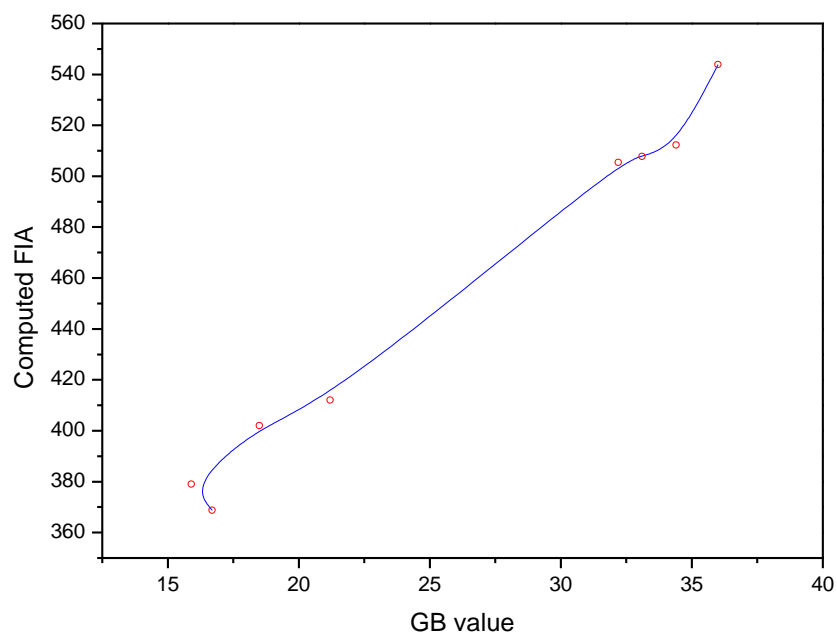


Figure S1. The correlation model established by AI-compound GB-FIA database.

Matlab program for establishing the GB-FIA model and hypothesis test:

```
function calfia
FIA=[368.8 379 402 412 505.4 507.8 512.2 543.8]';
GB=[16.7 15.9 18.5 21.2 32.2 33.1 34.4 36]';
n = length(GB);
p=polyfit(GB,FIA,2)
beta00=p(3)
```

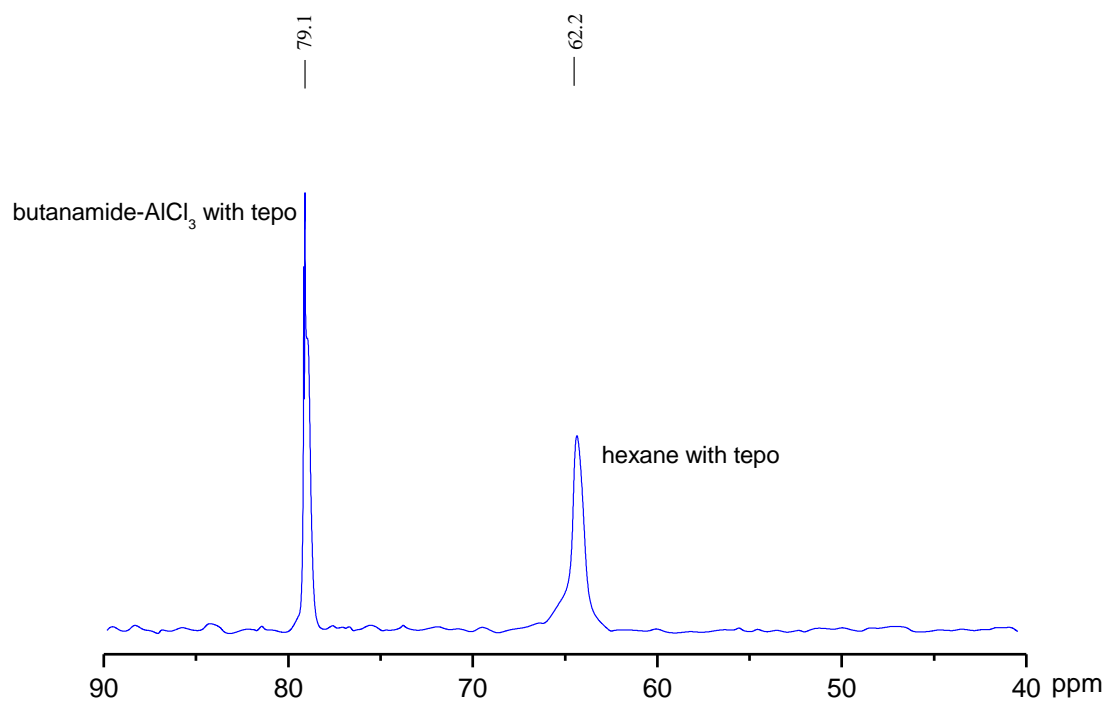
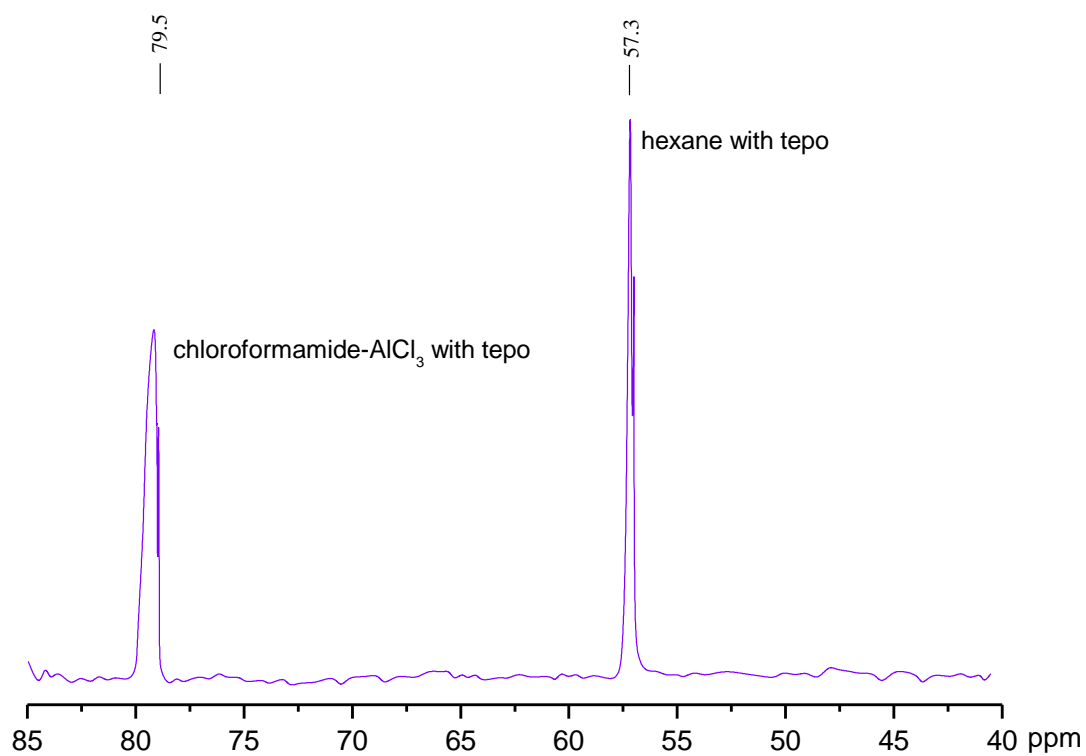
```

beta1=p(2)
beta2=p(1)
X = [ones(n, 1), GB, GB.^2]
Y = FIA
beta = (X' * X) \ (X' * Y) % Calculate the regression coefficients.
nbeta=inv(X'*X)*X'*Y
beta0 = beta(1)
beta1 = beta(2)
beta2 = beta(3)
predicted_FIA = beta0 + beta1 * GB + beta2 * GB.^2; % Calculate the predicted value
residuals = FIA - predicted_FIA; % Calculate the residuals.
new_FIA = 400;
% new_GB = 19.9;
new_GB=[23.1 34.5 34.8 35.0 35.3 35.7 37.0];
predicted_FIA_new = beta0 + beta1 * new_GB + beta2 * new_GB.^2; % Calculate the
predicted value of the new data point.
[M,N]=size(new_GB)
for i=1:N
    if predicted_FIA_new(i) >500
        predicted_FIA_new(i)=predicted_FIA_new(i)+70;
    end
end
predicted_FIA_new'
residual_new = new_FIA - predicted_FIA_new % Calculate the residual of the new
data point.
alpha = 0.05; % Confidence level.
residual_std = std(residuals); % Calculate the standard deviation of residuals.
z_critical = norminv(1 - alpha/2) % Critical values for a normal distribution.
confidence_interval = [predicted_FIA_new - z_critical * residual_std, pre-
dicted_FIA_new + z_critical * residual_std] % Calculate the confidence interval.

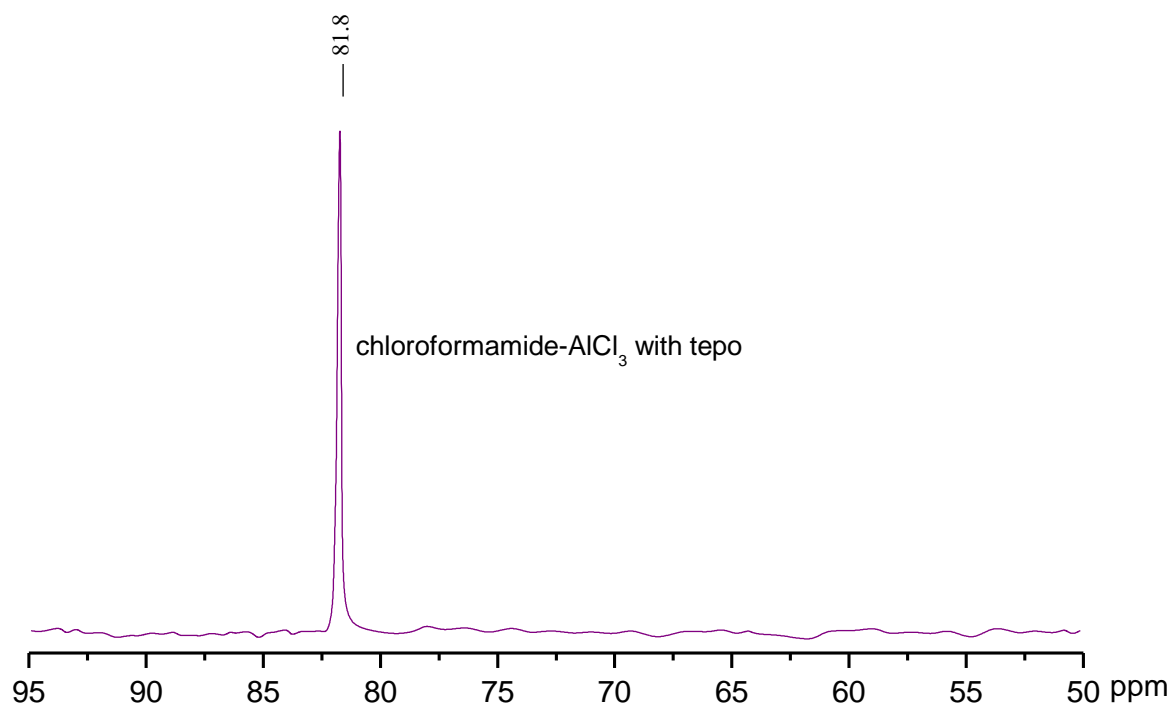
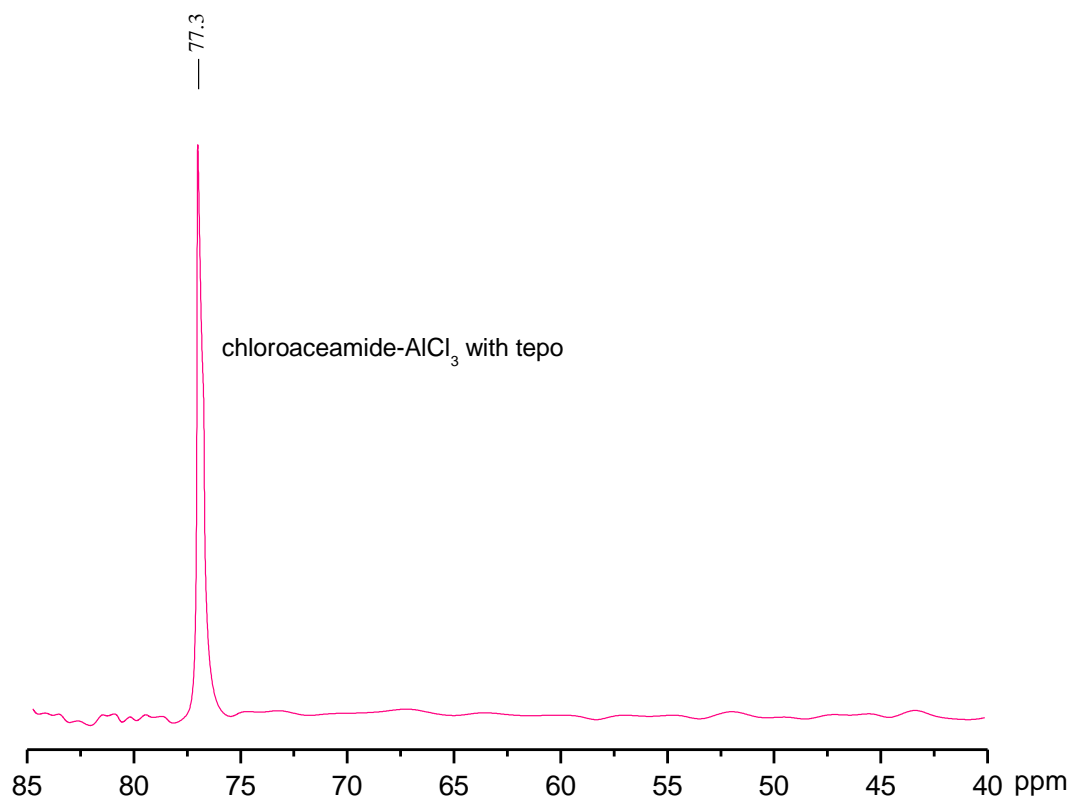
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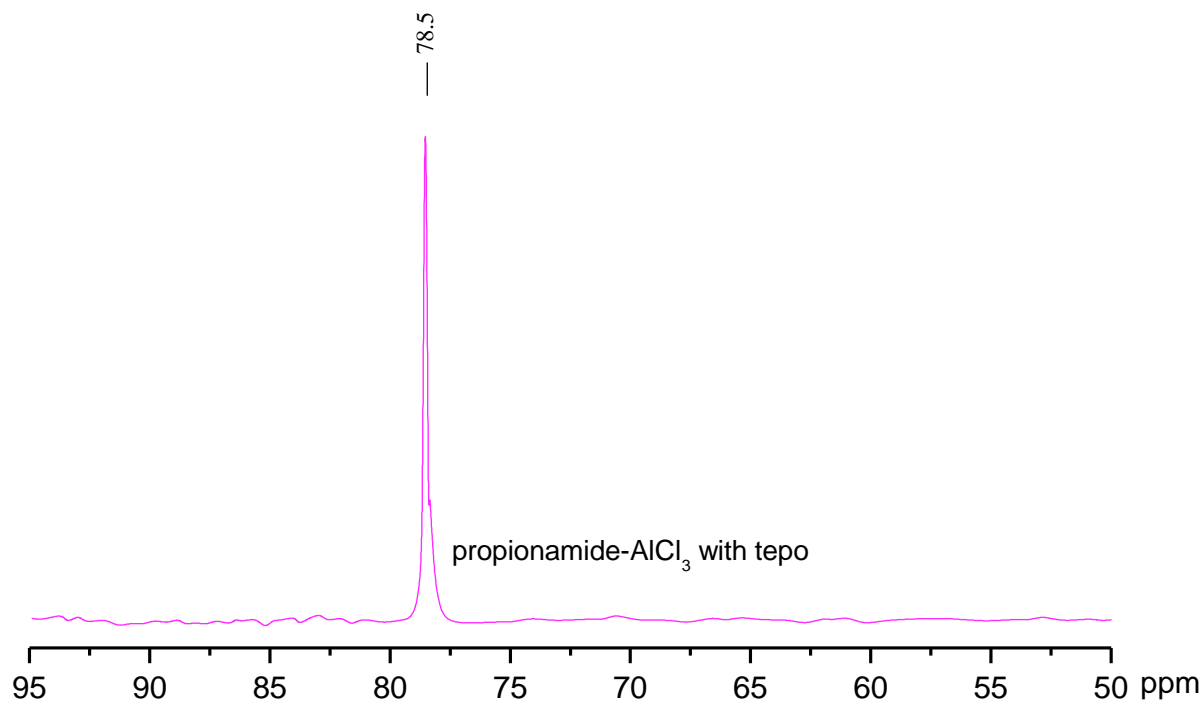
3.31. P NMR spectra and GC chromatogram

3.1.31. P NMR spectra for AN calculation



3.2.31. ^{31}P NMR spectra for GB value calculation





3.3. Typical GC chromatogram of the alkylate produced by acetamide- AlCl_3 LC

