Electronic Supporting Information

Mechanistic investigations of bifunctional squaramide organocatalyst in asymmetric Michael reaction, observation of stereoselective retro-Michael reaction

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I. Characterization of catalyst 3

A. Characterization of catalyst 3 by NMR spectroscopy

The preparation of 3-(benzylamino)-4-(((1S)-(6-methoxyquinolin-4-yl)((2S,4S,5R)-5-vinylquinuclidin-2-yl)methyl)amino)cyclobut-3-ene-1,2-dione (catalyst **3**)

To the solution of diethyl squarate (10.0 mmol, 1.702 g) in diethyl ether (10 ml) benzylamine (11.0 mmol, 1.179 g) was added drop wise over a period of 5 minutes at ambient temperature and stirred overnight. A small amount of white powder precipitated which was removed by filtration. 9-amino(9-deoxy)epihydroquinine (11 mmol) was added to the filtrate and stirred overnight at room temperature. The catalyst precipitated and was separated by filtration and washed with diethyl ether (2x 10 ml) to yield 4.424 g (87 % yield).

Nuclear Magnetic Resonance spectra (NMR) were acquired on a Bruker DRX-500 (500 MHz) instrument using TMS as internal standard. All assignments of the catalyst **3** are confirmed by 2D-COSY, 2D-HSQC, 2D-HMBC and NOESY measurements.

The structure of the catalyst **3** was determined in CDCl₃ solution by ¹H- and ¹³C NMR measurements at 310 and 320 K, respectively. Assignments were confirmed by 2D-COSY, 2D-HSQC, 2D-HMBC and NOESY methods.



¹H-NMR (500 MHz, CDCl₃, T = 300?): δ ppm 8.61 (d, *J*=4.4 Hz, 1H, H-2'), 8.01 (d, *J*=9.2 Hz, 1H, H-8'), 7.77 (br s, 1H, H-5'), 7.48 (br s, 1H, H-3'), 7.40 (dd, *J*=2.5, 9.2 Hz, 1H, H-7'), 7.11 (br m 3H, Ph-3"4"5"), 6.92 (br d, *J*=4.7 Hz, 2H, Ph-2"6"), 6.07 (br s, 1H, H-9), 5.78 (td, 1H, H-10), 5.01 (d, *J*=17.2 Hz, 1H, H^α-11), 4.96 (d, *J*=10.4 Hz, 1H, H^β-11), 4.35 (dd, *J*=14.3, 61.2 Hz, 2H, H^{αβ}-12), 3.92 (s, 3H, H-13, -OCH₃), 3.52 (br s, 1H, H-8), 3.40 (s, 1H, H^α-6), 3.20 (dd, *J*=10.4, 13.5 Hz, 1H, H^{exo}-2), 2.79 (br d, *J*=13.4 Hz, 1H, H^{endo}-2), 2.72 (m, 1H, H^β-6), 2.29 (br s, 1H, H-3), 1.66 (br s, 1H,

H-4), 1.59 (br s, 2H, H-5^{$\alpha\beta$}), 1.45 (t, *J*=11.0 Hz, 1H, H^{β}-7), 0.75 (s, 1H, H^{α}-7). ¹³C-NMR (500 MHz, CDCl₃, T = 310? K) δ ppm 167.5 (C-14), 159.2 (C-6'), 148.3 (C-4'), 147.5 (C-2'), 144.6 (C-8'a), 141.5 (C-10), 138 (C-1''), 130.6 (C-8'), 129.4 (C-4a'), 128 (C-2''), 122.3 (C-7'), 120.2 (C-3'), 115.2 (C-11), 102.1 (C-5'), 60.2 (C-8), 56.4 (C-13), 56.3 (C-2), 53.3 (C-9), 48.1 (C-12), 41 (C-6), 39.6 (C-3), 27.9 (C-4), 27.9 (C-5), 25.9 (C-7)

The effect of temperature to the conformation was investigated. Based on the comparison of the ¹H-NMR spectra of catalyst **3** at two different temperatures, it can be seen that the molecule can freely rotate around the C8-C9 bond and the quinuclidine and quinoline ring is sensitive to this conformation changes.



Figure 1. ¹H spectra of catalyst **3** at different temperatures.

The chemical shifts of ¹³C NMR signals were obtained and assigned in the 2D spectra on the basis of the cross-peaks discernible.



Figure 2. COSY spectra of catalyst **3** in CDCl₃.



Figure 3. HSQC spectra of catalyst **3** in CDCl₃.



Figure 4. HMBC spectra of catalyst **3** in CDCl₃.

B. Characterization of catalyst 3 by X-Ray study

Additionally, the solid state structure of the catalyst **3** was determined by X-ray diffraction. The X-ray crystal structure is in consistence with the structures in solution determined by NMR spectroscopy. On the other hand, it is not sure that the conformer in solid phase determined by molecular packing fully corresponds to that adopted in solution relevant to catalytic activity.

X-ray crystallographic data were collected on a Rigaku R-AXIS-RAPID diffractometer¹ (graphite monochromator). Crystallographic data (excluding structure factors) for the structure in this paper have been deposited with the Cambridge Crystallographic Data Centre as supplementary publication CCDC no. **1412465**. Copies of the data can be obtained, free of charge, on application to CCDC, 12 Union Road, Cambridge CB2 1EZ, UK, (fax: t44-(0)1223 336033 or e-mail: deposit@ccdc.cam.ac.uk).

Crystal data: $C_{31}H_{32}N_4O_3$, *Fwt.*: 508.60, colourless, platelet, size: 0.25 x 0.25 x 0.05 mm, orthorhombic, space group *P*2₁2₁2₁, *a* = 8.7790(2)Å, *b* = 11.6400(2)Å, *c* = 26.194(1)Å, α = 90°, β = 90°, γ = 90°, *V* = 2676.7(1)Å³, *T* = 295(2)K, *Z*= 4, *F*(000) = 1080, *D*_x = 1.262 Mg/m³, μ 0.659mm⁻¹.

A crystal of SHELX was mounted on a loop. Cell parameters were determined by least-squares using 21339 ($6.50 \le \theta \le 71.50^{\circ}$) reflections.

Intensity data were collected on a RAXIS-RAPID diffractometer (monochromator; Cu- $K\alpha$ radiation, $\lambda = 1.54178$ Å) at 295(2) K in the range $6.538 \le \theta \le 56.906^2$. A total of 23228 reflections were collected of which 3579 were unique [R(int) = 0.0293, $R(\sigma) = 0.0304$]; intensities of 2886 reflections were greater than $2\sigma(I)$. Completeness to $\theta = 0.995$.

A numerical absorption correction³ was applied to the data (the minimum and maximum transmission factors were 0.826 and 0.941).

The structure was solved by direct methods⁴ (and subsequent difference syntheses).

Anisotropic full-matrix least-squares refinement⁴ on F^2 for all non-hydrogen atoms yielded $R_1 = 0.0333$ and $wR^2 = 0.0657$ for 1332 [$I > 2\sigma(I)$] and $R_1 = 0.0547$ and $wR^2 = 0.0900$ for all (3579) intensity data, (number of parameters = 345, goodness-of-fit = 1.192, the maximum and mean shift/esd is 0.000 and 0.000). The absolute structure parameter is 0.01(5). (Friedel coverage: 0.723, Friedel fraction max.: 0.990, Friedel fraction full: 0.990).

The maximum and minimum residual electron density in the final difference map was 0.18 and -0.19 e.Å⁻³.

The weighting scheme applied was $w = 1/[\sigma^2(F_o^2) + (0.02400.9144P)^2 + 0.9144P]$ where $P = (F_o^2 + 2F_c^2)/3$.

Hydrogen atomic positions were calculated from assumed geometries except H1A, H1B that were located in difference maps. Hydrogen atoms were included in structure factor calculations but they were not refined. The isotropic displacement parameters of the hydrogen atoms were approximated from the U(eq) value of the atom they were bonded to.

²CrystalClear SM 1.4.0 (Rigaku/MSC Inc., 2008).

¹ A diffractometer purchase grant from the National Office for Research and Technology (MU-00338/2003) is gratefully acknowledgement.

³NUMABS: Higashi, T. (1998), rev. 2002. (Rigaku/MSC Inc.)

⁴G.M. Sheldrick, Acta Cryst. (2008). A64, 112-122.

ORTEP style molecular structure diagram can be found in Figure 5, while crystallographic data are in Table 1. 5



Figure 5. The crystallographically independent molecule in the asymmetric unit of the crystal **3** with the atomic labelling. Displacement ellipsoids are drawn at the 30% probability level.

⁵ PLATON: Spek, A.L. J.Appl.Cryst. (2003), 36, 7-13.

Table 1. Crystal data and structure refinement

Empirical formula	$C_{31}H_{32}N_4O_3$
Formula weight	508.60
lemperature	295(2)
	Cu-Ka,
wavelength (A)	λ =1.54178
Crystal system	orthorhombic
Space group	<i>P</i> 21 21 21
	0.7700(0)
	8.7790(2)
	11.6400(2)
	20.194(1)
α (°)	90
β(°)	90
$\gamma \sqcup^{\sim} \sqcup$	90
Volume (A ³)	2676.7(1)
	4
Density (calculated) (Mg/m ³)	1.262
Absorption coefficient, μ (mm ⁻¹)	0.059
F(000)	
Crystal colour, description	
Absorption correction	
Max and min transmission	$\begin{array}{c} \text{numerical} \\ 0.826 \text{ and } 0.041 \end{array}$
$\Omega \Box$ range for data collection (°)	6.528 < 0 < 56.006
	$0.550 \le 0 \le 50.900$
Index ranges	$-9 \ge \Box \ge 9$,
	$-12 \leq 11 \leq 12$,
Deflections collected	-27 ≤ UU≤ 28
Completeness to 20	23220
Absolute etructure perometer	0.995
Elack (x)	0.01(5)
Hooft (V)	0.01(3)
Darsons $(z)^6$	0.01(8)
Friedel coverage	0.723
Friedel fraction max	0.990
Friedel fraction full	0.990
Independent reflections	3579 [<i>R</i> (int) =0.0293]
Reflections $l \ge 2\sigma(l)$	2886
Refinement method	full-matrix least-squares on F^2
Data / restraints / parameters	3579 /0 /345
Goodness-of-fit on F2	1.192
Final R indices $[P 2\sigma(I)]$	$R_1 = 0.0333$,
	$wR^2 = 0.0657$
R indices (all data)	<i>R</i> ₁ =0.0547,
	wR ² =0.0900
Max. and mean shift/esd	0.000;0.000
Largest diff. peak and hole	0.18 and -0.19 e.Å ⁻³

⁶ S. Parsons, H. Flack, Acta Cryst. (2004), A60, s61.

Comparison of crystal structure of molecule **3** and other quinine-squaramide structures from the literature

The molecule **3** stands by the hydrogen bond acceptor isoquinoline and quinuclidine groups, and by the hydrogen bond donor squaramide groups. The aromatic attribution of the phenyl groups and isoquinoline groups are important organiser forces in the crystal lattices as well.

Two similar conformation squaramide crystal structures were found in the literature with similar isoquinoline, quinuclidine and phenyl groups⁷,⁸. One of them (**UBADUY**)⁷ has the same *R*, *S*, *S*, *S* configuration, the case of **NOLRIQ**⁸ the atoms have *R*, *S*, *R*, *R* configuration.

In the crystal lattices of **NOLRIQ** and **3** there are hydrogen bonds interactions between the squaramides moieties of the neighbour molecules. These interactions are the strongest secondary interaction in the lattices because of the high acidity of squaramide. In consequence these contacts are the prime organiser forces in the crystal lattice⁹. In the **NOLRIQ** there are additional π ···F and H···F secondary interactions (Figure 6.).



Figure 6. The layers of molecules (NOLRIQ) connected by strong squaramides...squaramides hydrogen bonds secondary interaction and an other aromatic layer can be observed in the crystal lattice (showed by using program Mercury10).

In the **UBADUY** structure a water molecule is connected with N-H···O type hydrogen bonds to the squaramides.

Molecule **3**, **NOLRIQ** and **UBADUY** molecules have similar conformation (Figure 7.), only the placement of the phenyl rings and isoquinoline rings differ. The phenyl ring turned at the CH_2 atoms by $87(2)^\circ$, the isoquinoline group turned by $43(3)^\circ$.

⁷ L. Dai, H. Yang, F. Chen, *Eur. J. Org. Chem.*, **2011**, 5071–5076

⁸ J. P. Malerich, K. Hagihara, V. H. Rawal, J. Am. Chem. Soc., 2008, 130, 14416–14417

⁹ R. I. Storer, C. Aciro, L. H. Jones, Chem. Soc. Rev., 2011,40, 2330-346

¹⁰ C. F. Macrae, I. J. Bruno, J. A. Chisholm, P. R. Edgington, P. McCabe, E. Pidcock, L. Rodriguez-Monge, R. Taylor, J. van de Streek, P. A. Wood, *J. Appl. Cryst.*, **2008**, 41, 466-470



Figure 7. Molecular overlay of structure **3** and UBADUY to observe conformation differences¹⁰.

UBADUY structure lattice enclathrates a toluene molecule (Figure 8.). Its placement is enclosed by $\pi \cdots \pi$ and CH $\cdots \pi$ interactions.



Figure 8. The aromatic interactions of the enclosed toluene in the UBADUY crystal lattice.

In absence of the enclathrated solvent the toluene, the phenyl groups turn to the neighbouring isoquinoline groups as it happens in the case of structure of **3**. (Figure 9.).



Figure 9. In the measured crystal **3** the molecules form a layer of alternating molecules connected by the squaramides...squaramides hydrogen bonds, strengthened by $\pi \cdots \pi$ interaction as well.

II. Detailed results of kinetic studies

A. Calibration of IR measurements

The starting materials and the product were dissolved in $CHCl_3$ and measured separately by FTIR and a calibration set was carried out. In all case the most intensive and nonoverlapping bands were selected and the calibration factor were determined based on Lambert-Beer's law (A= ϵ^*c^*l). The following bands were selected: Acetylacetone (**2**): 1621 cm⁻¹ (v C=O); β -Nitrostyrene (**1**): 1345 cm⁻¹ (v NO₂); 3-(2-nitro-1-phenylethyl)pentane-2,4-dione (**4**): 1556 cm-1 (v NO₂)



Figure 10. The recorded IR spectra of the 1 M solution of the reactants and the solvent.



Figure 11. Calibration curves of FTIR absorbance: a) Absorbance of β -nitrostyrene (1) (c = 0.05-0.75 M), b) Absorbance of acetylacetone (2) (c = 0.05-1.0 M) c) Absorbance of the Michael adduct (4) (c = 0.05-0.75 M) M)

β-Nitrostyrene (1): (ε^{*}I) = **0.968**, at 1345 cm⁻¹ (ν NO₂) Acetylacetone (2): (ε^{*}I) = **2.896**, at 1621 cm⁻¹ (ν C=O) Product (4): 3-(2-nitro-1-phenylethyl)-pentane-2,4-dione: (ε^{*}I) = **1.286**, at 1556 cm⁻¹ (ν NO₂)

B. General procedure for in situ IR experiments

Infrared spectra were recorded with a ReactIR 1000 Reaction Analysis System attached to atmospheric pressure silicon (SiCompTM) probe head. *In situ* IR experiments were performed in a three necked 25 ml glass flask. In every 2 minutes 64 scan were recorded between 4000-600 cm⁻¹, with resolution of 4 cm⁻¹ and the mean was converted to a single spectrum. The reaction was stopped after 1-3 h depending on the conversion. In some cases the spectra were recorded in every 1 minute registering 32 scans. From the selected bands were generated a quick profile of the reaction (Figure 12.) and the concentrations were calculated based on calibration.



Figure 12. Quick profile of a representative reaction using ReactIR 1000.

C. Method of reaction rate calculation

The reaction rate was generated from the concentration versus time plot by mathematically differentiating the fifth–order best fit polynomial f(t). (Since this is empirical method, df(*t*)/dt is valid only over the range of reactants for which experimental data was collected) There are three representative plot on Figure 13.

To test the reproducibility of the measurement, three parallel measurements were carried out at same conditions. The deviation of calculated initial rate was 18 %, and this value was applied for further calculations.



Figure 13. a) Reactant concentrations during the reaction, $[1]_0 = 0.6$ M, $[2]_0 = 0.3$ M, [3] = 0.006 M; b) Reaction rate vs. time plot by the same reaction, c) Conversion vs. time plot by the same reaction

D. Calculation of observed rate constant

The rate equation of this reaction could be written in the form: $v = k \cdot [1]^n \cdot [2]^m \cdot [3]^{\circ} \cdot [4]^p$

Determination the order of the catalyst (3):

By measuring the reaction rate at different catalyst loadings (from 0.003 to 0.021 M), it was determined that the reaction is first-order in catalyst **3**. The data according to Fig. 3 in the article is presented on Table 2.

Nr.	Catalyst loading / mol%	[3] / M	v₀ / M*h⁻¹	In [1]₀	In v₀				
A: [1] ₀ = 0.3 M, [2] ₀ = 0.6 M									
1	1	0.0030	0.351	-5.809	-1.047				
2	1.5	0.0045	0.569	-5.404	-0.564				
3	2	0.0060	1.205	-5.116	0.186				
4	4	0.0120	1.944	-4.423	0.665				
5	5	0.0150	2.962	-4.200	1.086				
6	6	0.0180	3.466	-4.017	1.243				
7	7	0.0210	4.280	-3.863	1.454				
B: [1] ₀ = 0.6	M, [2] ₀ = 0.3 M								
8	1	0.0030	0.308	-5.809	-1.179				
9	1.5	0.0045	1.004	-5.404	0.004				
10	2	0.0060	1.474	-5.116	0.388				
11	4	0.0120	2.847	-4.423	1.046				
12	5	0.0150	3.055	-4.200	1.117				
13	6	0.0180	3.782	-4.017	1.330				
14	7	0.0210	5.007	-3.863	1.611				

Table 2. Reaction conditions and reaction rate using different catalyst (3) concentration

The order of the reactants was determined by changing the initial concentration of the given substrate, while the other parameters were kept constant. Accordingly, the rate equation could be further simplified. The apparent rate constant was calculated based on logarithmic transformation.

The order of the nucleophile (**2**) was determined in the non-saturated regime I. and the reaction rate dependence was found less than first-order in nucleophile concentration. The data according to Fig. 5 in the article is presented on Table 3.

The order of the nucleophile (2) was determined based on following equations: $v = k_{obs} * [2]^m$

 $\ln v = m^* \ln[2] + \ln k_{obs}$

Table 3.: Reaction conditions and reaction rate using different acetylacetone concentration

	Nr.	[2] ₀ / M	v ₀ / M*h ⁻¹	In [2]₀	In v ₀			
[1] ₀ = 0.3 M, catalyst: [3] = 0.006 M								
Non-saturated regi	mel 1	0.15	0.267	-1.897	-1.321			
	2	0.30	0.500	-1.204	-0.693			
	3	0.38	0.664	-0.981	-0.409			
	4	0.60	0.860	-0.511	-0.151			
	5	0.75	1.171	-0.288	0.158			
Regime II.	6	0.90	1.189					
	7	1.50	1.106					
	[1] ₀ = 0.6 M, ca	atalyst: [3]	= 0.006 M					
Non-saturated regi	mel 8	0.15	1.010	-1.897	0.010			
	9	0.20	1.110	-1.609	0.104			
	10	0.30	1.295	-1.204	0.258			
	11	0.40	1.785	-0.916	0.579			
	12	0.50	1.849	-0.693	0.615			
	13	0.60	2.092	-0.511	0.738			
	14	0.75	2.300	-0.288	0.833			
Regime II.	15	0.90	2.190					
	16	1.20	2.232					
	17	1.50	2.054					
[1] ₀	Fitted line	1	m	k _{obs} (h⁻¹)				
low (0.3 M)	y = 0.8894x + 0.38	351 0).9 k _{obs} ,	$_{1} = 1.47 \pm 0$.27 h ⁻¹			
nign (0.6 M)	y = 0.5424x + 0.99	199 0	.5 K _{obs} ,	$_2 = 2.72 \pm 0$.49 N°'			

In the same way, the order of the electrophile (1) was determined in the non-saturated regime I, and the reaction rate dependence was found first-order in electrophile concentration. The data according to Fig. 6 in the article is presented on Table 4. The order of the electrophile (1) was determined based on following equations:

 $v = k_{obs}^{*}[1]^{n}$ In $v = n^{*}ln[1] + ln k_{obs}$

			Nr.	[1] ₀ / M	v ₀ / M*h ⁻¹	ln [1]₀	In v₀
[2] ₀ = 0.3 N	l, catalyst: [3] = 0.006 M						
Non	-saturated regime I.		1	0.30	0.480	-1.204	-0.734
			2	0.40	0.610	-0.916	-0.494
			3	0.50	0.860	-0.693	-0.151
			4	0.60	1.070	-0.511	0.068
			5	0.75	1.200	-0.288	0.182
	Regime II.		6	0.90	1.160		
			7	1.00	1.101		
[2] ₀ = 0.6 N	l, catalyst: [3] = 0.006 M						
Non	-saturated regime I.		8	0.3	0.856	-1.204	-0.156
			9	0.4	1.339	-0.916	0.292
			10	0.5	1.668	-0.635	0.512
			11	0.7	2.092	-0.357	0.738
	Regime II.		12	0.8	2.156		
			13	0.9	1.987		
			14	1.0	1.924		
			15	1.1	1.799		
[2] ₀	Fitted line	n		k _{obs} (h⁻¹)			
low (0.3 M) high (0.6 M)	y = 1.0682x + 0.5458 y = 1.0292x + 1.1472	1 1	k _{obs,} k _{obs,}	$_{3} = \overline{1.73 \pm 0.37}$ $_{4} = 3.15 \pm 0.57$	1 h ⁻¹ 7 h ⁻¹		

Table 4.: Reaction conditions and reaction rate using different β -nitrostyrene concentration

In the same way, the order of the product (4) was determined.

In order to investigate the possible inhibition effect of the product, the reaction was monitored in the presence of different amount of the product (**R-4**) and different amount of enantiomer (**S-4**). The data according to Fig. 7 in the article is presented on Table 5. and the data according to enantiomer (**S-4**) is presented on Table 6.

The order of the product (4) was determined based on following equations:

$$v = k_{obs}^* [4]^p$$

 $\ln v = p^{*}\ln[4] + \ln k_{obs}$

Table 5. Reaction conditions and reaction rate adding different amount R-product at the beginning of the reaction

Nr.	[R-4] ₀ / M	v ₀ / M*h ⁻¹	In [R-4]₀	$\ln v_0$
1	0.1	0.899	-2.303	-0.107
2	0.2	0.757	-1.609	-0.278
3	0.3	0.480	-1.204	-0.735

Table 6. Reaction conditions and reaction rate adding	different amount S-product at the beginning of the
reaction	

Nr.	[S-4] ₀ / M	v ₀ / M*h ⁻¹	In [S-4]₀	ln v _o
4	0.1	0.798	-2.303	-0.226
5	0.2	0.655	-1.609	-0.423
6	0.3	0.480	-1.204	-0.735
[4] ₀	F	itted line	р	k _{obs} (h⁻¹)
product (R) enantiomer (S)	y = -0. y = -0.	537x - 1.289 454x - 1.218	-0.5 -0.5	$k_{obs,5} = 0.28 \pm 0.05 h^{-1}$ $k_{obs,6} = 0.30 \pm 0.05 h^{-1}$

E. Calculation of activation energy

Activation energy was acquired by Arrhenius plot analysis. The reactions rates were determined at T = 274.15, 298.15 and 318.15 K. Reaction conditions: $[1]_0 = 0.3$ M, $[2]_0 = 0.6$ M; [3] = 0.006 M; solvent: CHCl₃. The data according to Fig. 9 in the article is presented on Table 7.

	Table 7.	Reaction	rates a	at different	temperature
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Τ/Κ	(10³/T) / (1/K)	v/ M*h ⁻¹	In (v/c _{cat}) / (s ⁻¹)
274.15	3.6476	0.486	-3.8
298.15	3.3540	1.122	-3.0
318.15	3.1432	1.700	-2.5

F. Calculation of association constant by NMR measurement using Job plot

The binding constants were evaluated by fitting the data to a 1:1 binding isotherm, and 1:1 binding was verified by Job plot. The binding constants were calculated from the changes in chemical shifts of catalysts protons. Nonlinear curve fitting was carried out with the Solver program in Excel. The binding constants K and the asymptotic change in chemical shift $\Delta \delta_{max}$ were chosen as the free parameters for fitting.

Complex formation of catalyst (4) and acetylacetone (2)

The complex of the nucleophile (acetylacetone, **2**) and the catalyst (**4**) was investigated. The catalyst concentration was varied between 0.02 and 0.2 M, while the concentration of the nucleophile was adjusted to get 0.2 M sum of concentration of each samples. The signals of catalyst were significantly shifted when adding the substrate. Signals of the

quinuclidine and some of the quinoline ring were changed. The most relevant peak belongs to the methoxy methylene of the catalyst (3H, δ_0 = 3.92 ppm, see on Figure 14.).



Figure 14. Chemical shifts of catalyst during Job-plot compared to free catalyst and the selected peaks for calculation of binding constant

Job plot calculation for catalyst (3) - acetylacetone (2)complex formation

Equations for determining binding constants:

 $[Q]_0$ = initial concentration of catalyst **3**

 $[A]_0$ = initial concentration of acetylacetone **2**

[QA] = actual concentration of catalyst-acetylacetone complex

$$\begin{split} \delta_{obs} &= X_Q \delta_0 + X_{QA} \delta_{max} \\ \delta_{obs} &= \frac{[Q]_0 - [QA]}{[Q]_0} \delta_0 + \frac{[QA]}{[Q]_0} \delta_{max} = \delta_0 + \frac{[QA]}{[Q]_0} (\delta_{max} - \delta_0) = \delta_0 + \frac{[QA]}{[Q]_0} \Delta \delta_{max} \\ K_A &= \frac{[QA]}{([Q]_0 - [QA])([A]_0 - [QA])} \\ ([Q]_0 - [QA])([A]_0 - [QA]) &= \frac{[QA]}{K_A} \\ [QA]^2 - ([Q]_0 + [A]_0 + \frac{1}{K_a})[QA] + [Q[_0[A]_0 = 0 \\ \\ &= \frac{[Q]_0 + [A]_0 + \frac{1}{K_a} \pm \sqrt{([Q]_0 + [A]_0 + \frac{1}{K_a})^2 - 4[Q]_0[A]_0}}{2} \end{split}$$

[Q] ₀ / M	X _Q	δ _{obs} (H-13)	[QA] _{calc}	δ_{calc}	$(\delta_{obs}-\delta_{calc})^2 \ (res^2)$	Sum res ²
0.020	1.0	3.924	0.0000	3.924		

	3.48E-06	3.930	0.0012	3.932	0.9	0.018
	5.35E-07	3.937	0.0023	3.936	0.8	0.016
	7.12E-06	3.944	0.0031	3.941	0.7	0.014
	7.46E-06	3.951	0.0036	3.948	0.6	0.012
	2.68E-06	3.958	0.0038	3.956	0.5	0.010
	3.57E-07	3.964	0.0036	3.964	0.4	0.008
	4.44E-06	3.970	0.0031	3.972	0.3	0.006
3.039E-05	4.31E-06	3.975	0.0023	3.977	0.2	0.004

 $c_{total} = [Q]_0 + [A]_0 = 0.02 M$

The calculated parameters: K = 98.89±1.02 M⁻¹ and $\Delta \delta_{max}$ = 0.086±0.002 ppm Based on this parameters, the Job plot was determined:

X _Q (Cat)	X _A (Nu)	Δδ _{obs} (<i>H</i>-13)	$\Delta \delta^* \mathbf{X}_{\mathbf{Q}}$
1,0	0.0	0.000	0.0000
0.9	0.1	0.006	0.0055
0.8	0.2	0.013	0.0102
0.7	0.3	0.020	0.0138
0.6	0.4	0.027	0.0160
0.5	0.5	0.034	0.0168
0.4	0.6	0.040	0.0160
0.3	0.7	0.046	0.0138
0.2	0.8	0.051	0.0102
0.1	0.9	0.069	0.0069
0.0	1.0	0.000	0.0000

Table 9.	Calculation	for	Job	plot
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