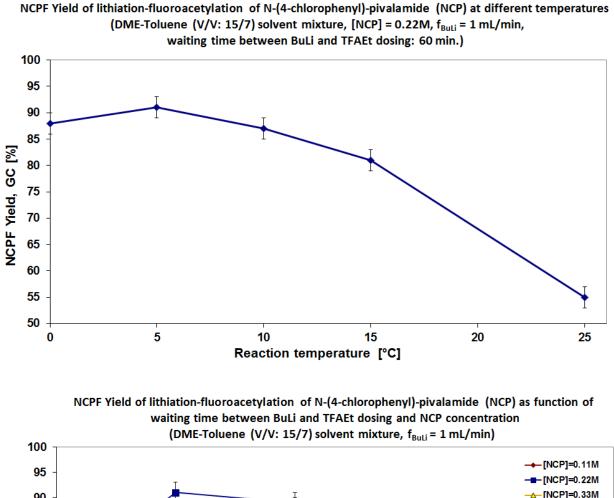
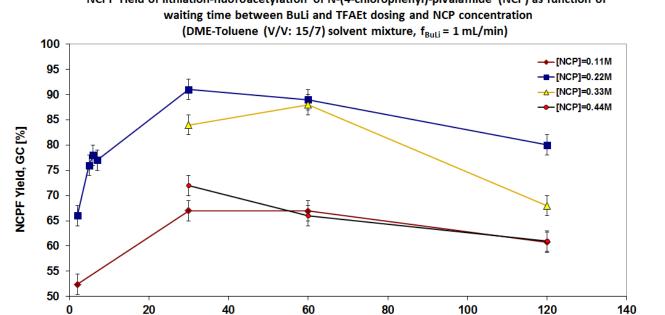
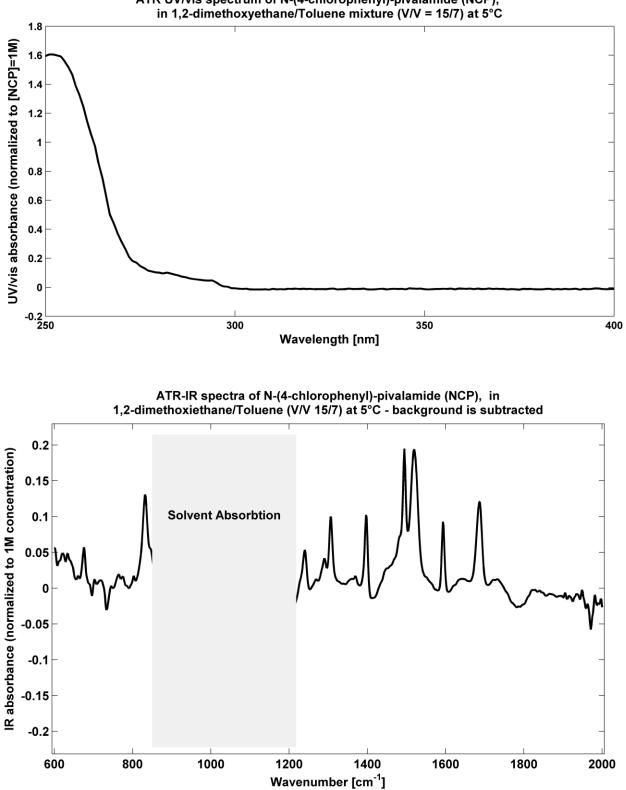
S.1 Optimization results



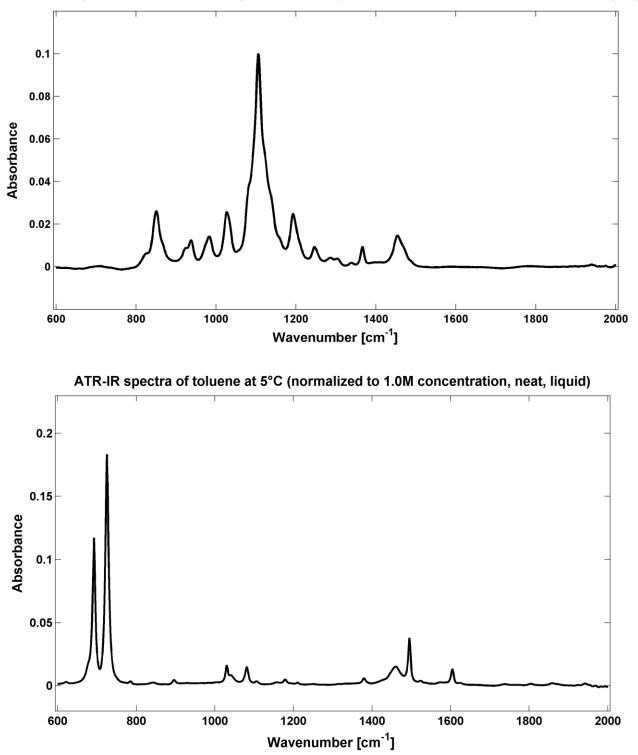


Waiting time between BuLi and TFAEt dosing [min]

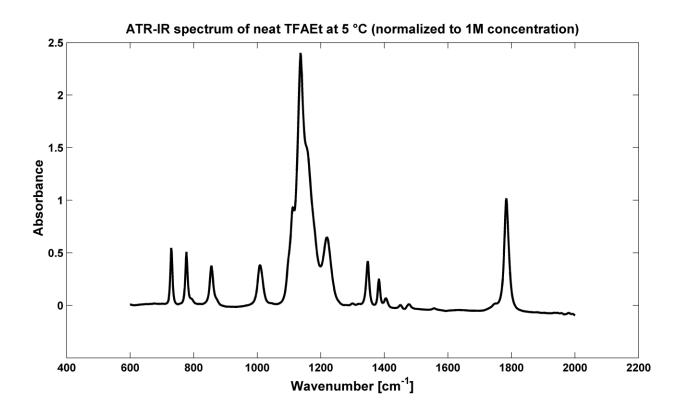
S.2: reference spectra



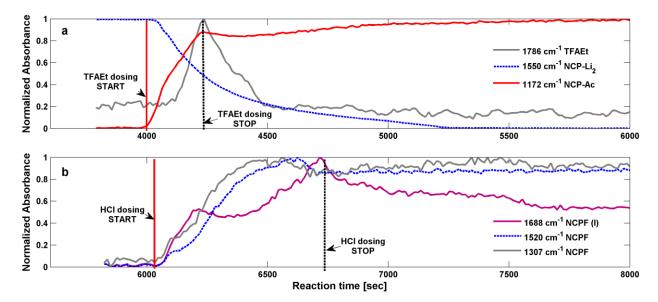
ATR UV/vis spectrum of N-(4-chlorophenyl)-pivalamide (NCP),



ATR-IR spectrum of 1,2-dimethoxyethane at 5°C (normalized to 1.0M concentration, neat, liquid)



S.3 IR absorbance profiles recorded during fluoroacetylation (a) and hydrolysis (b) latter indicating the NCPF product formation:



S.4 Sample Matlab code for fitting first order reaction model for the fluoroacetylation step:

Model:

 $[\text{NCP-Li}_2] = [\text{NCP-Li}_2]_0 \cdot e^{-k \cdot t} = > \text{Abs}_{\text{NCP-Li}_2} = (\text{baseline}) + \text{Abs}_{\text{NCP-Li}_2,0} \cdot e^{-k \cdot t}$

Where:

- k is the apparent rate constant
- [NCP-Li₂] is the NCP-Li₂ concentration at t seconds after the start of TFAEt dosing,
- $[NCP-Li_2]_0$ is the (initial) NCP-Li_2 concentration, at the start of TFAEt dosing,
- Abs_{NCP-Li₂} is the absorbance related to NCP-Li₂, measured at time t at IR-bands IIb, III, V, VI and IX (Table 1)
- Abs_{NCP-Li₂,0} is the (initial) absorbance related to NCP-Li₂, measured at IR-bands IIb, III, V, VI and IX (Table 1), at the start of TFAEt dosing,

Matlab:

1. Define objective function:

```
function ObjectiveFunction = fit_simp(x,X,Y)
A=x(1);
B=x(2);
C=x(3);
ObjectiveFunction = A + B.*exp(C.*X) - Y;
```

2. Fit and visualize data:

```
% Initialize coefficients of the function fit_simp:
X0=[mean(irmat2(i,:)) (max(irmat2(i,:))-min(irmat2(i,:))) -
2/(max(t) - min(t))]';
% Calculate the new coefficients using LSQNONLIN:
options = optimset('Largescale','off');
x=lsqnonlin(@fit_simp,X0,[],[],options,X,Y);
% Plot the original and experimental data.
y_new=fit_simp(x,X,Y)+Y;
plot(X,Y,'or',X,y_new,'b')
%Apparent first order rate constant:
k=-x(3);
```

S.5 Yield calculation from the GC/FID data via internal standard method

	Solution #1		Solution #2		Solution #3	
	mass [mg]*	Area (GC) [pA·s]	mass [mg]*	Area (GC) [pA·s]	mass [mg]*	Area (GC) [pA·s]
NCP	174	6784	520	16762	345	11408
NCPF	503	10501	200	4219	331	6955
Dibenzyl	104	5284	41	2106	19	948
Solution #4			Solution #5			
	mass [mg]*	Area (GC) [pA·s]	mass [mg]*	Area (GC [pA⋅s]	C)	
NCP	0	1334	489	15581		
NCPF	497	10950	0	0		
Dibenzyl	49.8	2659	23	1172		

Five standard solutions of NCP and NCPF in 10 mL DME/Toluene (V/V=15/7) mixture were measured using GC/FID for calibration, the solutions were:

* added weight, the NCPF standard contains about 10% NCP

S.5/a determination of NCP content of the NCPF standard via standard addition.

As seen from the GC/FID results for solution #4 (no added NCP), the NCPF standard contains a significant amount of NCP. Therefore the amount of NCP impurity has to be determined, based on the standard addition method (equations 1.1 - 1.3).

For a choosen solution "i":

$$\frac{A_{\text{NCP, i}}}{A_{\text{NCPF, i}}} = \frac{y_{\text{NCP}}}{y_{\text{NCPF}}} \cdot \frac{c_{\text{NCP, i}}}{c_{\text{NCPF, i}}} = y_{\text{rel}} \cdot \frac{M_{\text{NCPF}} \cdot m_{\text{NCP, i}}^{\text{total}}}{M_{\text{NCP}} \cdot m_{\text{NCPF, i}}^{\text{total}}} = y_{\text{rel}} \cdot \frac{M_{\text{NCPF}} \cdot (m_{\text{NCP, i}} + r \cdot m_{\text{NCPF, i}})}{M_{\text{NCPF, i}}(1-r)} \quad 1.1$$

For an other choosen solution "j":

$$\frac{A_{\text{NCP,}j}}{A_{\text{NCPF,}j}} = y_{\text{rel}} \cdot \frac{M_{\text{NCPF}} \cdot (m_{\text{NCP,}j} + r \cdot m_{\text{NCPF,}j})}{M_{\text{NCP}} \cdot m_{\text{NCPF,}j} (1-r)} \quad 1.2$$

Where:

- A_{xxx} is the corresponding area measured with GC
- m_{xxx}^{total} is the total weight of the corresponding species in the solution
- m_{xxx} is the added weight of the corresponding species
- M_{xxx} is the molecular weight of the corresponding species
- y_{xxx} is the sensitivity of the GC system to the corresponding species
- y_{rel} is the relative sensitivity of the GC system to NCP versus NCPF
- r is the weight percentage of NCP in the NCPF standard

If $L = \frac{A_{\text{NCP, i}}}{A_{\text{NCPF, i}}} \cdot \frac{A_{\text{NCPF, j}}}{A_{\text{NCP, j}}}$, after division of equation 1.1 by equation 1.2:

$$\mathbf{r} = \frac{\mathbf{m}_{\text{NCP, i}} \cdot \mathbf{m}_{\text{NCPF, j}} - \mathbf{L} \cdot \mathbf{m}_{\text{NCPF, i}} \cdot \mathbf{m}_{\text{NCP, j}}}{\mathbf{L} \cdot \mathbf{m}_{\text{NCPF, i}} \cdot \mathbf{m}_{\text{NCPF, i}} - \mathbf{m}_{\text{NCPF, i}} \cdot \mathbf{m}_{\text{NCPF, j}}}$$
1.3

Considering all the $\binom{5}{2} = 10$ combinations, we received that there is $100 \cdot r = 8.3 \pm 1 \% w/w$

<u>NCP content in the NCPF standard</u> (68% confidence level). Since there are no other significant impurities, purity of the NCPF used for internal standard calibration is $100 \cdot (1-r) = 91.7 \pm 1$ %.

S.5/b Determination of the response factor y* of NCPF and NCP versus dibenzyl for the internal standard calibration

For a choosen solution "i":

$$\frac{\mathbf{A}_{\mathrm{S,i}}}{\mathbf{A}_{\mathrm{DB,i}}} = \mathbf{y}_{\mathrm{S}} \cdot \frac{\mathbf{M}_{\mathrm{DB}} \cdot \mathbf{m}_{\mathrm{S,i}}}{\mathbf{M}_{\mathrm{S}} \cdot \mathbf{m}_{\mathrm{DB,i}}}$$
 1.4

Where:

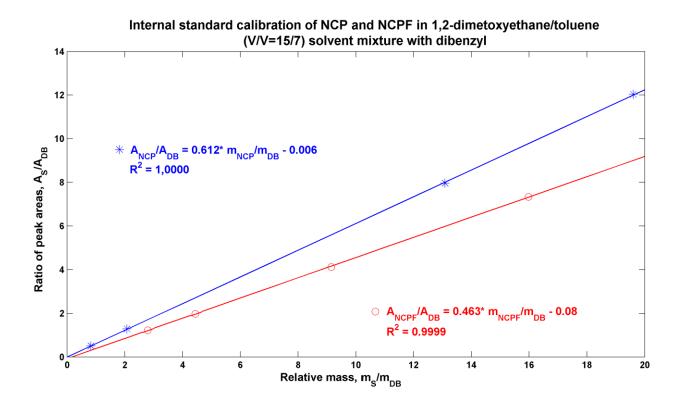
- y_S is the response factor of species S (NCPF or NCP) versus dibenzyl
- A_{DB} is the peak area corresponding to dibenzyl
- M_{DB} is the molecular weight of dibenzyl
- M_{DB} is the molecular weight of species S (NCPF or NCP)
- m_S is the total weight of species S in the solution (in case of NCPF $m_{NCPF, \text{ total}} = (1-r) \cdot m_{NCPF}$, in case of NCP, $m_{NCP, \text{ total}} = m_{NCP} + r \cdot m_{NCPF}$)

For convenience, we incorporate the molecular weights into the response factor, thus we can calculate with weights:

$$\mathbf{y}_{\mathrm{S}}^{*} = \mathbf{y}_{\mathrm{S}} \cdot \frac{\mathbf{M}_{\mathrm{DB}}}{\mathbf{M}_{\mathrm{S}}}$$
 1.5

As shown below, y_s^* values can be determined as slopes of the $A_s/A_{DB} = y_s^* mS/m_{DB} + b$ functions. The corresponding values to NCP and NCPF:

 $y_{NCPF}^* = 0.463$ and $y_{NCP}^* = 0.612$



S.5/c Calculation of the yield and the remaining NCP content

The mass of species S (NCP or NCPF) in the reaction mixture, after hydrolysis and neutralization:

$$\mathbf{m}_{\mathrm{S}} = \frac{\mathbf{A}_{\mathrm{S}} \cdot \mathbf{m}_{\mathrm{DB}}}{\mathbf{y}_{\mathrm{S}}^* \mathbf{A}_{\mathrm{DB}}}$$
 1.6

Example GC result:

Measurement ID: GT_110310, measurement date: 11.03.2010

Conditions: reagent amounts BuLi/BuLi/TFAET/HCl = 2.15/0.15/1.3/2.25 NCP equivalents, [NCP] = 0.22M, solvent: DME/Toluene (V/V = 15/7), waiting times after dosing steps: 60/10/30/10 min, $f_{BuLi}/f_{TFAEt}/f_{HCl} = 1.0/0.2/0.2$ mL/min, $T_{react} = 5$ °C, stirrer speed: 600 rpm.

Species	Retention time [min]	Area [pA*s]	Weight [mg]	Y (mol/mol initial NCP)
Dibenzyl	8.852	1527	100.6	
NCPF	9.294	10422	1483*	96.45
NCP	9.71	308	33.2*	3.13

* calculated according to equation 1.6