

Supplementary Information

Development of Improved Spectrophotometric Assays for Biocatalytic Silyl Ether Hydrolysis

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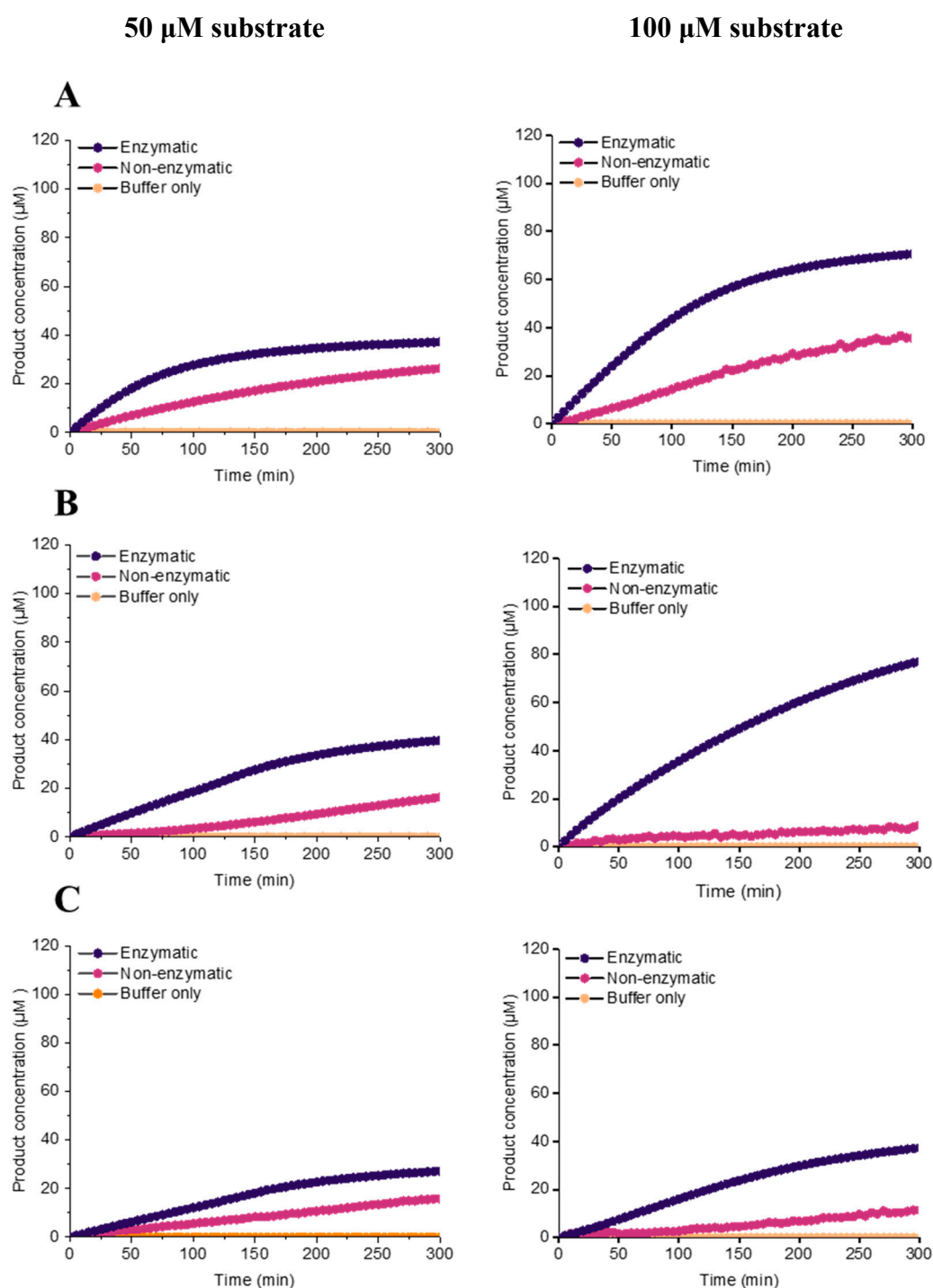


Figure S1. Graph of concentration of nitrophenolate produced against time for enzymatic and non-enzymatic hydrolysis measured at the λ_{max} of their corresponding 4-nitrophenolate ions. (A) substrate **1** at 405 nm, (B) substrate **4** at 414 nm, (C) substrate **5** at 394 nm, (D, next page) substrate **6** at 398 nm, (E) substrate **7** at 294 nm. Data were quantified using their corresponding calibration curves (Supporting Information Figures S4). Reactions were carried out with 6.7 μM TF-Sil α -Strep, 50 μM substrate (Left), 100 μM substrate (Right), 10 % v/v 1,4-dioxane, 50 mM Tris buffer at the pH 8.5 and 100 mM NaCl.

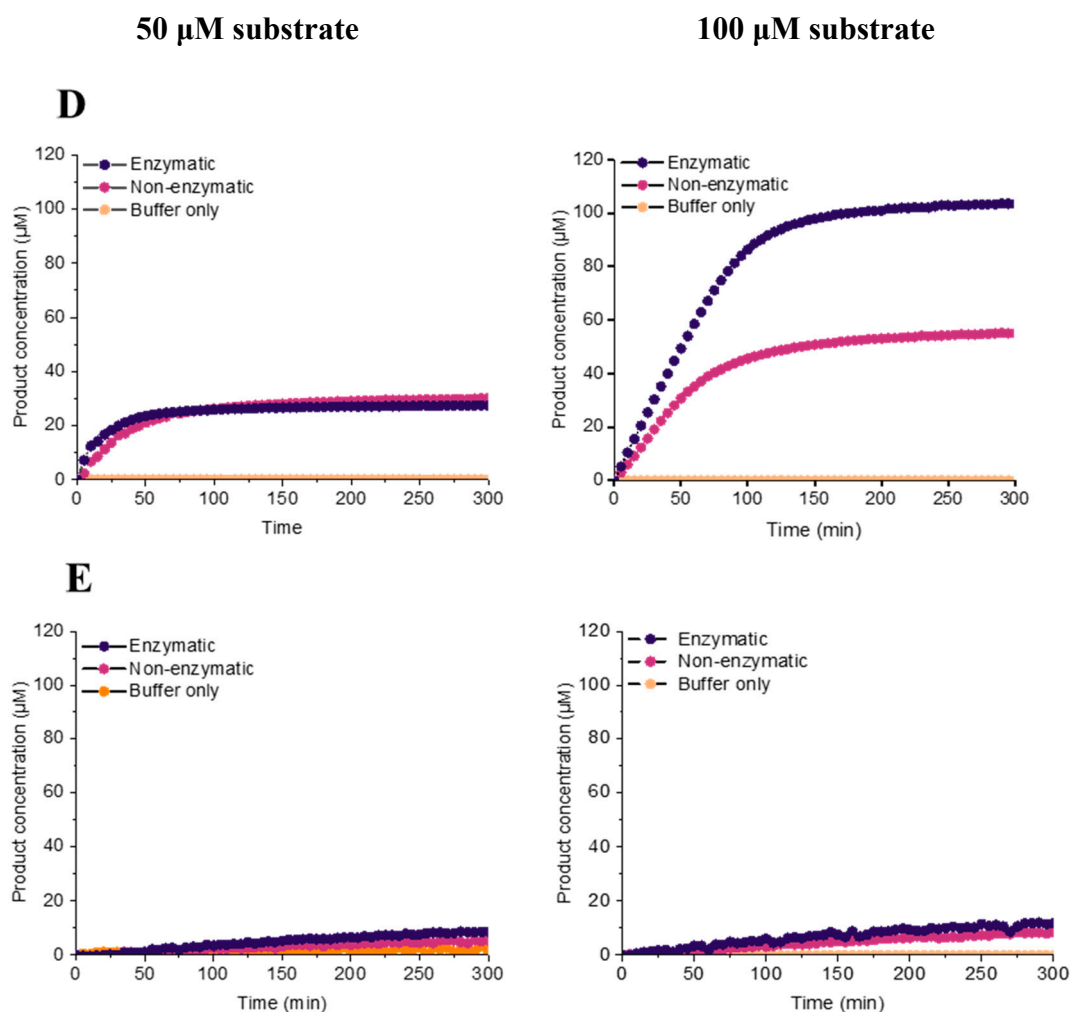


Figure S1 (continued). Graph of concentration of silanols produced showing enzymatic and non-enzymatic (background) hydrolysis measured at the λ_{max} of their corresponding 4-nitrophenolate ions after 300 min. Buffer only is used as the blank. (A) substrate **1** at 405 nm, (B) substrate **4** at 414 nm, (C) substrate **5** at 394 nm, (D, next page) substrate **6** at 398 nm, (E) substrate **7** at 294 nm. Data were quantified using their corresponding calibration curves (Supporting Information Figures S4). Reactions were carried out with 6.7 μ M TF-Sil α -Strep, 50 μ M substrate (Left), 100 μ M substrate (Right), 10 % v/v 1,4-dioxane, 50 mM Tris buffer at the pH 8.5 and 100 mM NaCl.

Table S1. Rates of enzymatic vs. background hydrolysis showing the initial rates and fold increase of enzyme-catalysed hydrolysis of different silyl ether substrates. Enzymatic reactions were carried out with 6.7 μM TF-Sil α -Strep, 50 or 100 μM substrate, 10 % v/v 1,4-dioxane, 50 mM Tris buffer at pH 8.5 and 100 mM NaCl.

Substrate	Initial rate at 50 μM substrate ($\mu\text{M min}^{-1}$)			Fold difference [†]	Initial rate at 100 μM substrate ($\mu\text{M min}^{-1}$)			Fold difference [†]
	Enzymatic	Background	Net*		Enzymatic	Background	Net*	
1	0.44	0.15	0.29	2.9	0.80	0.37	0.43	2.2
4	0.19	0.03	0.16	6.2	0.35	0.03	0.32	11.6
5	0.12	0.06	0.06	2.0	0.14	0.05	0.09	2.8
6	0.72	0.54	0.18	1.3	1.36	0.79	0.57	1.7
7	0.04	0.027	0.013	1.5	0.04	0.03	0.01	1.3

*Net initial rate is measured as the difference in initial rate between the enzymatic and background hydrolysis.

[†]Fold difference (rate ratio) calculated as the ratio of the enzymatic reaction relative to background (non-enzymatic) reaction.

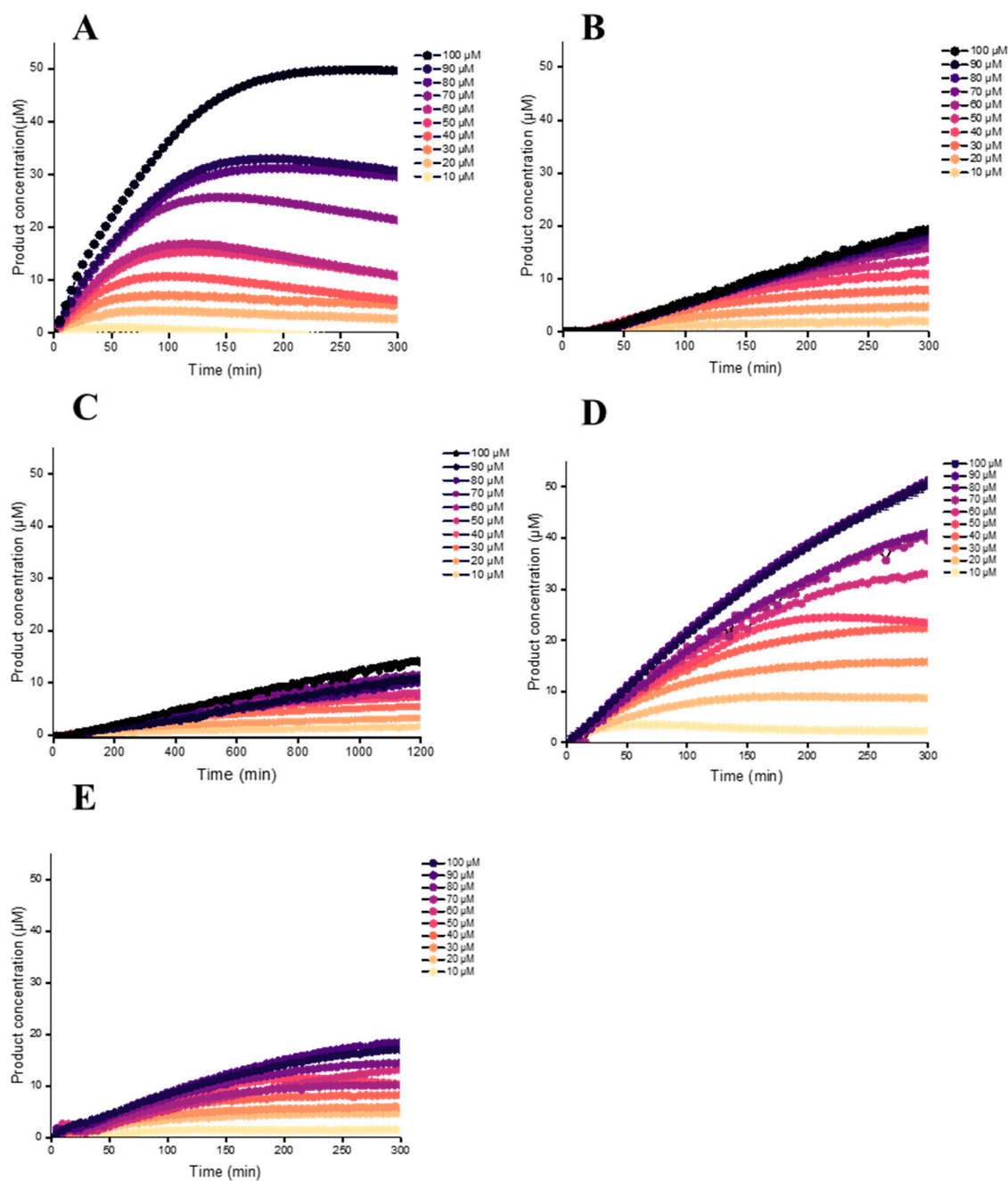


Figure S2. Graph of net concentration of corresponding phenoxide products measured by UV-Vis absorbance at their respective λ_{max} . Substrate **1** (A), **2** (B), and **3** (C) at 405 nm; **4** (D) at 414 nm and **5** (E) at 394 nm. Data were calibrated using data from Supporting Information Figures S4, and normalised with respect to the x- and y-axes.

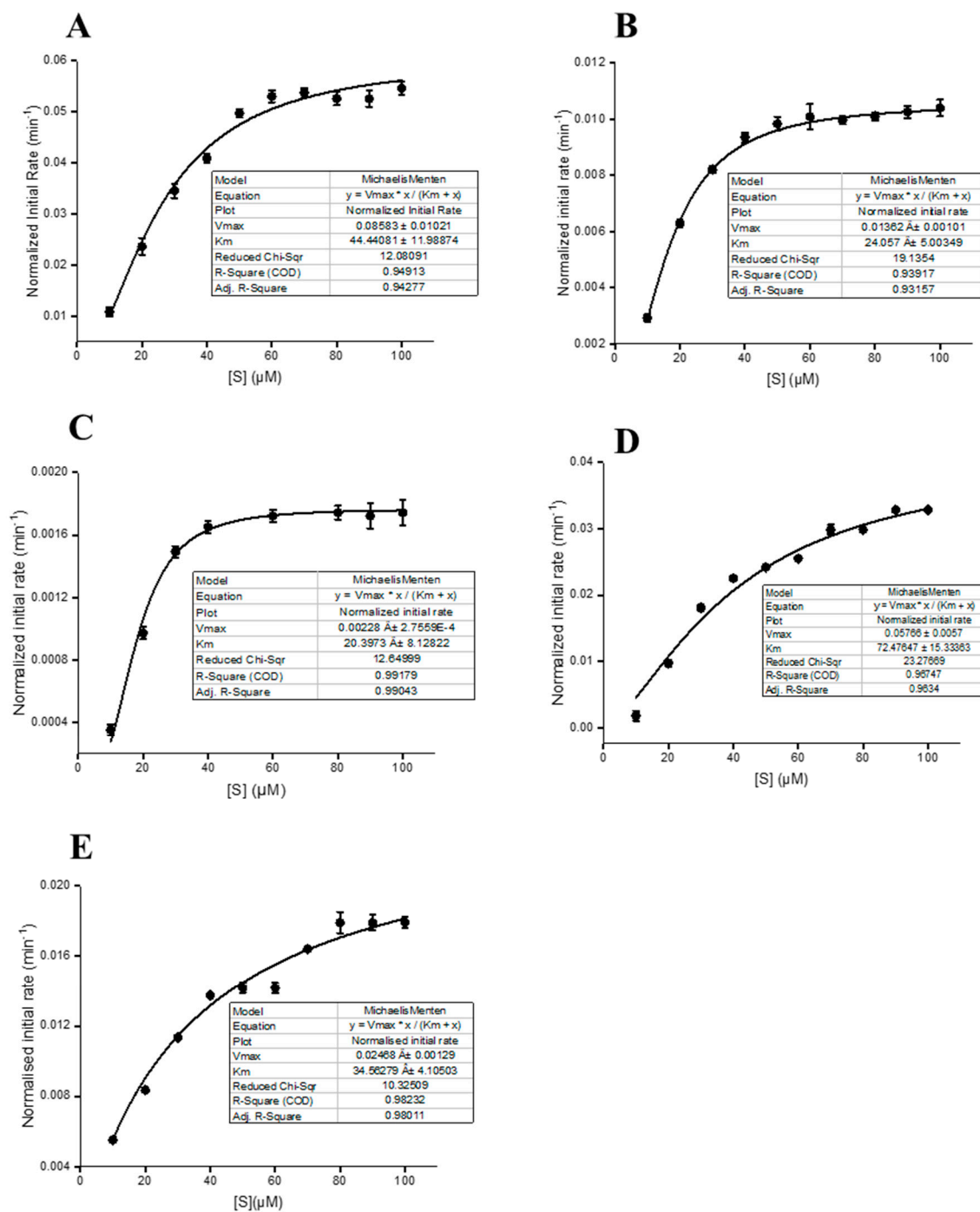


Figure S3. Graph of best fit Michaelis–Menten curves for the hydrolysis of silyl ether substrates by TF-Sil α -Strep against a range of substrate concentrations. Substrate **1** (A), **2** (B), **3** (C), **4** (D) and **5** (E).

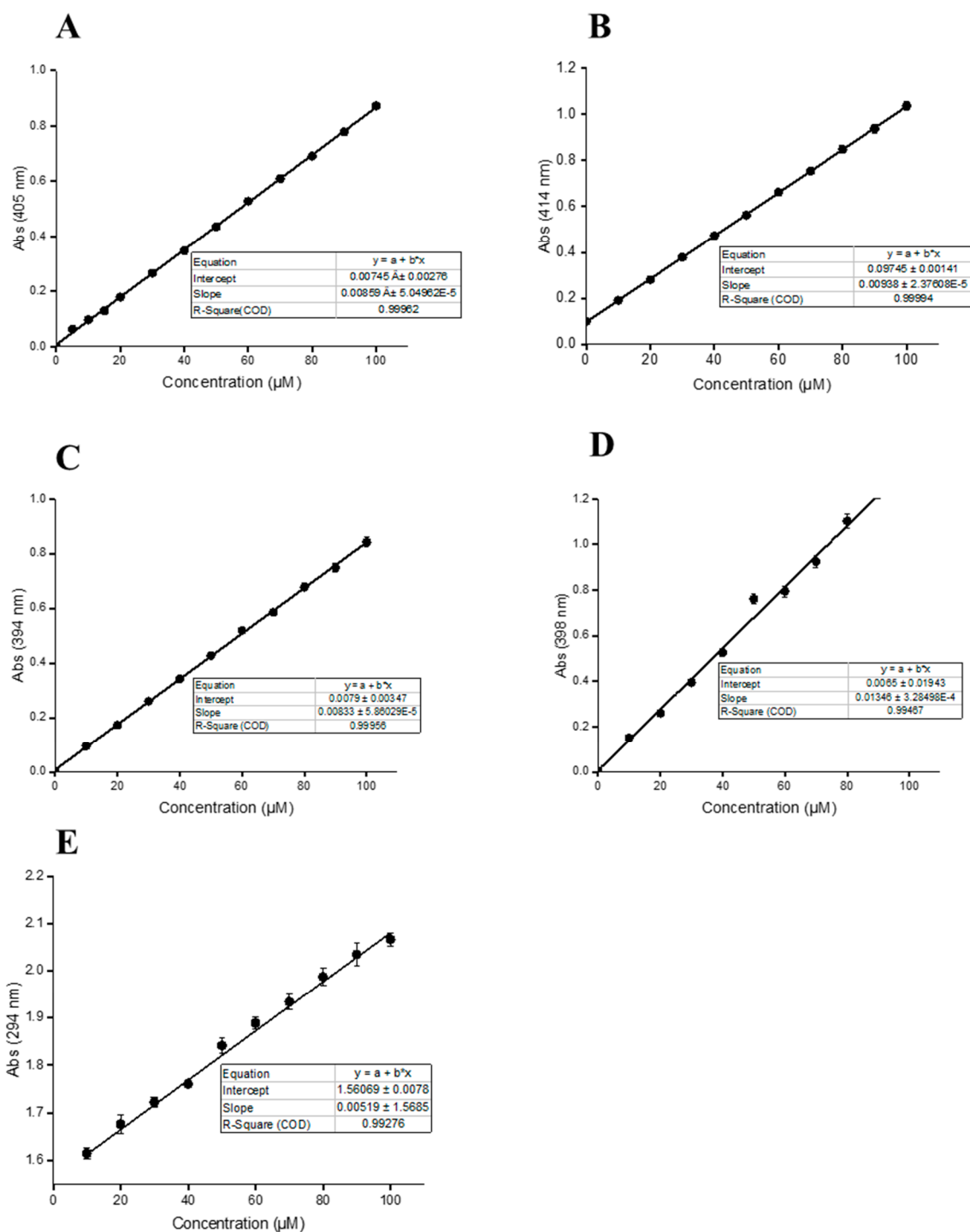


Figure S4. Calibration graph of UV-Vis absorption against concentration of (A) 4-nitrophenol for **1**, (B) 2-methyl-4-nitrophenol for **4**, (C) 3-methyl-4-nitrophenol for **5**, (D) 3-methoxy-4-nitrophenol for **6**, (E) 4-cyanophenol for **7**, in buffer (Tris (50 mM), NaCl (100 mM), pH 8.5, 10% (v/v) 1,4-dioxane).

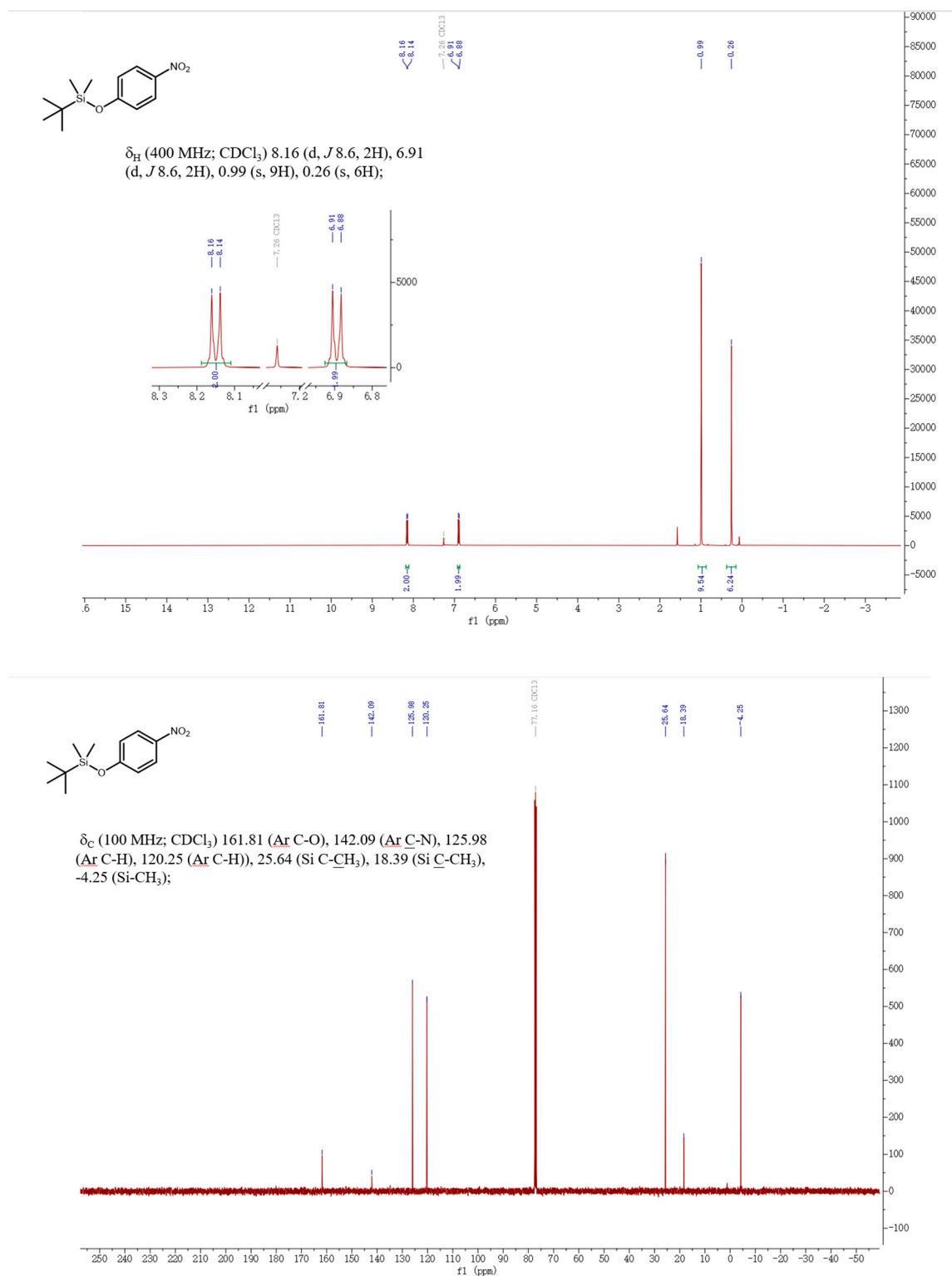


Figure S5. Calibrated NMR spectra for **1** showing ^1H (**top**) and ^{13}C (**bottom**) chemical shifts.

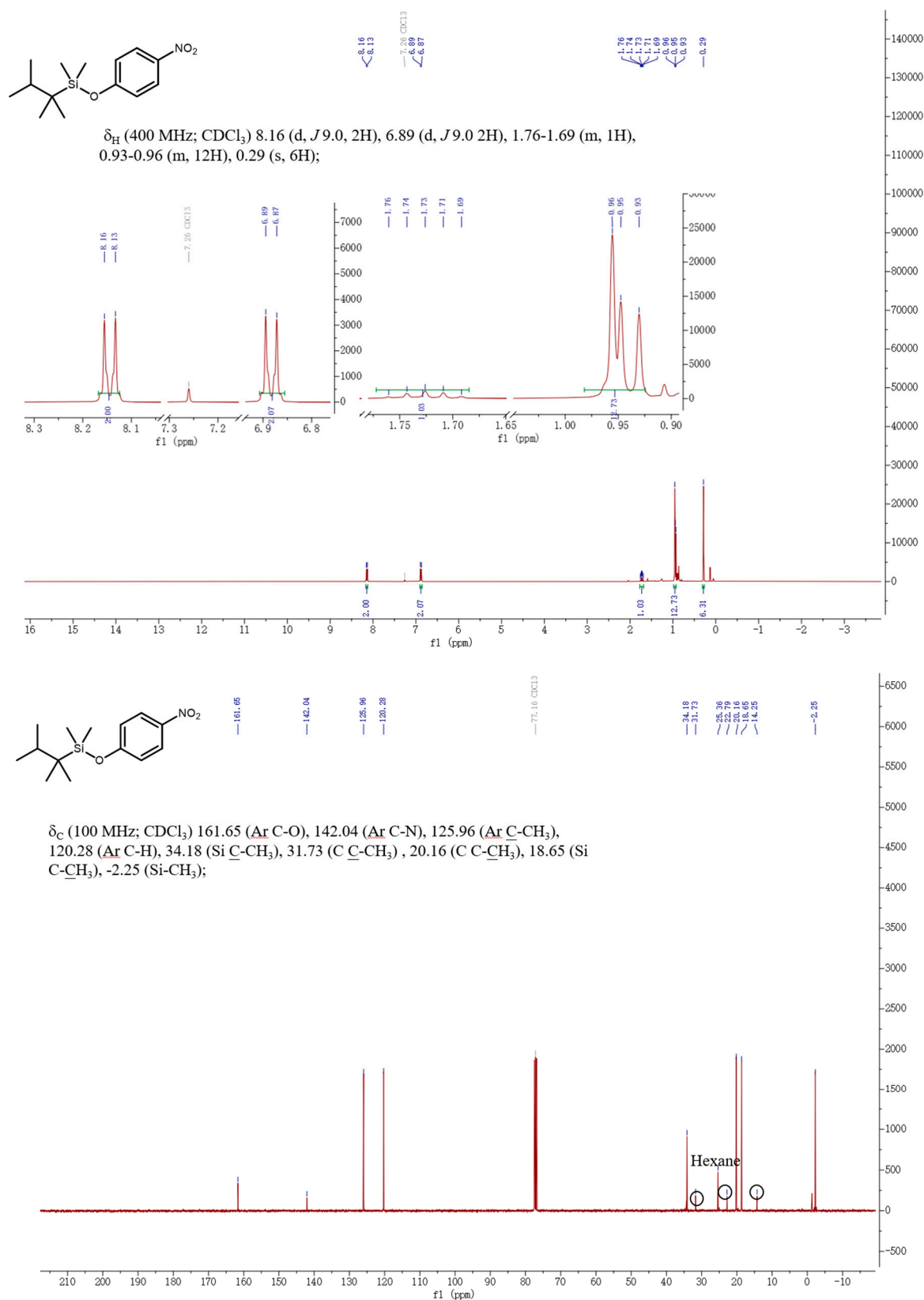


Figure S6. Calibrated NMR spectra for **2** showing ¹H (top) and ¹³C (bottom) chemical shifts.

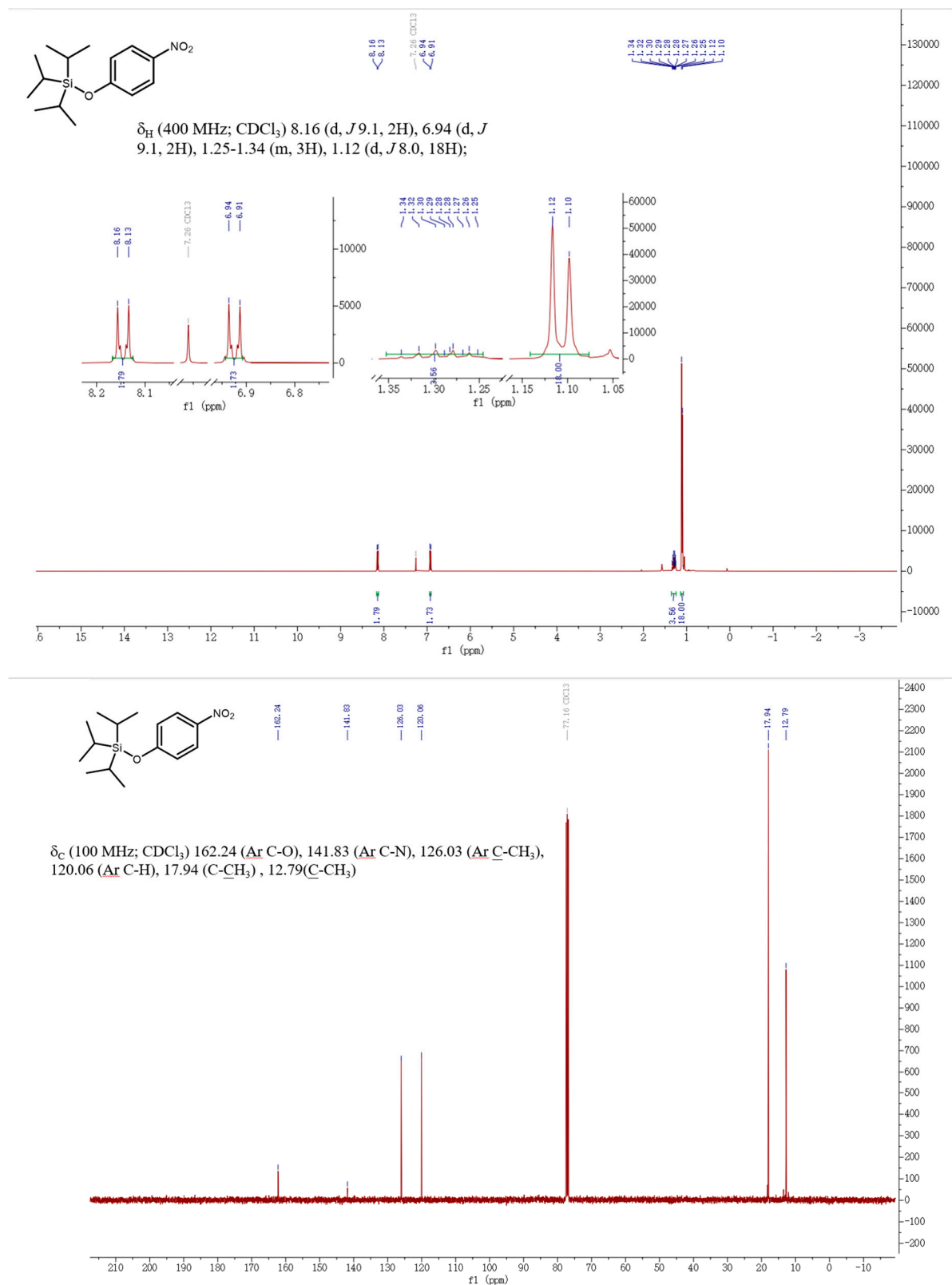


Figure S7. Calibrated NMR spectra for **3** showing 1H (top) and ^{13}C (bottom) chemical shifts.

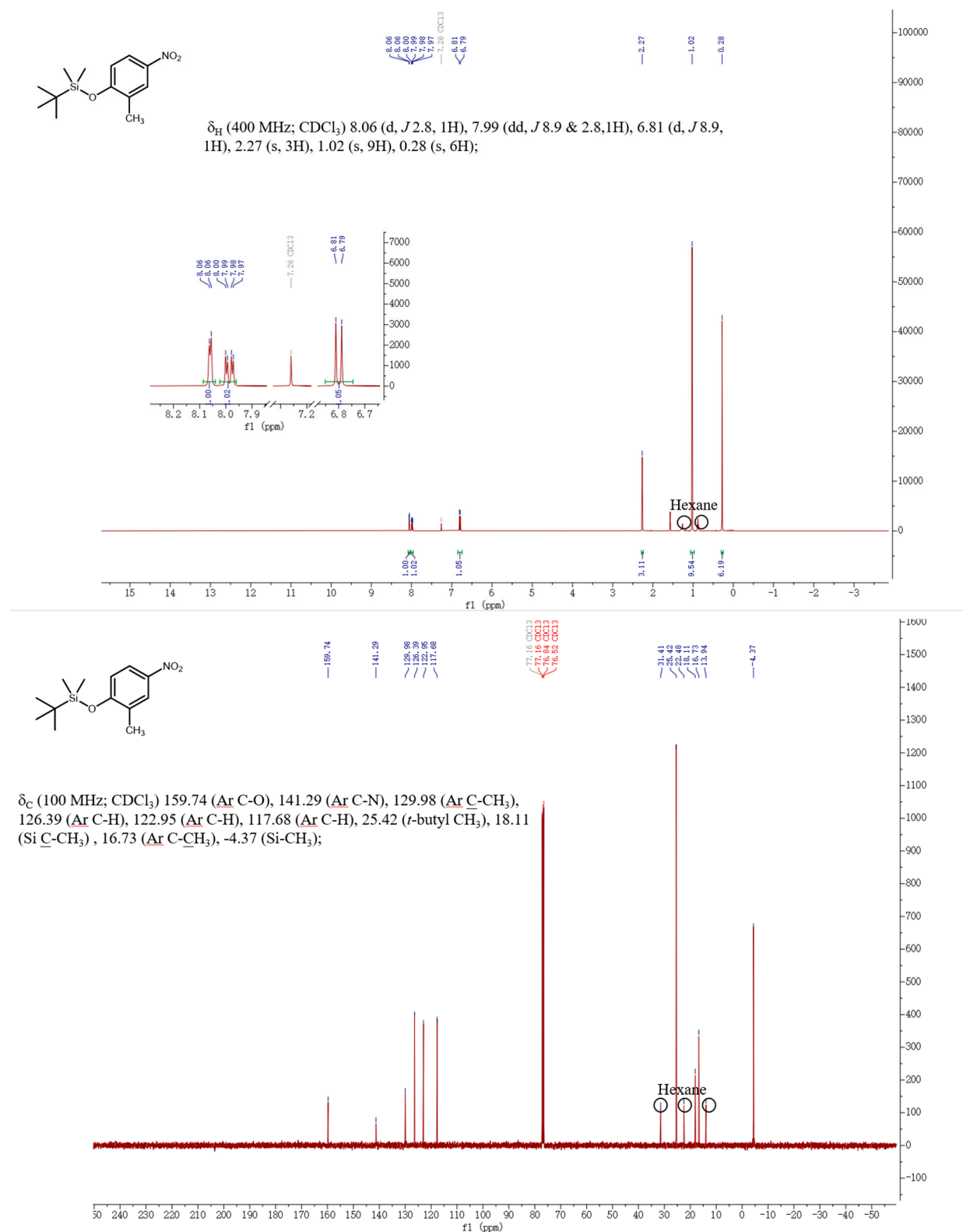


Figure S8. Calibrated NMR spectra for **4** showing ¹H (top) and ¹³C (bottom) chemical shifts.

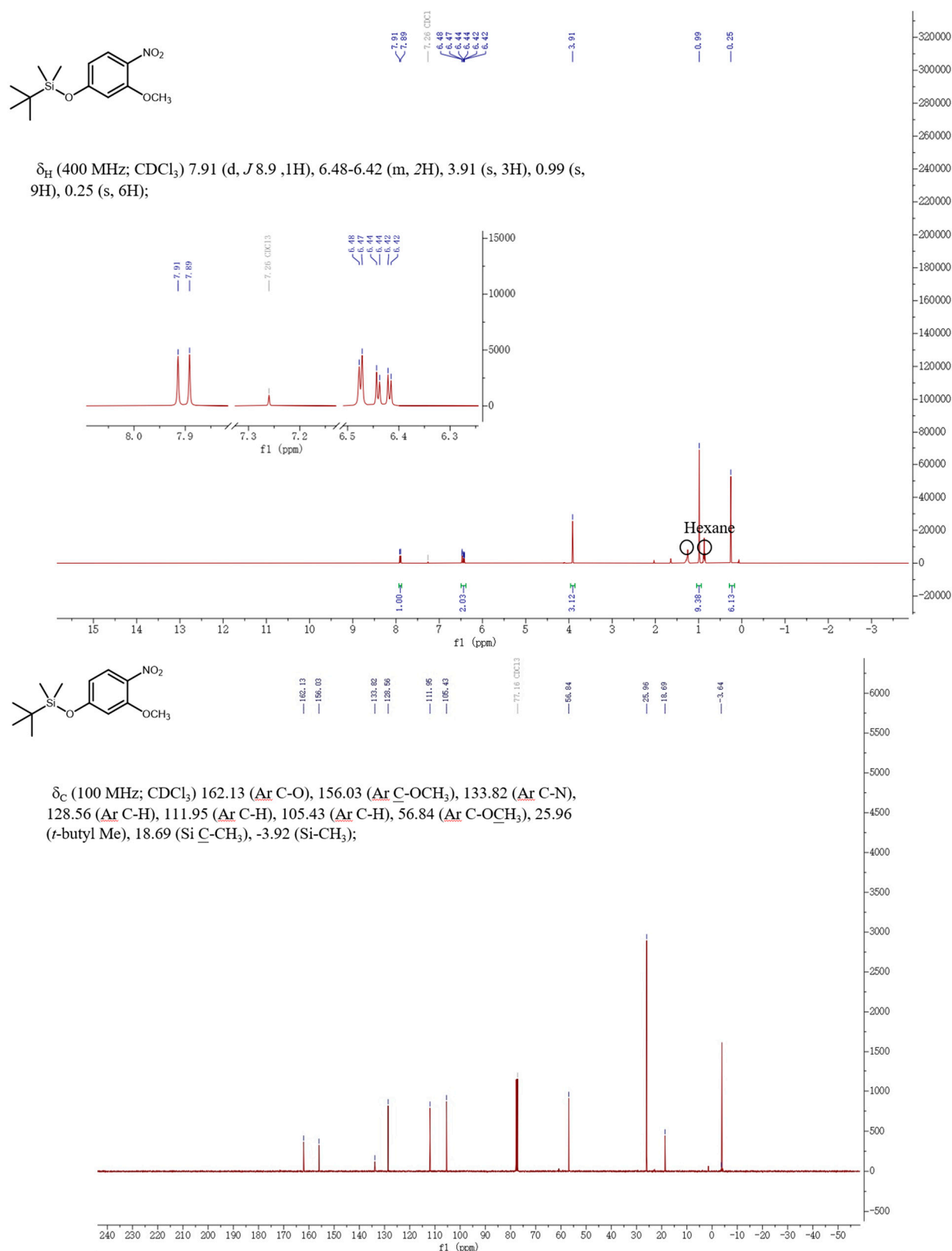


Figure S10. Calibrated NMR spectra for **6** showing ¹H (top) and ¹³C (bottom) chemical shifts.

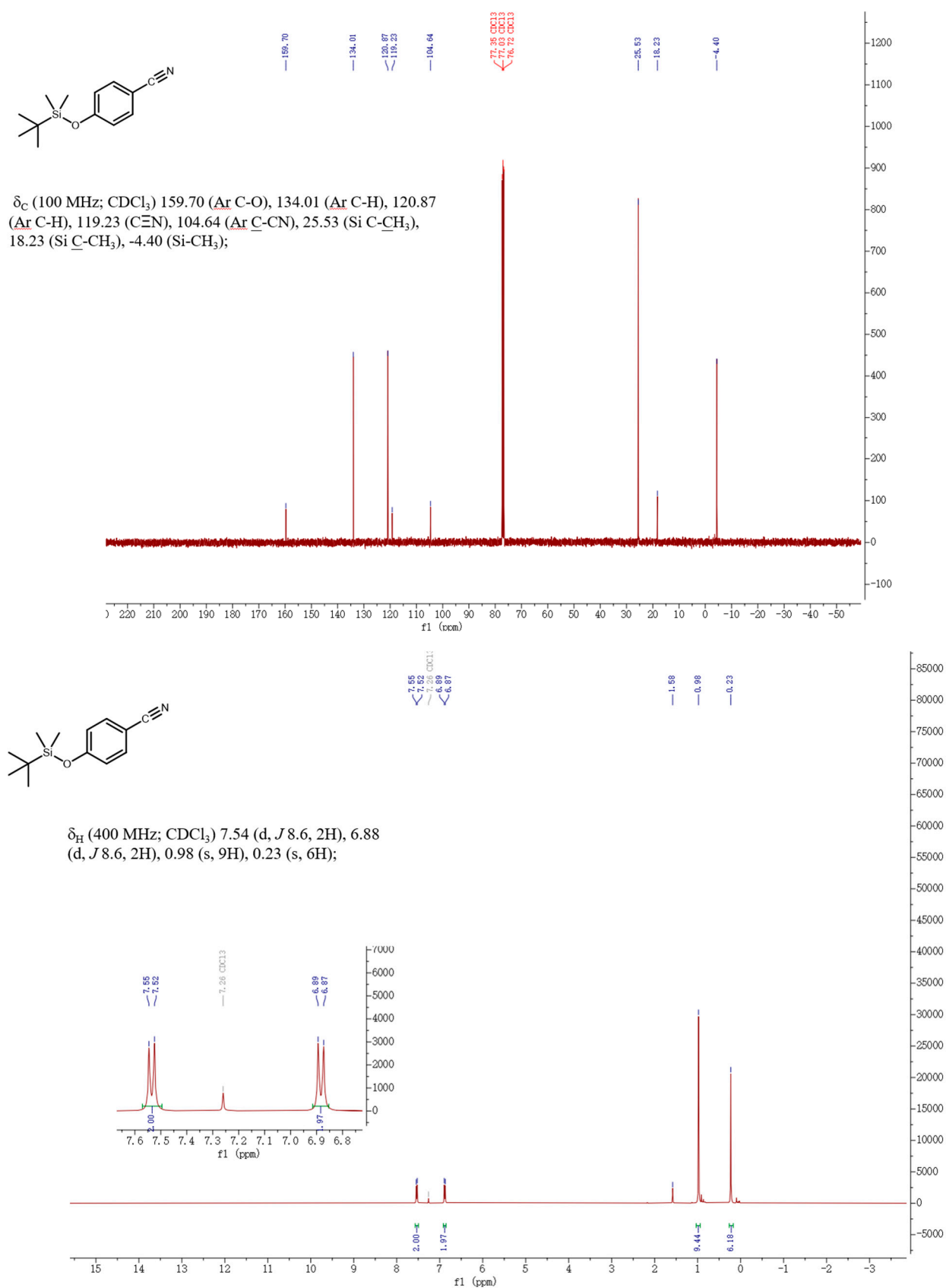


Figure S11. Calibrated NMR spectra for **7** showing 1H (top) and ^{13}C (bottom) chemical shifts.

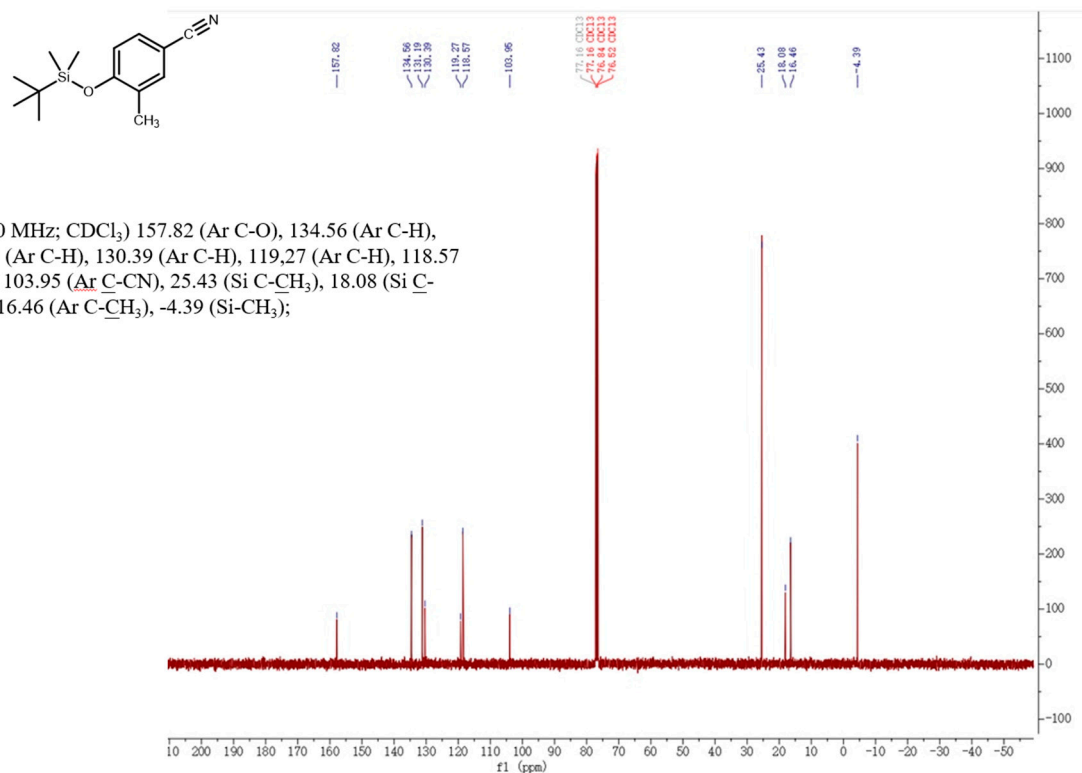
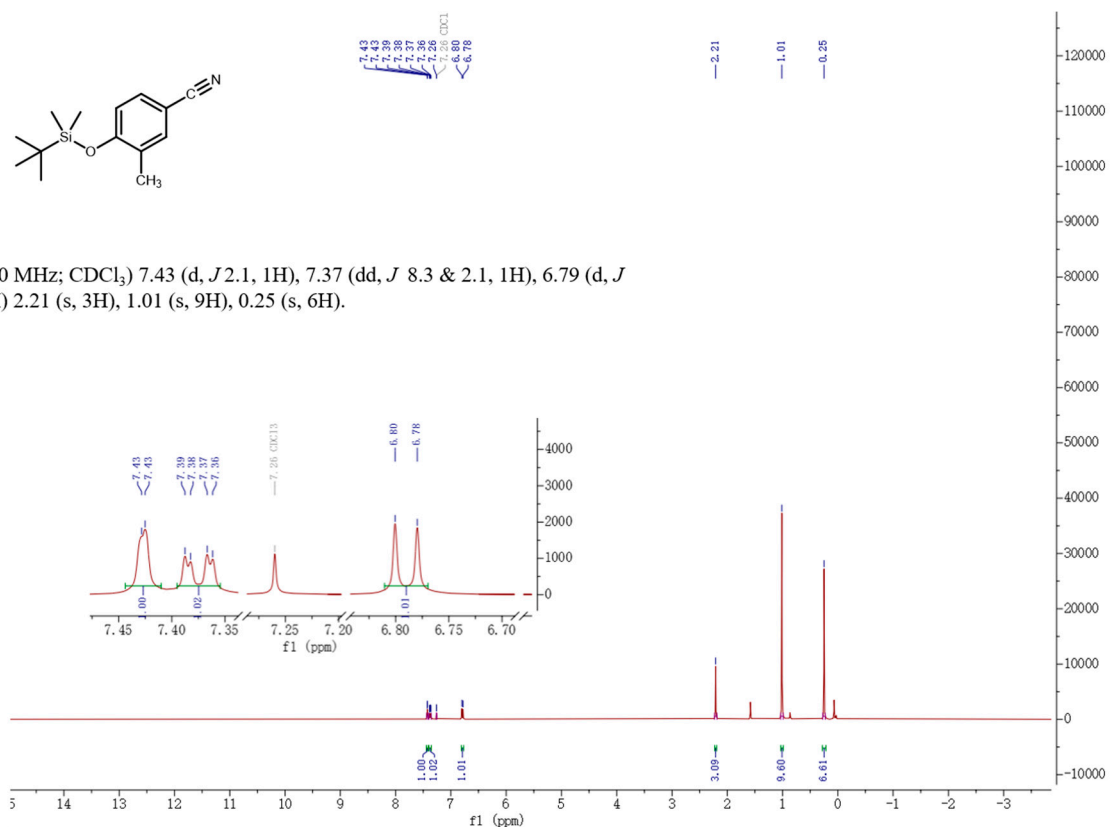


Figure S12. Calibrated NMR spectra for **8** showing 1H (top) and ^{13}C (bottom) chemical shifts.

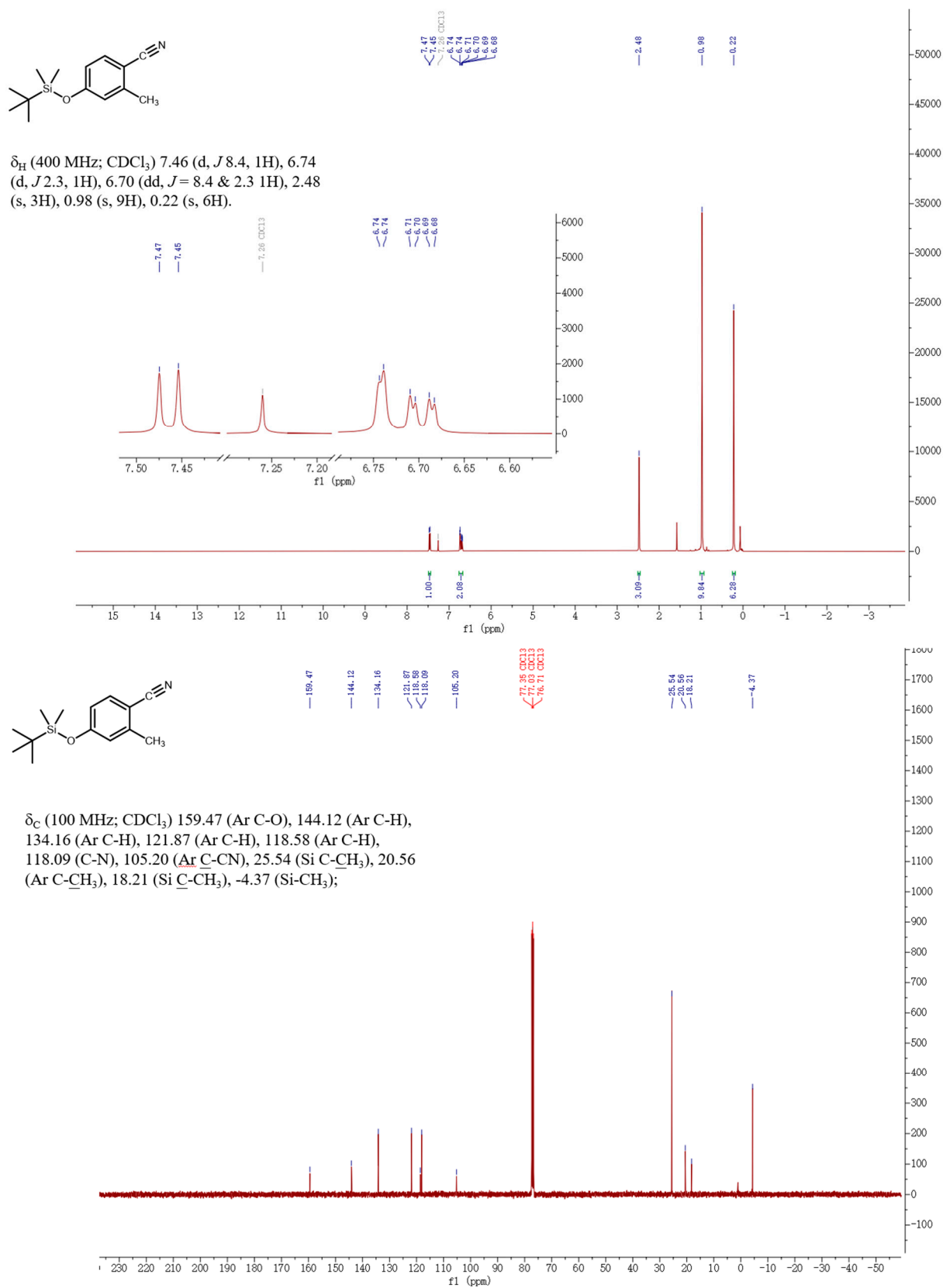


Figure S13. Calibrated NMR spectra for **9** showing ¹H (top) and ¹³C (bottom) chemical shifts.

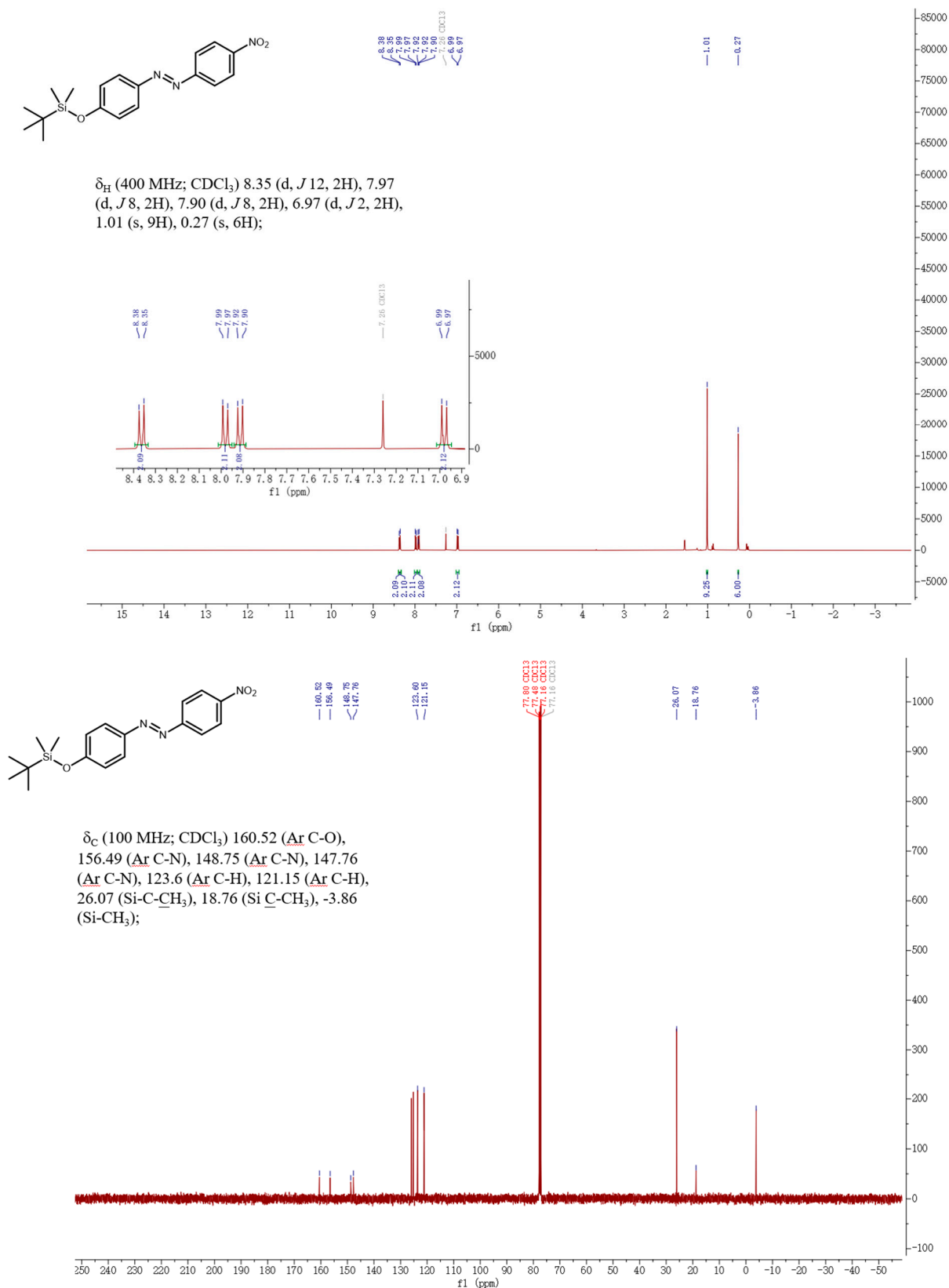


Figure S14. Calibrated NMR spectra for **10** showing ¹H (top) and ¹³C (bottom) chemical shifts.

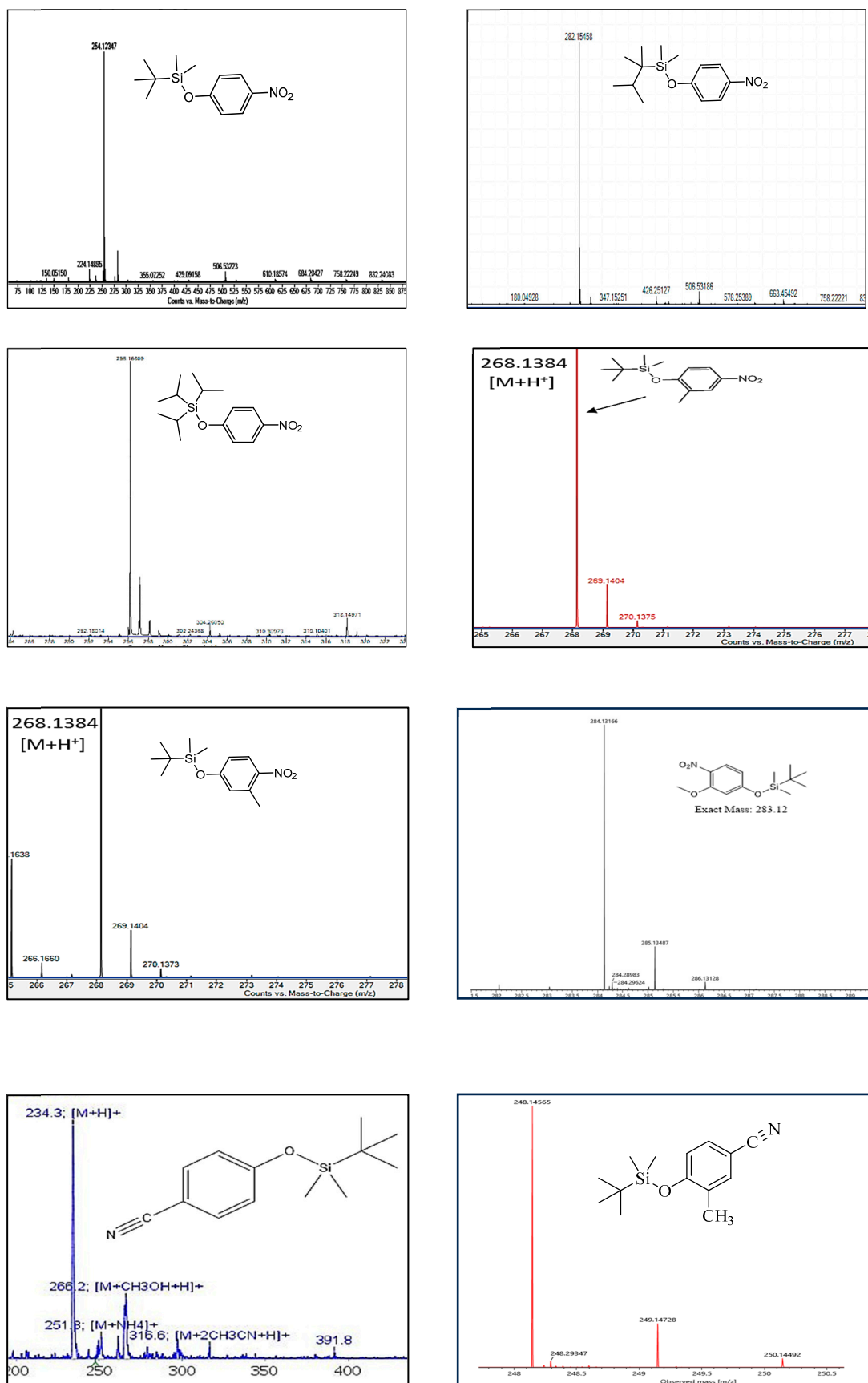


Figure S15. ESI+ mass spectra of substrates 1-10.

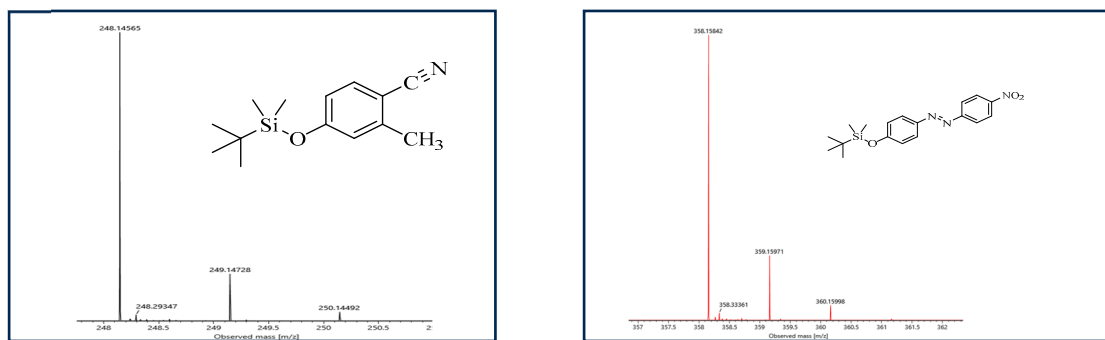


Figure S15 (continued). ESI+ mass spectra of substrates **1-10**.