Supporting information for

Platinum(II) Complexes with Novel Diisocyanide Ligands: Catalysts in Alkyne Hydroarylation

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Experimental Section.

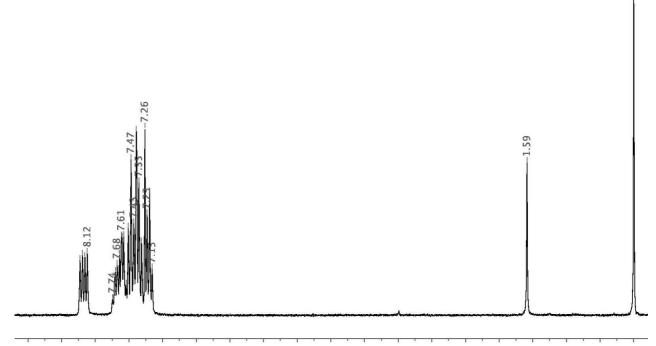
General remarks

NMR spectra were recorded at 298 K on a Bruker Avance-III 200 MHz (200.13 MHz for ¹H, 50.32 for ¹³C, 81.01 for ³¹P) or on a Bruker Avance DX-400 (400.13 MHz for ¹H and 100.61 for ¹³C); chemical shifts (δ) are reported in units of ppm relative to Si(CH₃)₄ or the residual solvent signals for ¹H and ¹³C NMR spectra, and to 85% H₃PO₄ for ³¹P NMR spectra. The FT-IR spectra were obtained with a Perkin Elmer Spectrum 100 spectrophotometer, with a resolution of 2 cm⁻¹. ESI-MS analyses were performed using a LCQ-Duo (Thermo-Finnigan) operating in positive ion mode. Instrumental parameters: capillary voltage 10 V, spray voltage 4.5 kV; capillary temperature 200 °C; mass scan range from 150 to 2000 amu; N2 was used as sheath gas; the He pressure inside the trap was kept constant. The pressure directly read by an ion gauge (in the absence of the N_2 stream) was 1.33×10^{-5} Torr. Sample solutions were prepared by dissolving the compounds in acetonitrile. Sample solutions were directly infused into the ESI source by a syringe pump at 8µL/min flow rate. Liquid chromatography (LC)-high resolution MS (HRMS) analyses were performed with an UHPLC system (Agilent Series 1200; Agilent Technologies, Palo Alto, CA, USA), consisting of vacuum degasser, auto-sampler, a binary pump coupled to a Quadrupole-Time of Flight mass analyzer (Agilent Series 6520; Agilent Technologies) with an ESI source, using nitrogen gas and operating in positive acquisition, with the following operation parameters: capillary voltage, 3500 V; nebulizer pressure, 35 psi; drying gas, 8 L min⁻¹; gas temperature, 350°C; fragmentor voltage in the range of 80-200 V; skimmer 65 V.

Analyses were performed injecting a total of 40 μ L (8 X 5 μ L injection volume) of samples by means of an injection program, varying each time the fragmentor voltage. The mobile phase was acetonitrile or acetonitrile 0.1 % HTFA at flow rate of 0.2 mL min⁻¹. Full scan mass spectra were recorded as centroid over the 50–3000 m/z range with a scan rate of 2 spectra/s. Mass spectra acquisition and data analysis was processed with Masshunter Workstation B 04.00 software (Agilent Technologies).

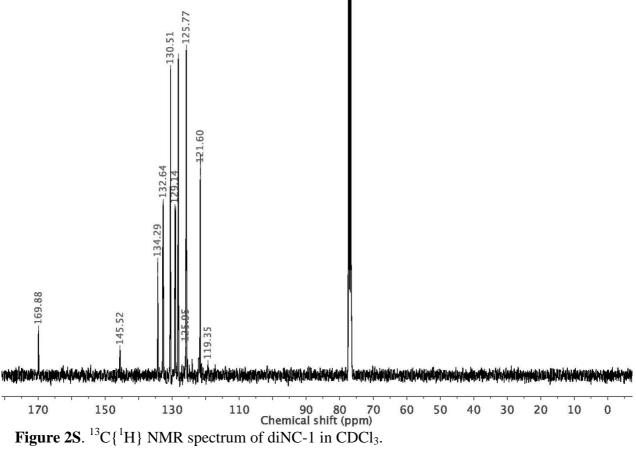
Complex	2	1'	2'
Formula	$C_{22} H_{12} Cl_2 N_2 O_4 Pt$	C ₂₂ H ₁₉ N ₂ O ₃ P Pt	$C_{48}H_{36}N_4O_8Pt_2\cdot CH_2Cl_2$
Molecular weight	634.33	585.45	1271.91
Crystal system	Monoclinic	Monoclinic	Monoclinic
Space group	P2 ₁ /c	P2 ₁ /n	C 2/c
a/Å	14.7441(14)	11.871(1)	26.097(4)
<i>b</i> /Å	8.6683(8)	8.358(1)	12.1200(14)
c/Å	16.881(3)	21.583(2)	20.158(3)
α/°	90.00	90.00	90.00
β/°	98.386(14)	93.142(5)	130.299(9)°
γ/°	90.00	90.00	90.00
Volume, Å ³	2134.4(5)	2138.2(4)	4862.8(13)
Ζ	4	4	4
$D_{calc}/g \text{ cm}^{-3}$	1.974	1.819	1.737
F(000)	1208	1128	2456
μ(Mo- Kα)/mm ⁻¹	6.857	6.66	5.91
Reflections collected	42931	40575	49583
Unique reflections	4652	3976	5306
Observed reflections [I>2 σ (<i>I</i>)]	3468 [R _{int} = 0.047]	3368 [R _{int} = 0.027]	3170 [R _{int} = 0.078]
$R\left[I > 2\sigma(I)\right]$	$R_1(F) = 0.0308,$ $wR_2(F^2) = 0.0318$	$R_1(F) = 0.0290,$ $wR_2(F^2) = 0.0720$	$R_1(F) = 0.0507,$ $wR_2(F^2) = 0.0504$
R [all data]	$R_1(F) = 0.0677,$ $wR_2(F^2) = 0.0363$	$R_1(F) = 0.0369,$ $wR_2(F^2) = 0.0763$	$R_1(F) = 0.1421,$ $wR_2(F^2) = 0.0645$
Goodness of fit on F ²	0.990	1.053	1.059

 Table S1. Selected crystallographic data for complexes 2, 1' and 2'



-0.00

7.5 7.0 5.0 4.5 4.0 3.5 Chemical shift (ppm) 9.0 8.5 8.0 6.5 6.0 5.5 3.0 2.5 2.0 1.5 1.0 0.5 0.0 Figure 1S. ¹H NMR spectrum of diNC-1 in CDCl₃.



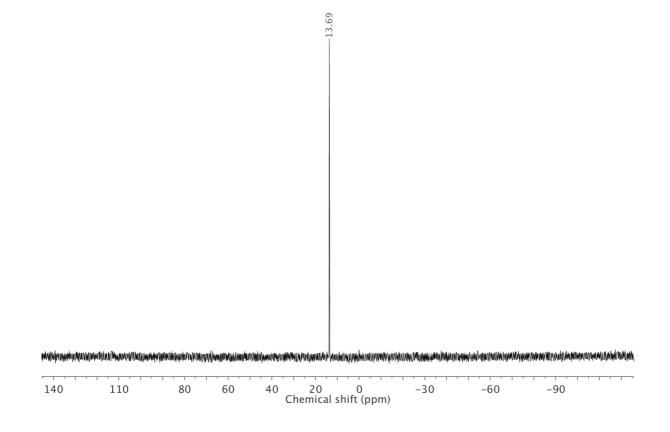
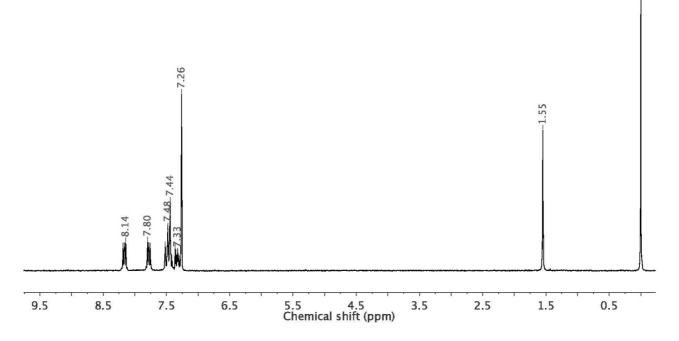
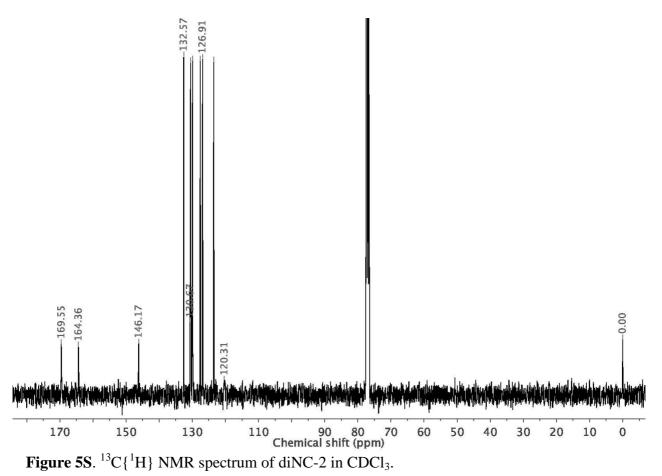


Figure 38. ³¹P{¹H} NMR spectrum of diNC-1 in CDCl₃.



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Figure 4S. ¹H NMR spectrum of diNC-2 in CDCl₃.



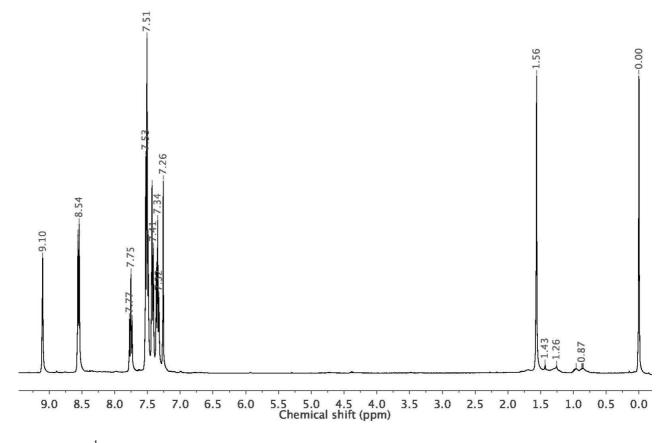
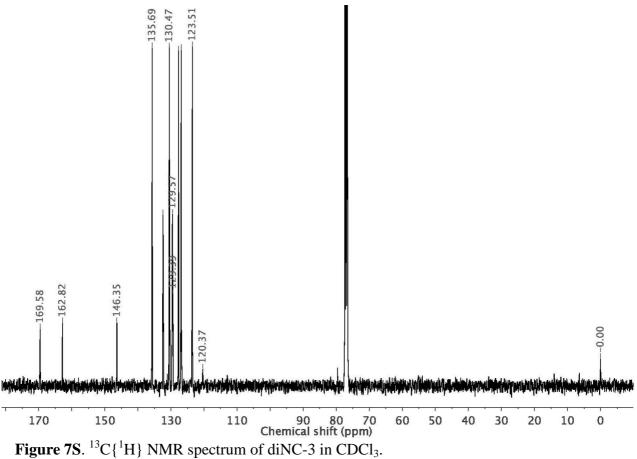


Figure 6S. ¹H NMR spectrum of diNC-3 in CDCl₃.



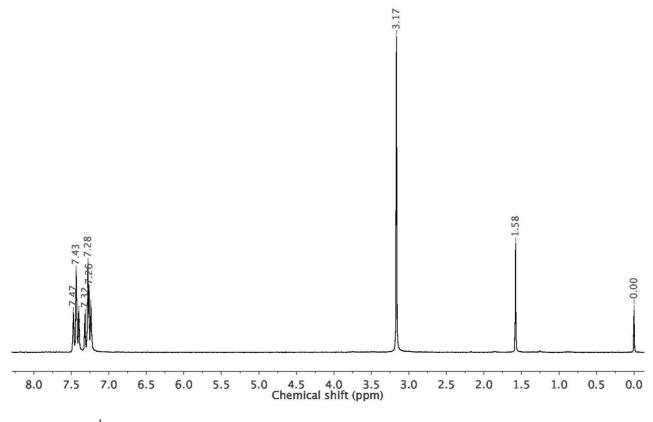


Figure 8S. ¹H NMR spectrum of diNC-4 in CDCl₃.

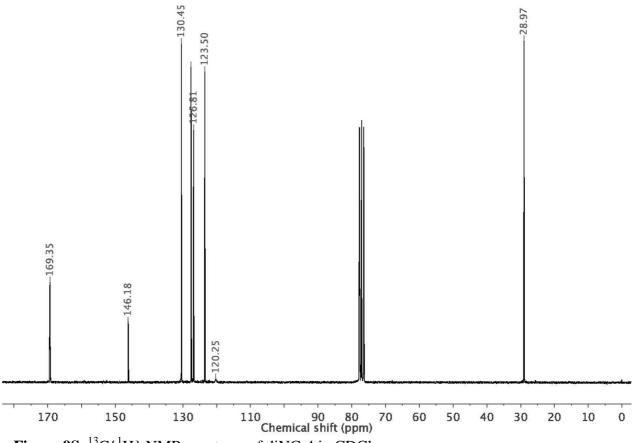


Figure 9S. ¹³C{¹H} NMR spectrum of diNC-4 in CDCl₃.

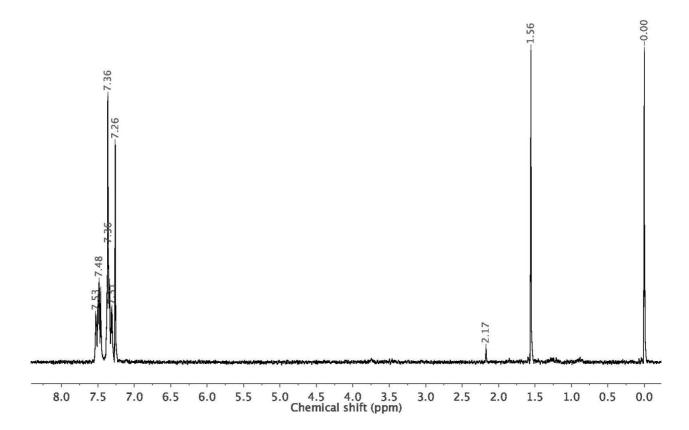
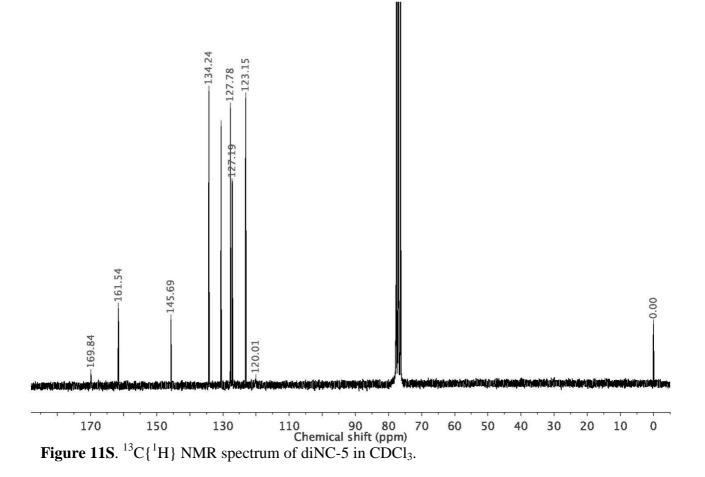
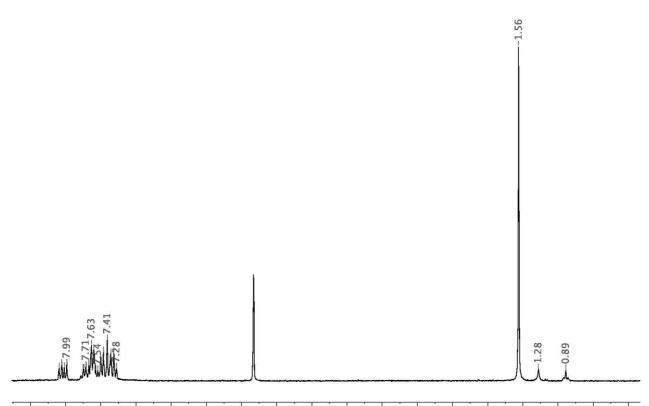
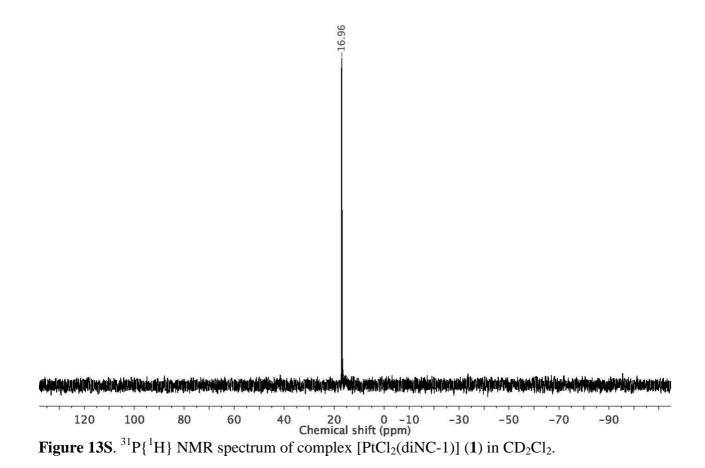


Figure 10S. ¹H NMR spectrum of diNC-5 in CDCl₃.





5.0 4.5 4.0 3.5 Chemical shift (ppm) 7.5 8.5 8.0 7.0 6.5 6.0 5.5 2.0 1.5 3.0 2.5 1.0 0.5 0.0 Figure 12S. ¹H NMR spectrum of complex [PtCl₂(diNC-1)] (1) in CD₂Cl₂.



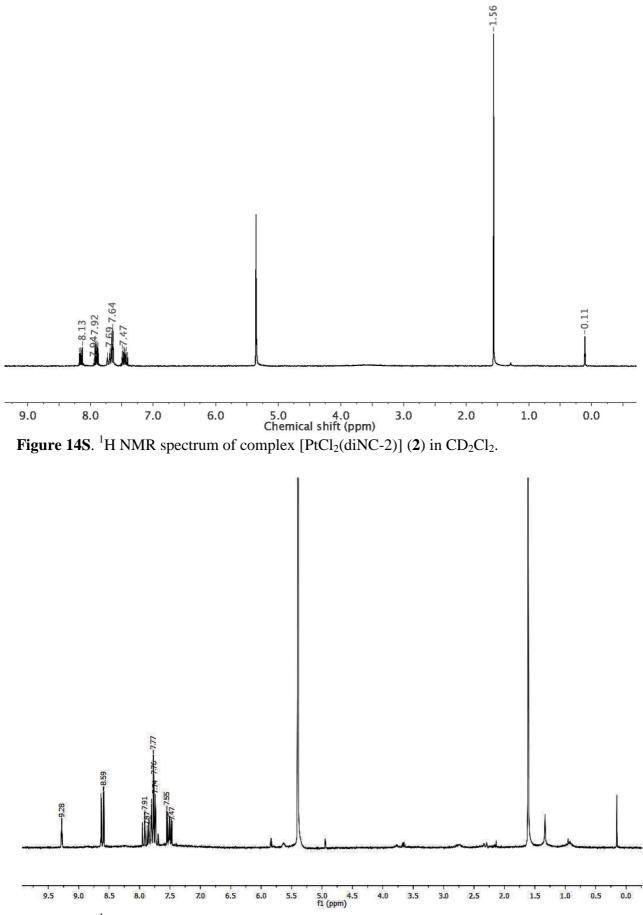
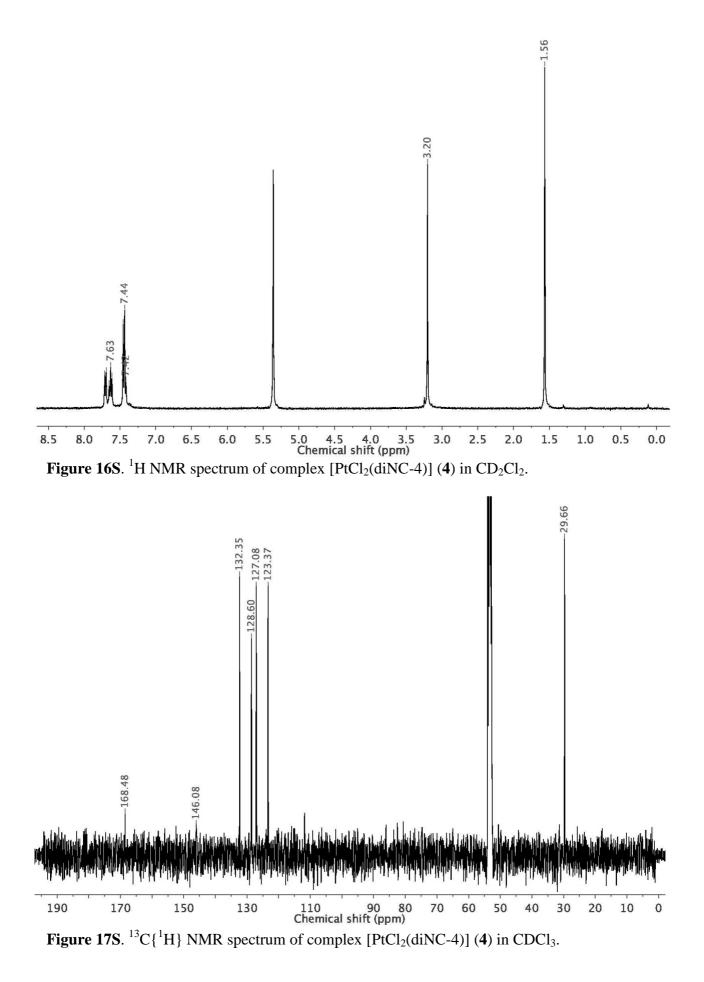
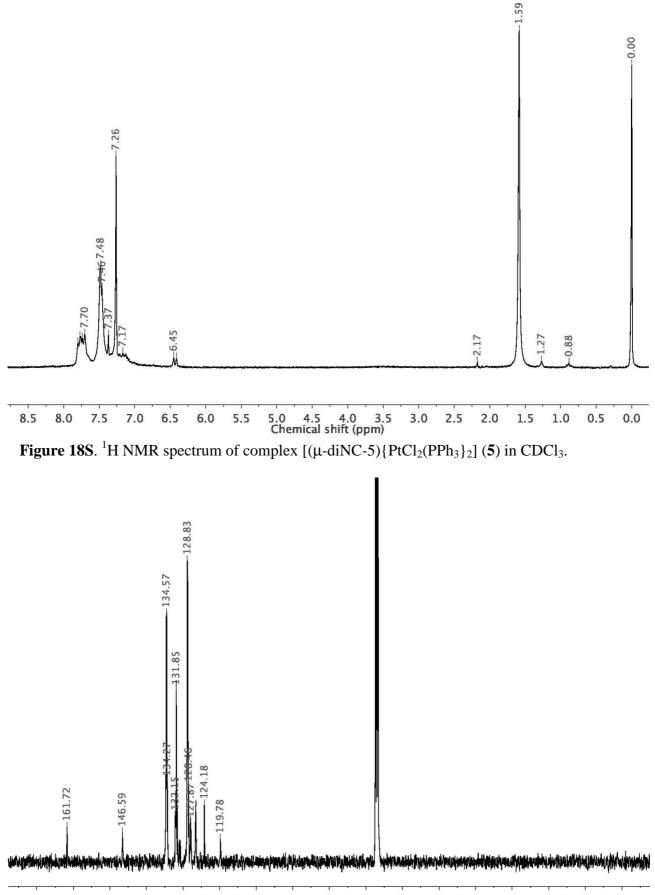
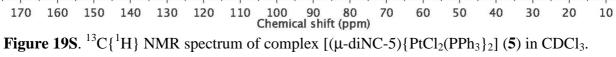
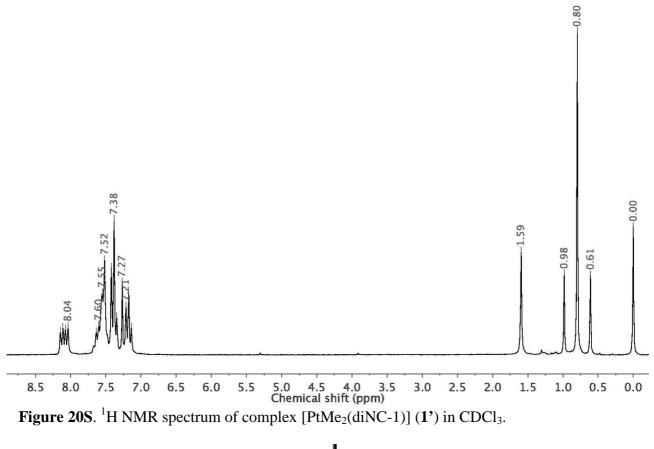


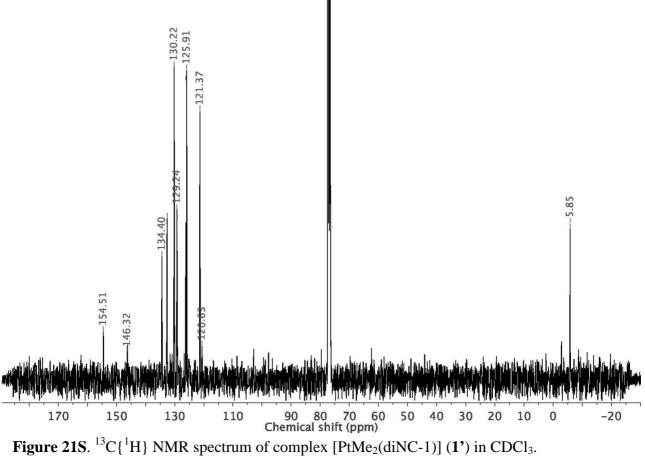
Figure 15S. ¹H NMR spectrum of complex [PtCl₂(diNC-3)] (3) in CD₂Cl₂.











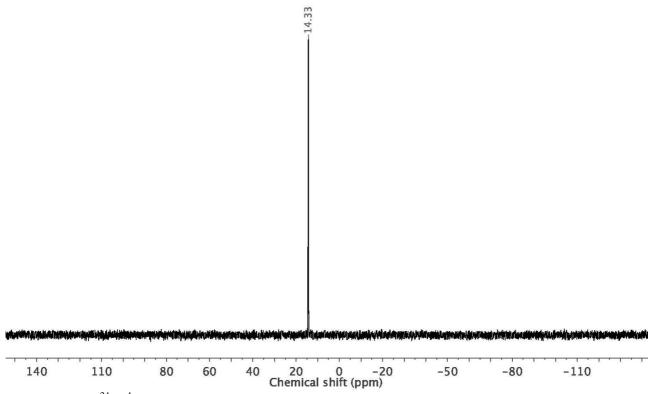


Figure 22S. ³¹P{¹H} NMR spectrum of complex [PtMe₂(diNC-1)] (1') in CDCl₃.

