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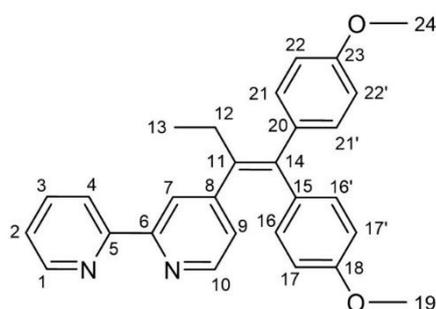
Metallo drugs against Breast Cancer: Combining the Tamoxifen Vector with Platinum(II) and Palladium(II) Complexes

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1. Characterisation



1.1. 4-[1,1-Bis(4-methoxyphenyl)but-1-en-2-yl]-2,2'-bipyridine (L) (4)

$M(C_{28}H_{26}Cl_2N_2O_2) = 422.53 \text{ g mol}^{-1}$. $^1\text{H NMR}$ (400 MHz, CDCl_3), δ (ppm): 8.67 (m, 1H, 1-CH), 8.40 (d, $^3J_{\text{HH}} = 8.6 \text{ Hz}$, 1H, 10-CH), 8.35 (d, $^3J_{\text{HH}} = 8.0 \text{ Hz}$, 1H, 4-CH), 8.30 (s, 1H, 7-CH), 7.80 (td, $^3J_{\text{HH}} = 7.8$, $^3J_{\text{HH}} = 1.8 \text{ Hz}$, 1H, 3-CH), 7.29 (ddd, $^3J_{\text{HH}} = 7.6$, $^3J_{\text{HH}} = 4.7$, $^4J_{\text{HH}} = 1.2 \text{ Hz}$, 1H, 2-CH), 7.16 (d, $^3J_{\text{HH}} = 8.5 \text{ Hz}$, 2H, 21,21'-CH), 6.93 (dd, $^3J_{\text{HH}} = 5.0$, $^4J_{\text{HH}} = 1.7 \text{ Hz}$, 1H, 9-CH), 6.89 (d, $^3J_{\text{HH}} = 8.6 \text{ Hz}$, 2H, 22, 22'-CH), 6.84 (d, 2H, $^3J_{\text{HH}} = 8.6 \text{ Hz}$, 2H, 16,16'-CH), 6.56 (d, $^3J_{\text{HH}} = 8.6 \text{ Hz}$, 2H, 17,17'-CH), 3.83 (s, 3H, 24-CH₃), 3.66 (s, 3H, 19-CH₃), 2.59 (q, $^3J_{\text{HH}} = 7.3 \text{ Hz}$, 2H, 12-CH₂), 0.97 (t, $^3J_{\text{HH}} = 7.4 \text{ Hz}$, 3H, 13-CH₃). $^{13}\text{C}\{^1\text{H}\}$ NMR (101 MHz, CDCl_3), δ (ppm): 158.6 (23-C), 158.0 (18-C), 156.3 (5-C), 155.9 (6-C), 152.3 (8-C), 149.1 (1-CH), 148.7 (10-CH), 140.2 (14-C), 138.7 (11-C), 136.9 (3-CH), 135.5 (20-C), 134.8 (15-C), 132.0 (16, 16'-CH), 130.5 (21,21'-CH), 125.8 (9-CH), 123.6 (2-CH), 121.3 (7-CH), 121.1 (4-CH), 113.6 (22, 22'-CH), 113.1 (17, 17'-CH), 55.3 (24-CH₃), 55.0 (19-CH₃), 28.4 (12-CH₂), 13.6 (13-CH₂). **Elemental analysis:** $\text{C}_{28}\text{H}_{26}\text{N}_2\text{O}_2 \cdot \text{H}_2\text{O}$ calc. (%) C 76.34 H 6.41 N 6.36, found (%) C 76.75 H 6.38 N 6.17.

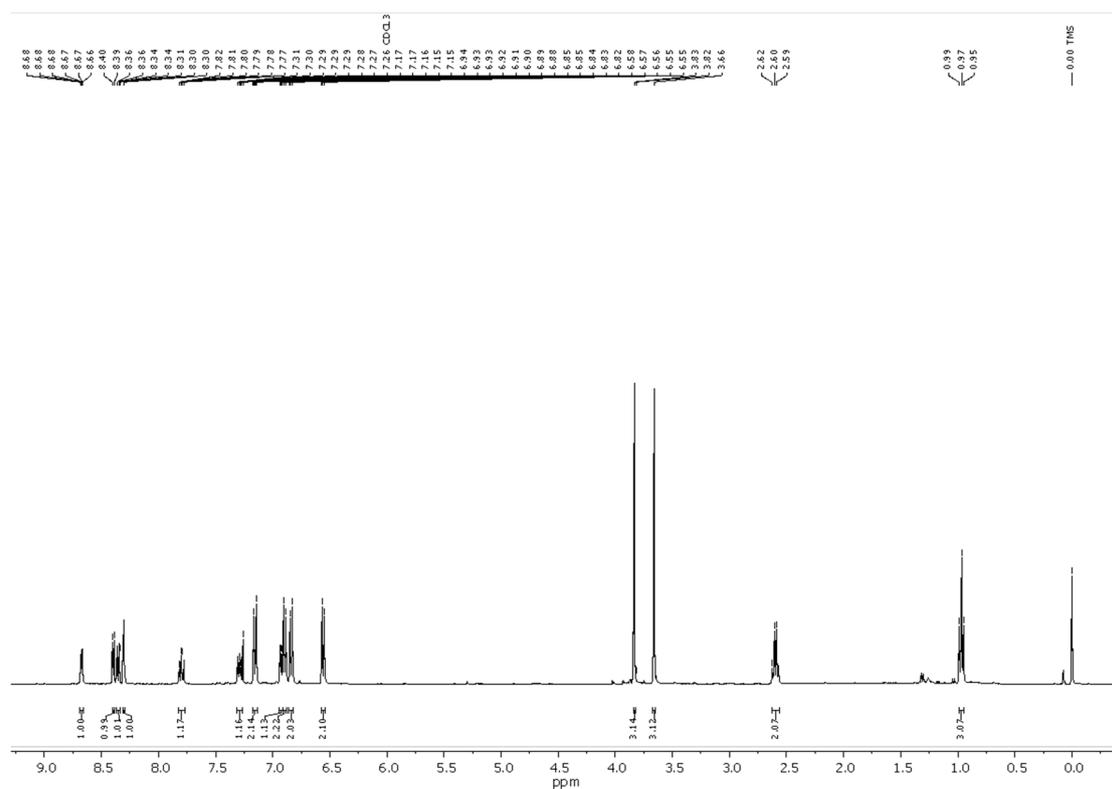


Figure S1. $^1\text{H NMR}$ spectrum of compound 4 in CDCl_3 .

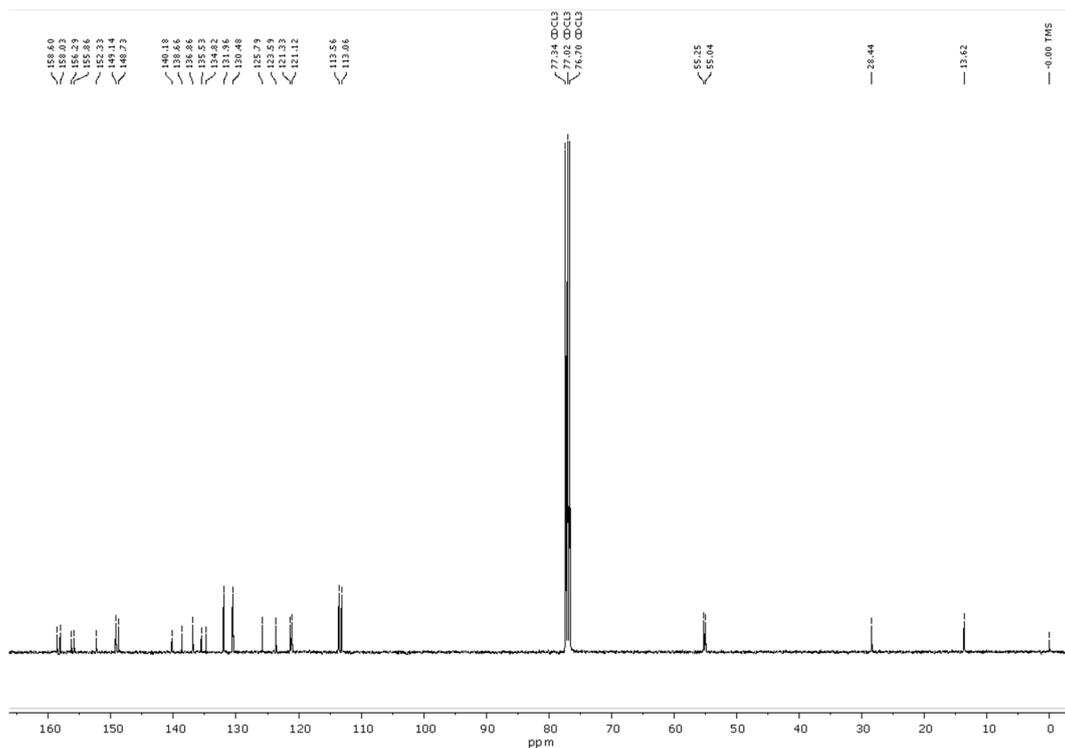


Figure S2. $^{13}\text{C}\{^1\text{H}\}$ NMR spectrum of compound **4** in CDCl_3 .

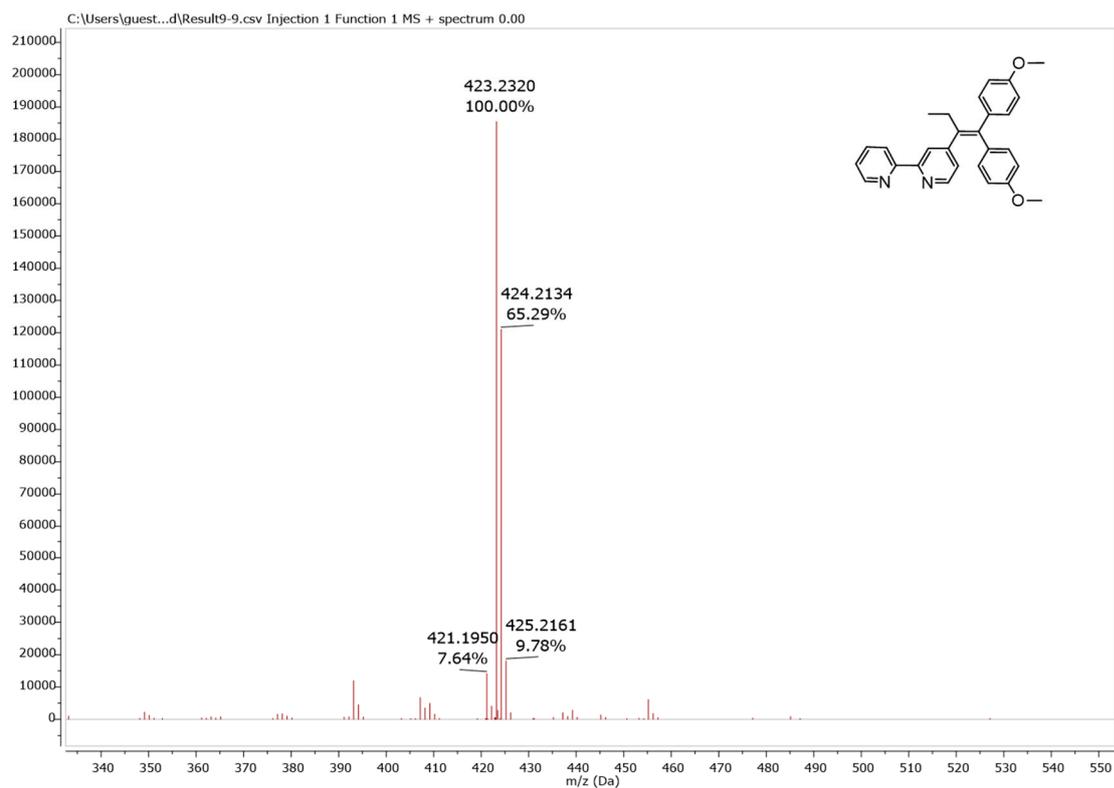


Figure S3. HR-ESI-MS (positive mode, acetonitrile) of **4**: m/z $[\text{M}+\text{H}]^+ = 423.2072$ (calc.), 423.2320 (found).

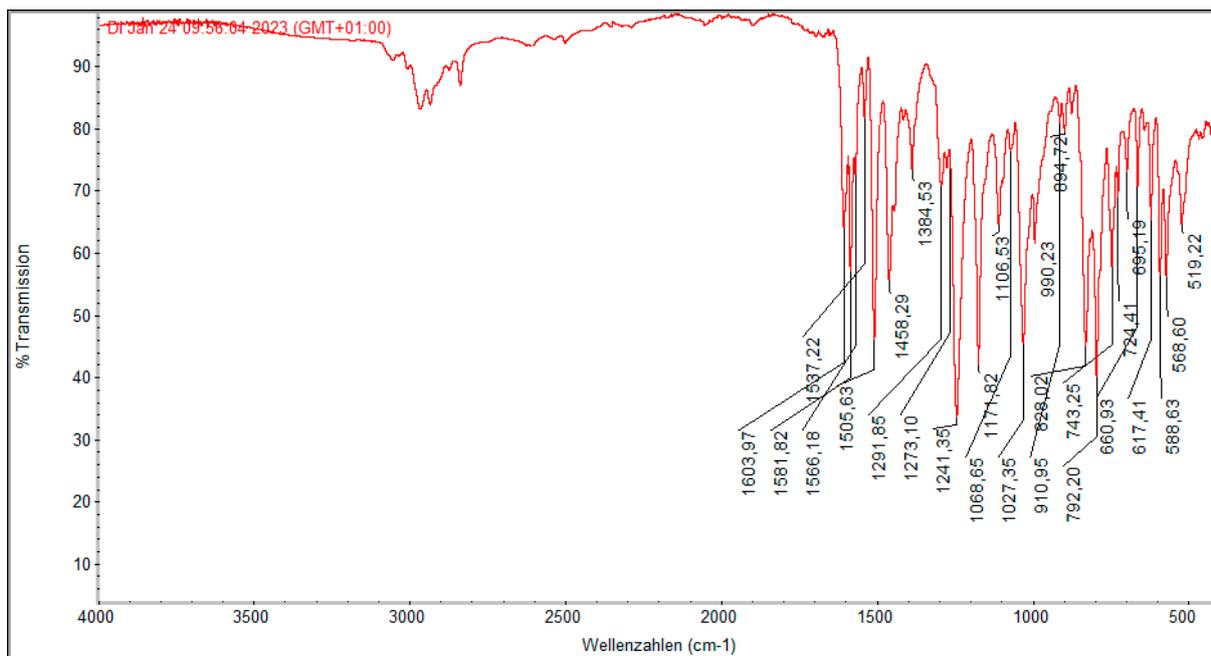


Figure S4. IR spectrum of **4**: 3050–2084 (w, $C_{alk}-H$), 1604 (m, $\nu(C=C)$), 1582 (m, $\nu(C=C)$), 1537 (w, $\nu(C=C)$), 1506 (s, $C_{arom}-H$ in-plane bending), 1458 (w, $C_{arom}-H$ in-plane bending), 1384 (w, $C_{arom}-H$ in-plane bending), 1292 (w), 1273 (w, $C_{arom}-O$), 1241 (s, $C_{arom}-O$), 1172 (s, $C_{arom}-O$), 1107 (m), 1069 (w), 1027 (s, $C_{alk}-O$), 828 (m, out-of-plane bending), 792 (m, out-of-plane bending) 743 (s, out-of-plane bending), 724 (w), 661 (w), 627(w), 589 (w), 567 (w), 519 (w).

1.2. $[PtCl_2(L-\kappa^2N,N')]$ (**5**)

$M(C_{28}H_{26}Cl_2N_2O_2Pt) = 687.10 \text{ g mol}^{-1}$, 1H NMR (400 MHz, $CDCl_3$), δ (ppm): 9.68 (d, $^3J_{HH} = 5.9$ Hz, 1H, 1-CH), 9.44 (d, $^3J = 6.1$ Hz, 1H, 10-CH), 8.07 (td, $^3J_{HH} = 7.8$, $^3J_{HH} = 1.5$ Hz, 1H, 3-CH), 7.66 (d, $^3J_{HH} = 8.0$ Hz, 1H, 4-CH), 7.63 (d, $^3J_{HH} = 1.9$ Hz, 1H, 7-CH), 7.47 (ddd, $^3J_{HH} = 7.4$, $^3J_{HH} = 5.8$, $^4J_{HH} = 1.3$ Hz, 1H, 2-CH), 7.30 (dd, $^3J_{HH} = 6.2$, $^4J_{HH} = 1.9$ Hz, 1H, 9-CH), 7.16 (d, $^3J_{HH} = 8.6$ Hz, 2H, 21,21'-CH), 6.91 (d, $J_{HH} = 8.7$ Hz, 2H, 22,22'-CH), 6.86 (d, $^3J_{HH} = 8.7$ Hz, 2H, 16,16'-CH), 6.66 (d, $^3J_{HH} = 8.7$ Hz, 2H, 17,17'-CH), 3.85 (s, 3H, 24-CH₃), 3.69 (s, 3H, 19-CH₃), 2.67 (q, $^3J_{HH} = 7.4$ Hz, 2H, 12-CH₂), 1.03 (t, $^3J_{HH} = 7.4$ Hz, 3H, 13-CH₃). $^{13}C\{^1H\}$ NMR (101 MHz, $CDCl_3$), δ (ppm): 159.3 (23-C), 159.1 (18-C), 157.2 (8-C), 156.0 (6-C), 155.3 (5-C), 149.6 (1-CH), 148.9 (10-CH), 144.6 (14-C), 139.2 (3-CH), 136.2 (20-C), 134.3 (15-C), 133.8 (11-C), 132.2 (16,16'-CH), 130.4 (21,21'-CH), 127.6 (9-CH), 126.7 (2-CH), 124.2 (7-CH), 122.3 (4-CH), 113.9 (22,22'-CH), 113.8 (17,17'-CH), 55.3 (24-CH₃), 55.2 (19-CH₃), 27.7 (12-CH₃), 13.8 (13-CH₂). 1H NMR (400 MHz, $DMSO-d_6$), δ (ppm): 9.47 (d, $^3J_{HH} = 5.8$ Hz, 1H, 1-CH), 9.14 (d, $^3J_{HH} = 6.2$ Hz, 1H, 10-CH), 8.50 (d, $^3J_{HH} = 8.1$ Hz, 1H, 4-CH), 8.43 (d, $^3J_{HH} = 1.9$ Hz, 1H, 7-CH), 8.38 (t, $^3J_{HH} = 7.8$ Hz, 1H, 3-CH), 7.82 (t, $^3J_{HH} = 6.8$ Hz, 1H, 2-CH), 7.40 (dd, $^3J_{HH} = 6.2$, $^4J_{HH} = 1.9$ Hz, 1H, 9-CH), 7.16 (d, $^3J_{HH} = 8.6$ Hz, 2H, 16,16'-CH), 6.99 (d, $^3J_{HH} = 8.6$ Hz, 2H, 17,17'-CH), 6.85 (d, $^3J_{HH} = 8.6$ Hz, 2H, 21,21'-CH), 6.70 (d, $^3J_{HH} = 8.5$ Hz, 2H, 22,22'-CH), 3.79 (s, 3H, 24-CH₃), 3.63 (s, 3H, 19-CH₃), 2.62 (q, $^3J_{HH} = 7.3$ Hz, 2H, 12-CH₂), 0.95 (t, $^3J_{HH} = 7.4$ Hz, 3H, 13-CH₃). **Elemental analysis:** $C_{28}H_{26}Cl_2N_2O_2Pt$ calc. (%) C 47.85 H 3.81 N 4.07, found (%) C 47.10 H 3.67 N 3.98; **UV-vis:** $\lambda_{max} = 240, 300, 430 \text{ nm}$.

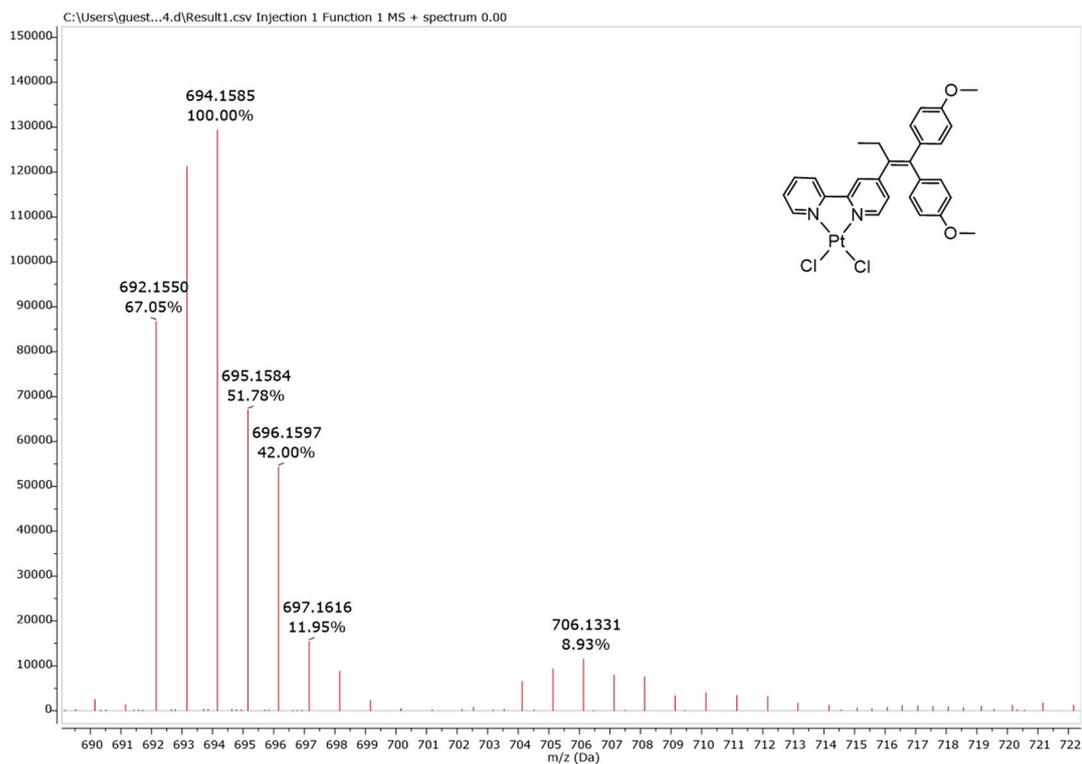


Figure S7. HR-ESI-MS (Positive mode, acetonitrile (MeCN)) of **5**: m/z $[M+NH_4]^+$ = 706.1363 (calc.), 706.1331 (found); m/z $[M-Cl+MeCN]^+$ = 694.1596 (calc.), 694.1585 (found).

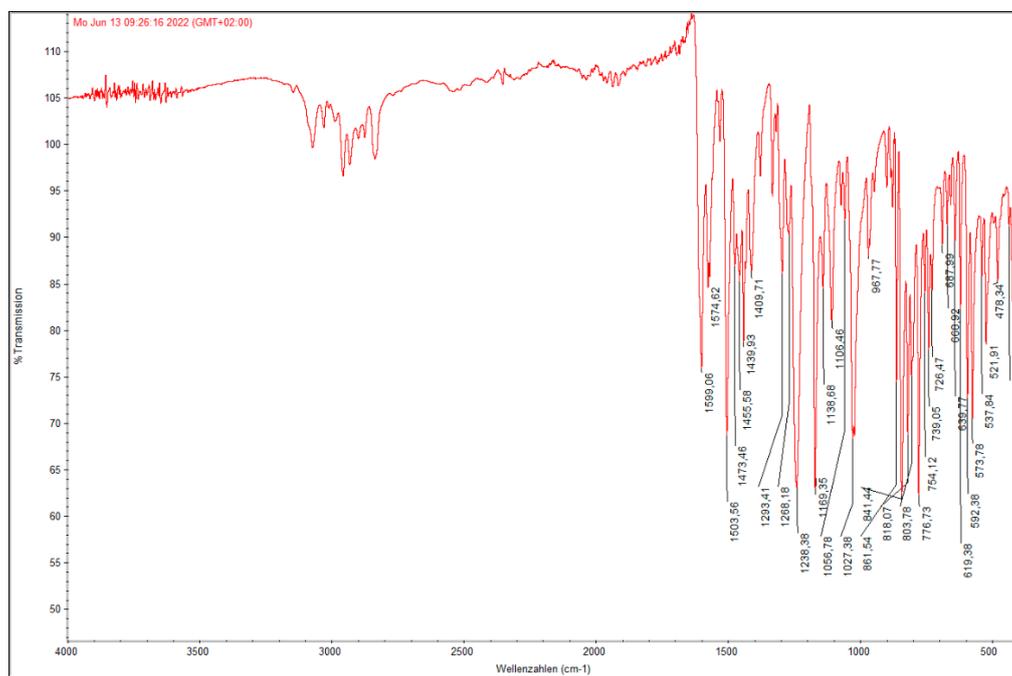


Figure S8. IR spectrum of **5**: 3071–2084 (w, C_{alk}-H), 1599 (m, ν(C=C)), 1575 (m), 1504 (s, C_{arom}-H in-plane bending), 1473 (w, C_{arom}-H in-plane bending), 1456 (w, C_{arom}-H in-plane bending), 1440 (m, C_{arom}-H in-plane bending), 1409 (w), 1293 (w), 1268 (w, C_{arom}-O), 1238 (s, C_{arom}-O), 1169 (s, C_{arom}-O), 1107 (m), 1027 (s, C_{alk}-O), 841 (m, out-of-plane bending), 804 (m, out-of-plane bending) 776 (s, out-of-plane bending), 739 (w), 661 (w), 574 (w), 522 (w), 435 (w).

1.3. [PdCl₂(L-κ²N,N')] (6)

M(C₂₈H₂₆Cl₂N₂O₂Pd) = 598.04 g mol⁻¹, ¹H NMR (400 MHz, CDCl₃), δ (ppm): 9.34 (dd, ³J_{HH} = 5.8, ³J_{HH} = 1.5 Hz, 1H, 1-CH), 9.11 (d, ³J_{HH} = 6.0 Hz, 1H, 10-CH), 8.03 (td, ³J_{HH} = 7.8 Hz, ³J_{HH} = 1.6 Hz, 1H, 3-CH), 7.69 (d, ³J_{HH} = 8.2 Hz, 1H, 4-CH), 7.65 (d, ³J_{HH} = 1.9 Hz, 1H, 7-CH), 7.47 (ddd, ³J_{HH} = 7.4, ³J_{HH} = 5.7, ⁴J_{HH} = 1.3 Hz, 1H, 2-CH), 7.30 (dd, ³J_{HH} = 6.1 Hz, ⁴J_{HH} = 1.9 Hz, 1H, 9-CH), 7.15 (d, ³J_{HH} = 8.6, 2H, 21,21'-CH), 6.91 (d, ³J_{HH} = 8.7 Hz, 2H, 22,22'-CH), 6.84 (d, ³J_{HH} = 8.4 Hz, 2H, 16,16'-CH), 6.66 (d, ³J_{HH} = 8.4 Hz, 2H, 17,17'-CH), 3.84 (s, 3H, 24-CH₃), 3.70 (s, 3H, 19-CH₃), 2.66 (q, ³J_{HH} = 7.4 Hz, 2H, 12-CH₂), 1.02 (t, ³J_{HH} = 7.4 Hz, 3H, 13-CH₃). ¹³C{¹H} NMR (101 MHz, CDCl₃), δ (ppm): 159.4 (23-C), 159.1 (18-C), 156.5 (5-C), 156.3 (6-C), 155.2 (8-C), 151.1 (1-CH), 150.4 (10-CH), 144.9 (14-C), 140.0 (3-CH), 136.0 (20-C), 134.2 (15-C), 133.7 (11-CH), 132.2 (16,16'-CH), 130.5 (21,21'-CH), 127.3 (9-CH), 126.4 (2-CH), 123.9 (7-CH), 121.9 (4-CH), 113.8 (22,22'-CH), 113.8 (17, 17'-CH), 55.3 (24-CH₃), 55.2 (19-CH₃), 27.9 (12-CH₂), 13.8 (13-CH₃). ¹H NMR (400 MHz, DMSO-d₆), δ (ppm): 9.47 (d, ³J_{HH} = 5.8 Hz, 1H, 1-CH), 9.14 (d, ³J_{HH} = 6.2 Hz, 1H, 10-CH), 8.50 (d, ³J_{HH} = 8.1 Hz, 1H, 4-CH), 8.43 (d, ³J_{HH} = 1.9 Hz, 1H, 7-CH), 8.38 (t, ³J_{HH} = 7.8 Hz, 1H, 3-CH), 7.82 (t, ³J_{HH} = 6.8 Hz, 1H, 2-CH), 7.40 (dd, ³J_{HH} = 6.2, ⁴J_{HH} = 1.9 Hz, 1H, 9-CH), 7.15 (d, ³J_{HH} = 8.6 Hz, 2H, 21,21'-CH), 6.99 (d, ³J_{HH} = 8.1 Hz, 2H, 22,22'-CH), 6.85 (d, ³J_{HH} = 8.6 Hz, 2H, 16,16'-CH), 6.70 (d, ³J_{HH} = 8.6 Hz, 1H, 17,17'-CH), 3.79 (s, 3H, 24-CH₃), 3.63 (s, 3H, 19-CH₃), 2.62 (q, ³J_{HH} = 7.5 Hz, 1H, 12-CH₂), 0.95 (t, ³J_{HH} = 7.4 Hz, 3H, 13-CH₃), **Elemental analysis:** C₂₈H₂₆Cl₂N₂O₂Pd calc. (%) C 56.07 H 4.37 N 4.67, found (%) C 55.74 H 4.77 N 4.36; UV-vis: λ_{max} = 245, 290, 400 nm

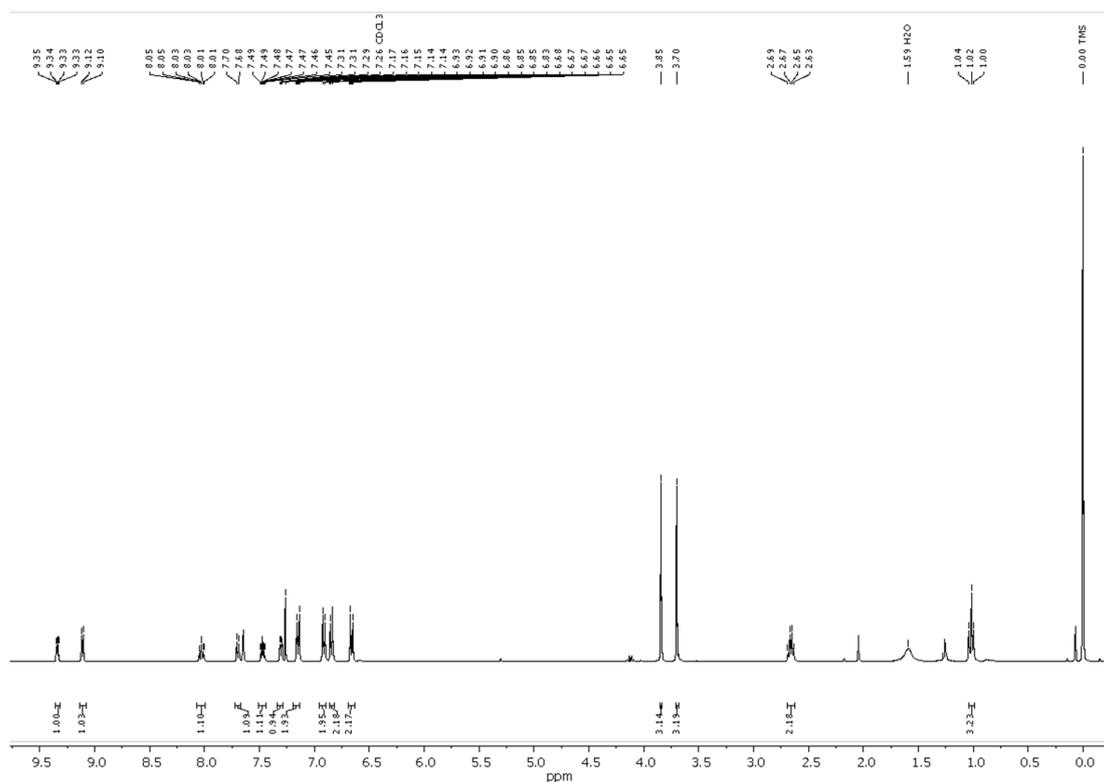


Figure S9. ¹H NMR spectrum of compound 6 in CDCl₃.

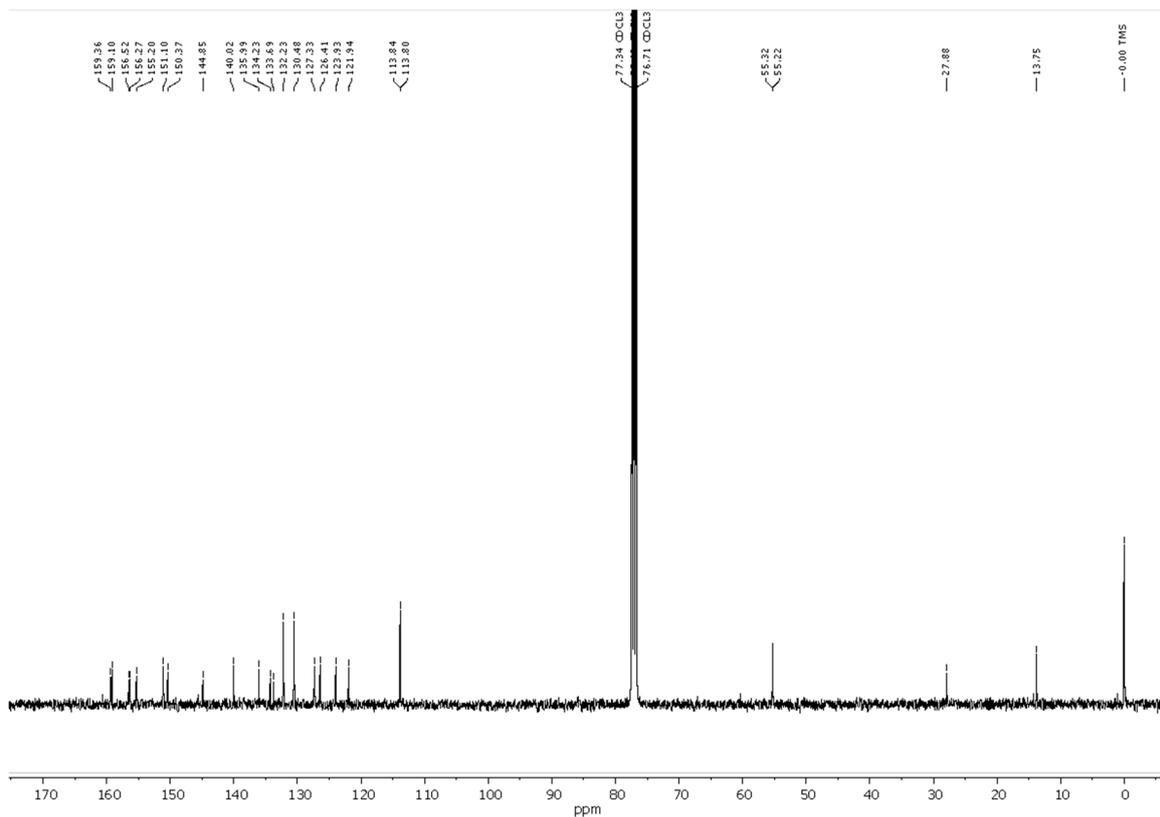


Figure S10. $^{13}\text{C}\{^1\text{H}\}$ NMR spectrum of compound **6** in CDCl_3 .

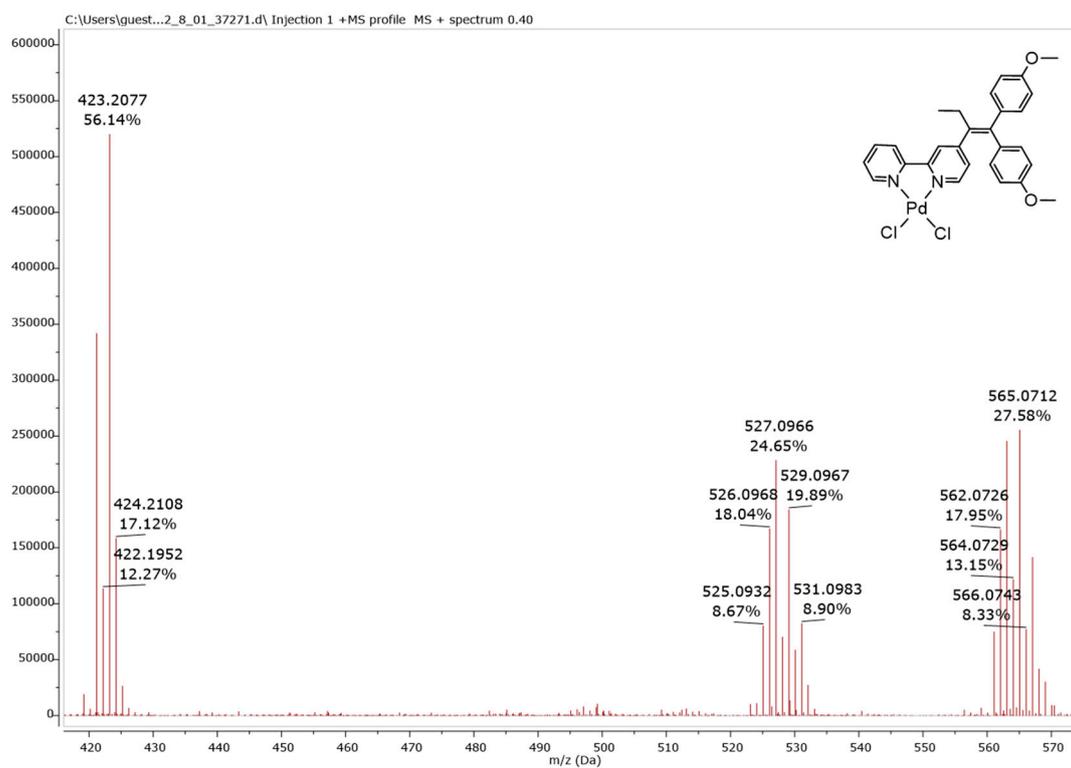


Figure S11. HR-ESI-MS (positive mode, acetonitrile) of **6**: m/z $[\text{M}-\text{Cl}]^+$ = 565.0722 (calc.), 565.0712 (found); m/z $[\text{M}-2\text{Cl}]^+$ = 527.1045 (calc.), 527.0966 (found); m/z $[\text{C}_{28}\text{H}_{26}\text{N}_2\text{O}_2+\text{H}]^+$ = 423.2073 (calc.), 423.2077 (found).

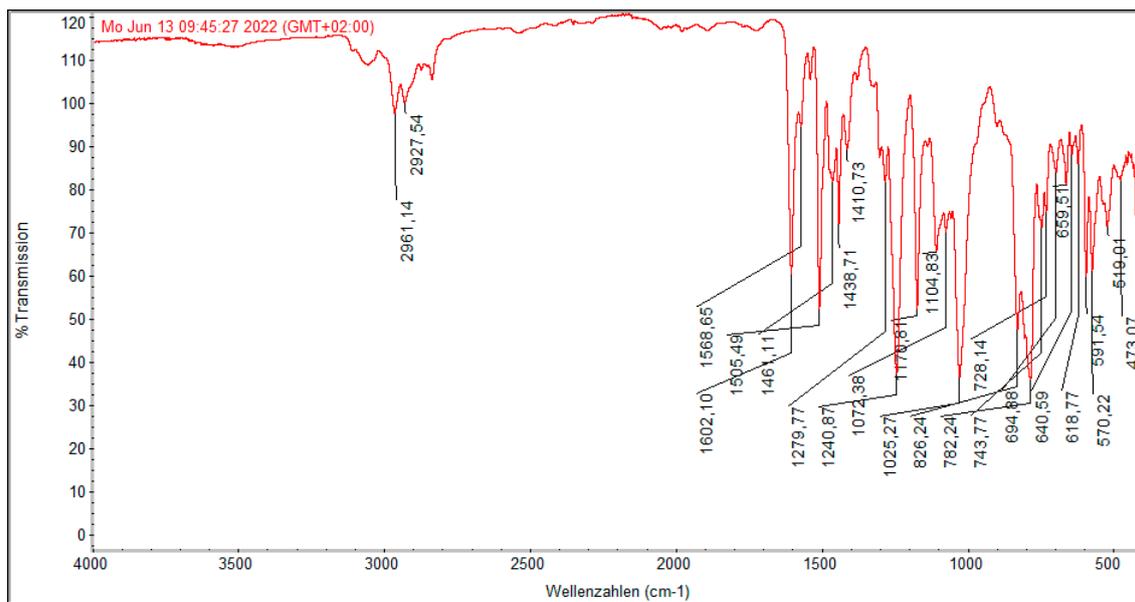


Figure S12. IR spectrum of **6**: 2961 (w, C_{alk}-H), 2927 (w, C_{alk}-H), 1602 (m, v(C=C)), 1569 (w), 1506 (m, C_{arom}-H in-plane bending), 1461 (w, C_{arom}-H in-plane bending), 1439 (m, C_{arom}-H in-plane bending), 1411(w), 1279 (w), 1241 (m, C_{arom}-O), 1171 (m, C_{arom}-O), 1105 (w), 1025 (m, C_{alk}-O), 826 (m, C_{arom}-H out-of-plane bending), 782 (m, C_{arom}-H out-of-plane bending), 744 (w), 660 (w), 619 (w), 591 (w), 570 (w), 519 (w), 473 (w).

1.4. [3-(L-κ²N,N')-3,1,2-PtC₂B₉H₁₁] (7)

R_f = 0.25 (DCM/*n*-hexane, 3:2), M(C₃₀H₃₇N₂O₂PtB₉) = 751.33 g mol⁻¹. ¹H NMR (400 MHz, CD₂Cl₂), δ (ppm): 9.12 (d, ³J_{HH} = 5.7 Hz, 1H, 1-CH), 8.88 (d, ³J_{HH} = 5.9 Hz, 1H, 10-CH), 8.02 (t, ³J_{HH} = 7.8, ³J_{HH} = 1.5 Hz, 1H, 3-CH), 7.88 (dd, ³J_{HH} = 5.1, ³J_{HH} = 3.2 Hz, 2H, 4-CH and 7-CH), 7.45 (t, ³J_{HH} = 6.7 Hz, 1H, 2-CH), 7.27 (dd, ³J_{HH} = 5.9, ⁴J_{HH} = 1.9 Hz, 1H, 9-CH), 7.18 (d, ³J_{HH} = 8.7 Hz, 2H, 21,21'-CH), 6.91 (dd, ³J_{HH} = 12.7, ³J_{HH} = 8.6 Hz, 4H, 22, 22'- and 16,16'-CH), 6.67 (d, ³J = 8.7 Hz, 2H, 17,17'-CH), 3.90 (br s, 2H, CH_{cluster}), 3.83 (s, 3H, 24-CH₃), 3.68 (s, 3H, 19-CH₃), 2.66 (q, ³J_{HH} = 7.4 Hz, 2H, 12-CH₂), 1.02 (t, ³J_{HH} = 7.4 Hz, 3H, 13-CH₃), 2.32–1.03 (m, 9H, BH_{cluster}). ¹³C{¹H} NMR (101 MHz, CD₂Cl₂), δ (ppm): 159.7 (23-C), 159.4 (18-C), 154.9 (5-C), 154.4 (6-C), 153.5 (8-C), 153.3 (1-CH), 152.8 (10-CH), 144.2 (14-C), 138.9 (3-CH), 137.0 (20-C), 135.1 (15-C), 134.5 (11-CH), 132.5 (21,21'-CH), 130.8 (16,16'-CH), 128.0 (9-CH), 127.0 (2-CH), 124.9 (7-CH), 122.8 (4-CH), 114.1 (17,17'-CH and 22,22'-CH), 59.1 (CH_{cluster}), 55.7 (24-CH₃), 55.6 (19-CH₃), 28.2 (12-CH₂), 13.8 (13-CH₂). ¹¹B NMR (128 MHz, CD₂Cl₂), δ (ppm): 15.5 (d, *J* = 147 Hz, 1B), -10.6 (m, 5B), -22.4 (d, *J* = 167 Hz, 1B), -28.6 (d, *J* = 168 Hz, 2B). ¹H NMR (400 MHz, DMSO-*d*₆), δ (ppm): 8.94 (d, ³J_{HH} = 5.4 Hz, 1H, 1-CH), 8.67 (s, 1H, 7-CH), 8.65 (d, ³J_{HH} = 2.9 Hz, 1H, 10-CH), 8.59 (d, ³J_{HH} = 1.9 Hz, 1H, 4-CH), 8.35 (d, ³J_{HH} = 2.9 Hz, 1H, 3-CH), 7.69 (ddd, ³J_{HH} = 7.4, ³J_{HH} = 5.6, ⁴J_{HH} = 1.3 Hz, 1H, 2-CH), 7.30 (dd, ³J_{HH} = 5.9, ⁴J_{HH} = 1.8 Hz, 1H, 9-CH), 7.16 (d, ³J_{HH} = 8.7 Hz, 2H, 21,21'-CH), 6.99 (d, *J* = 8.7 Hz, 2H, 22,22'-CH), 6.88 (d, *J* = 8.7 Hz, 2H, 16,16'-CH), 6.71 (d, *J* = 8.8 Hz, 2H, 17,17'-CH), 3.99 (br s, 2H, CH_{cluster}), 3.79 (s, 3H, 24-CH₃), 3.63 (s, 3H, 19-CH₃), 2.63 (q, ³J_{HH} = 7.5 Hz, 2H, 12-CH₂), 0.95 (t, ³J_{HH} = 7.4 Hz, 3H, 13-CH₃), 0.40–2.5 (m, 9-BH). **Elemental analysis:** C₃₀H₃₇N₂O₂PtB₉ calc. (%) C 48.04 H 4.97 N 3.74, found (%) C 47.68 H 5.26 N 3.93; **UV-vis:** λ_{max} = 240, 300, 450 nm.

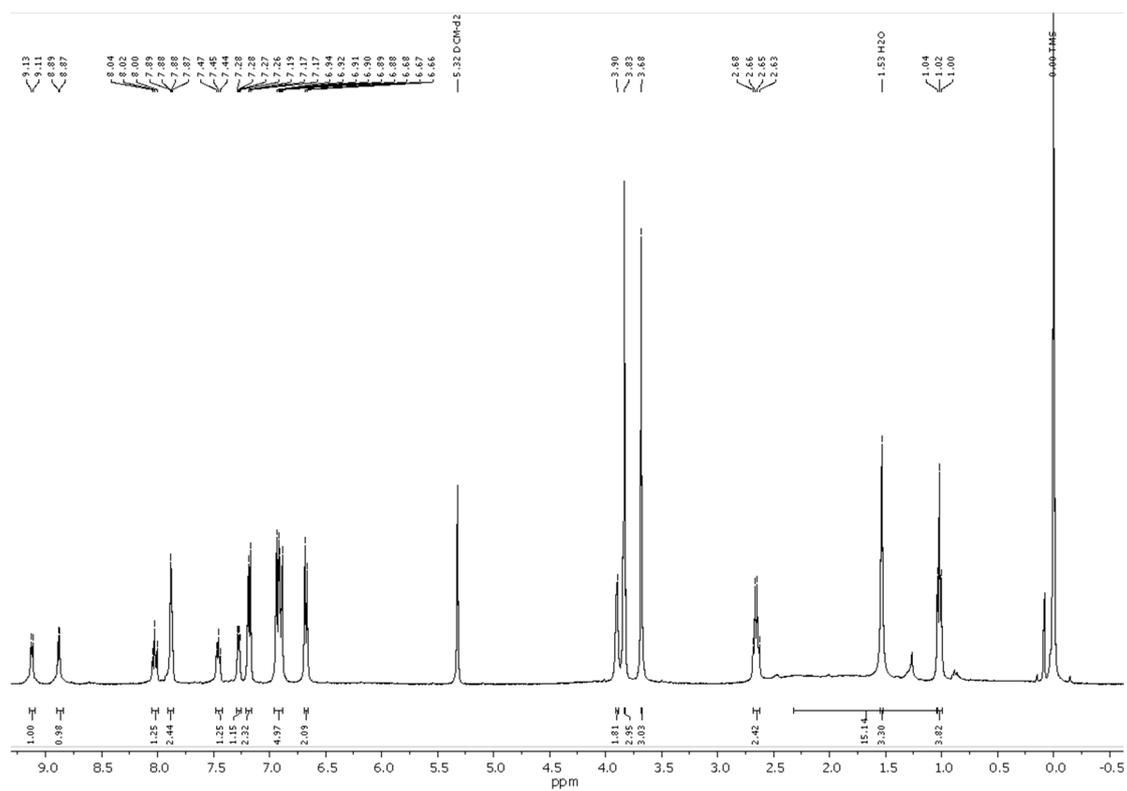


Figure S13. ¹H NMR spectrum of compound 7 in CD₂Cl₂.

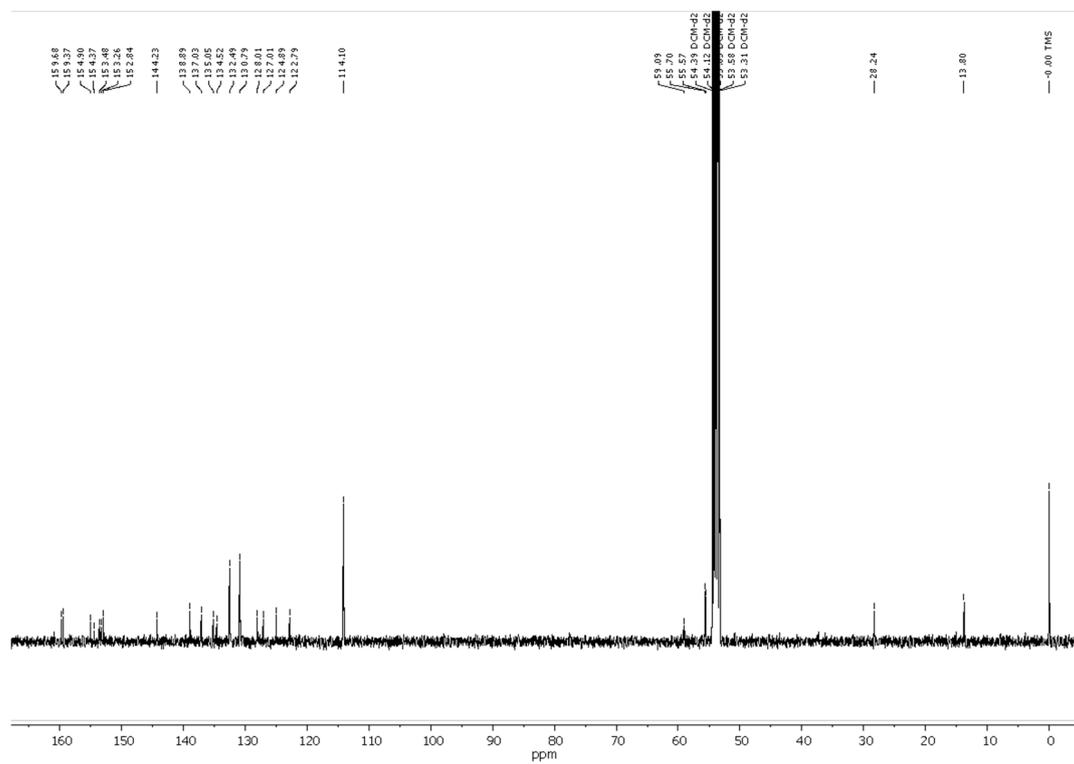


Figure S14. ¹³C{¹H} NMR spectrum of compound 7 in CD₂Cl₂.

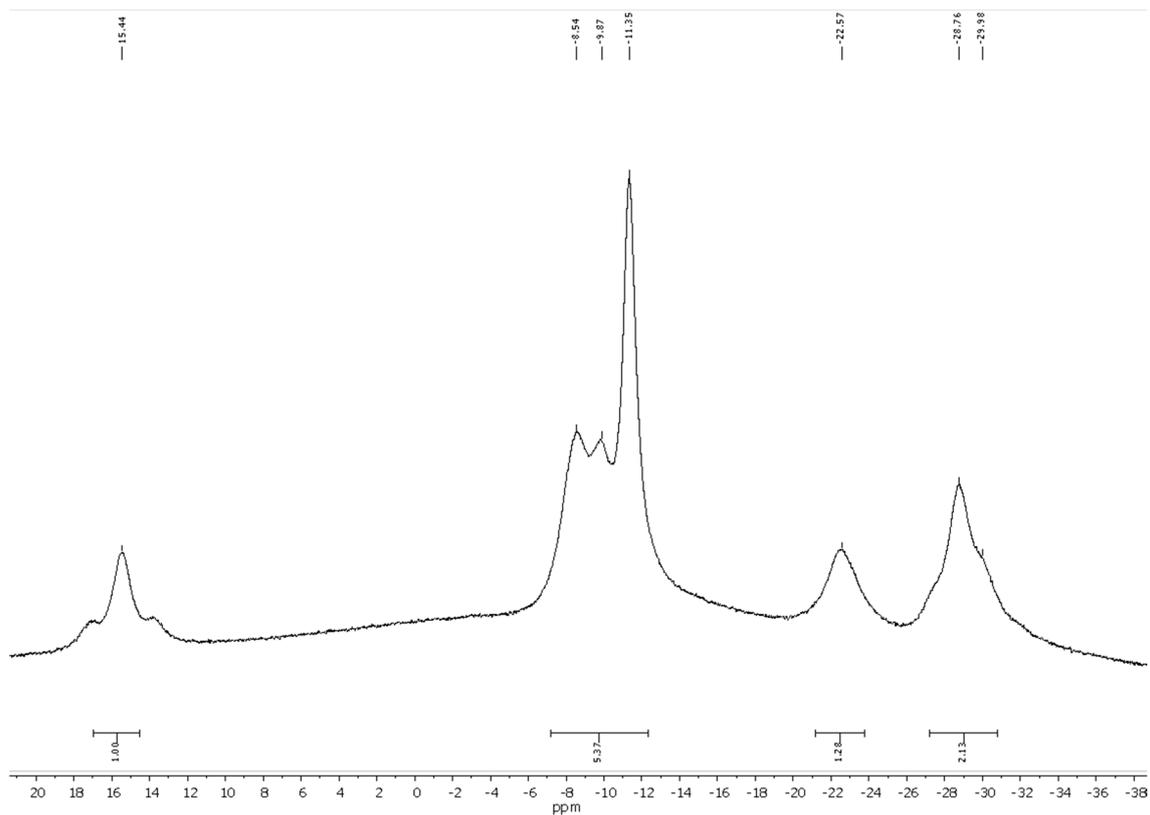


Figure S15. $^{11}\text{B}\{^1\text{H}\}$ NMR spectrum of compound **7** in CD_2Cl_2 .

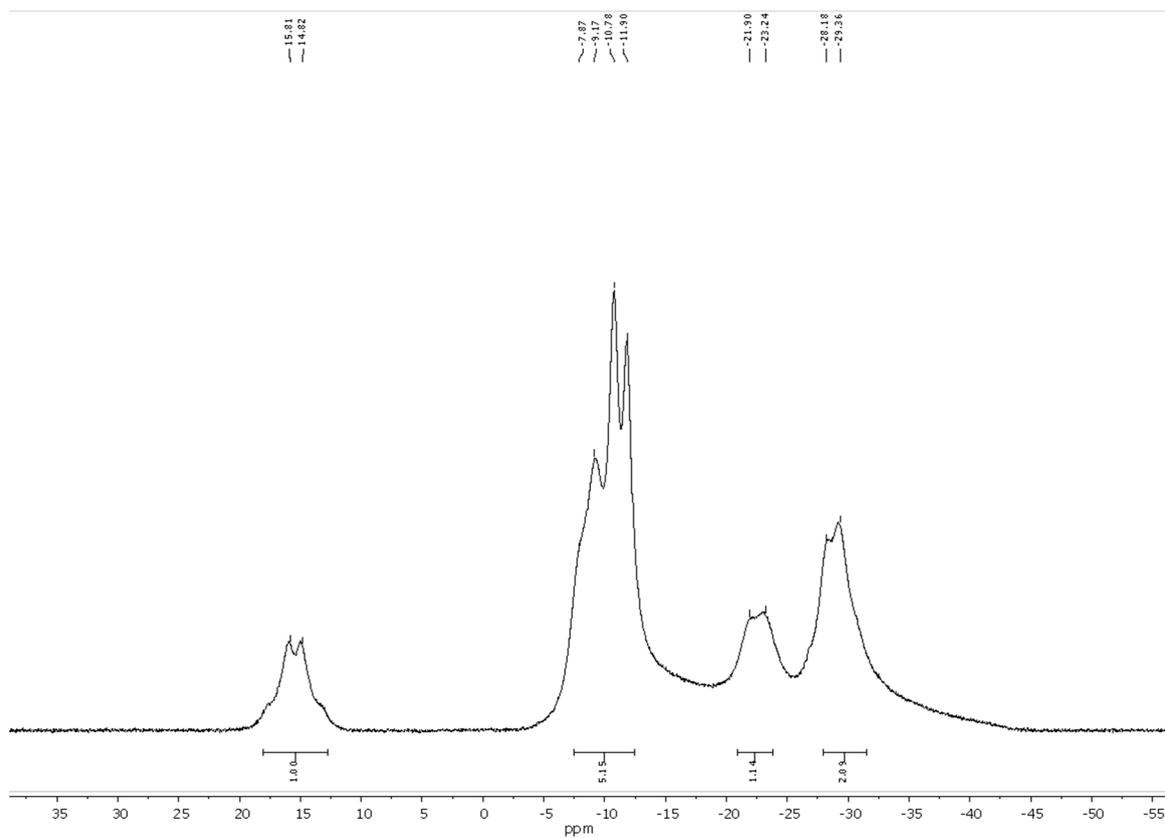


Figure S16. ^{11}B NMR spectrum of compound **7** in CD_2Cl_2 .

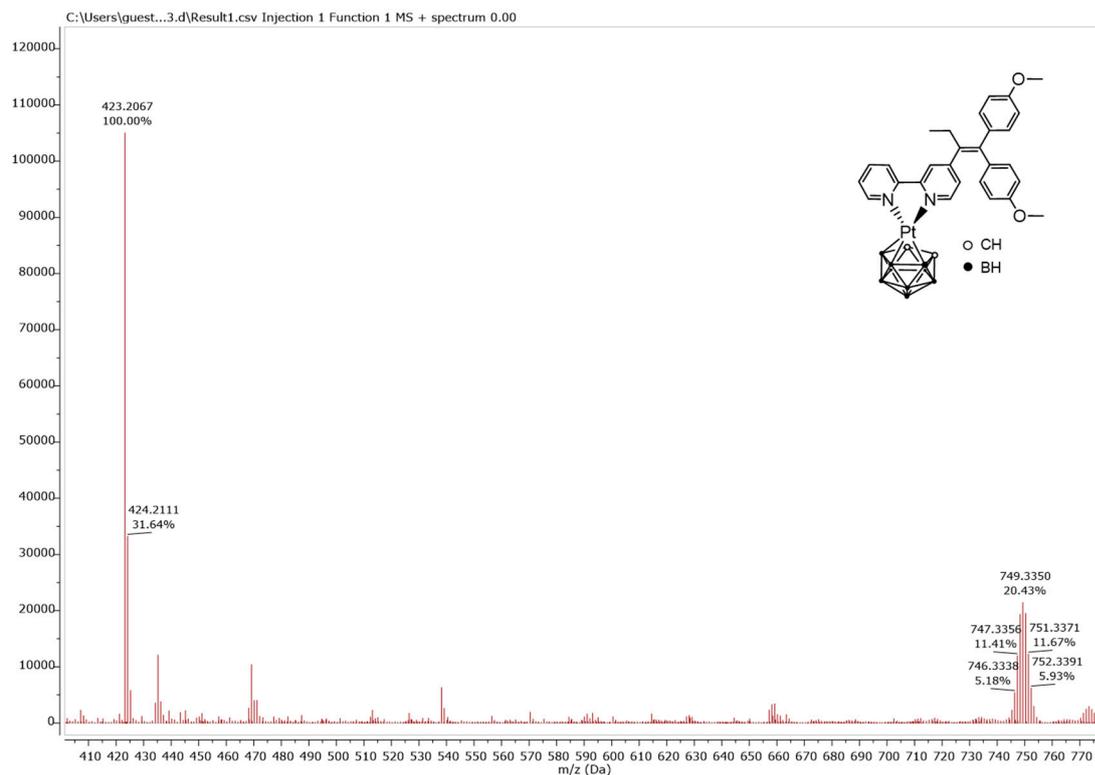


Figure S17. HR-ESI-MS (positive mode, acetonitrile) of **7**: m/z $[M+H]^+ = 749.3340$ (calc.), 749.3350 (found); m/z $[C_{28}H_{26}N_2O_2+H]^+ = 423.2073$ (calc.), 423.2067 (found).

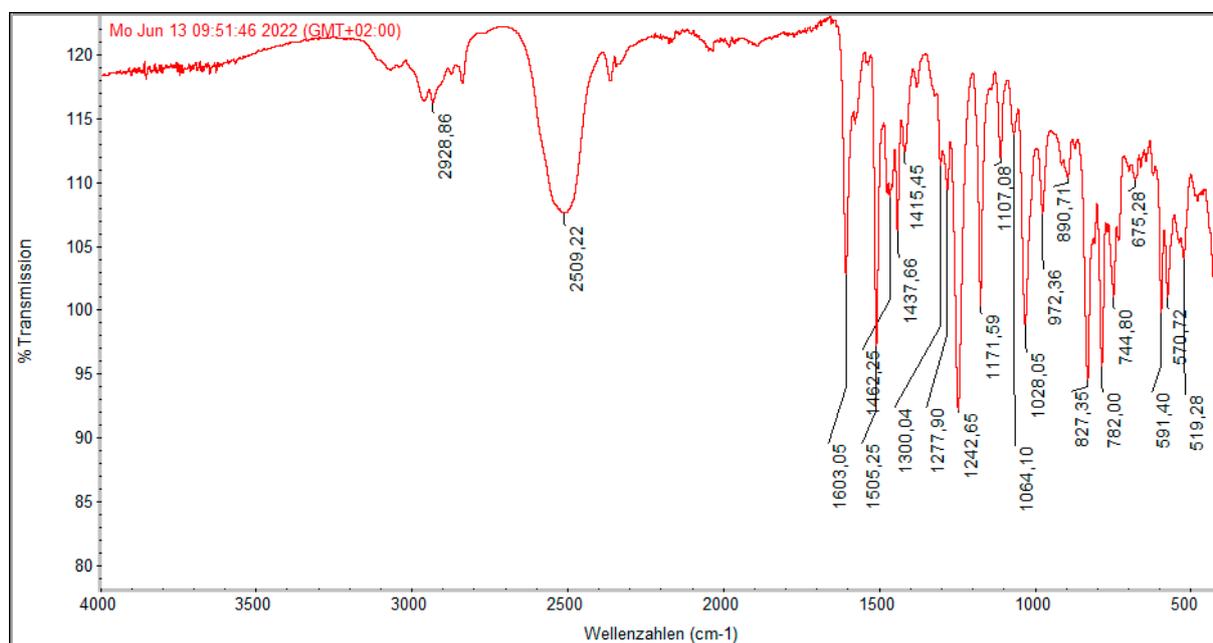


Figure S18. IR spectra of **7**: 2961(w, C_{alk-H}), 2929(w, C_{alk-H}), 2509(m, $\nu(B-H)$), 1603(m, $\nu(C=C)$), 1505(m, C_{arom-H} in-plane bending), 1462(w, C_{arom-H} in-plane bending), 1438(m, C_{arom-H} in-plane bending), 1277(w), 1243(s, C_{arom-O}), 1173(m, C_{arom-O}), 1028(m, C_{alk-O}), 826(m, C_{arom-H} out-of-plane bending), 782(m, C_{arom-H} out-of-plane bending), 591(w), 520(w).

1.5. [3-(*L*-κ²N,N')-3,1,2-PdC₂B₉H₁₁] (8)

$R_f = 0.40$ (DCM/*n*-hexane 3:1), $M(C_{30}H_{37}N_2O_2PdB_9) = 662.27$ g mol⁻¹. ¹H NMR (400 MHz, CDCl₃), δ (ppm): 8.66 (d, ³J_{HH} = 5.6, 1H, 1-CH), 8.45 (d, ³J_{HH} = 5.7 Hz, 1H, 10-CH), 7.94 (t, ³J_{HH} = 7.8 Hz, 1H, 3-CH), 7.83 (m, 2H, 4-CH and 7-CH), 7.41 (t, ³J_{HH} = 6.5 Hz, 1H, 2-CH), 7.23 (d, ³J_{HH} = 6.0, 1H, 9-CH), 7.16 (d, ³J = 8.6 Hz, 2H, 21,21'-CH), 6.92 (d, ³J = 8.7 Hz, 2H, 22,22'-CH), 6.86 (d, ³J = 8.7 Hz, 2H, 16,16'-CH), 6.67 (d, ³J = 8.7 Hz, 2H, 17,17'-CH), 3.84 (s, 3H, 24-CH₃), 3.76 (br s, 2H, CH_{cluster}), 3.70 (s, 3H, 19-CH₃), 2.65 (q, ³J = 7.4 Hz, 2H, 12-CH₂), 1.00 (t, ³J = 7.4 Hz, 3H, 13-CH₃), 0.80–2.35 (m, 9H, BH_{cluster}). ¹³C{¹H} NMR (101 MHz, CDCl₃), δ (ppm): 159.2 (23-C), 158.9 (18-C), 154.4 (5-C), 153.5 (6-C), 152.4 (8-C), 152.4 (1-CH), 151.9 (10-CH), 143.8 (14-C), 138.3 (3-CH), 136.4 (20-C), 134.5 (15-C), 134.0 (11-CH), 132.1 (21, 21'-CH), 130.4 (16,16'-CH), 126.8 (9-C), 125.8 (2-CH), 124.0 (7-CH), 121.9 (4-CH), 113.8 (17,17'-CH), 113.8 (22,22'-CH), 55.3 (24-CH), 55.2 (19-CH), 49.5 (CH_{cluster}), 28.0 (12-CH), 13.7 (13-CH). ¹¹B NMR (400 MHz, CDCl₃), δ (ppm): 17.9 (d, $J = 134$ Hz, 1B, BH), -9.7 (d, $J = 142$ Hz, 5B, BH), -20.7 (m, 3B, BH). ¹H NMR (400 MHz, DMSO-d₆), δ (ppm): 8.59 (d, ³J_{HH} = 8.2 Hz, 1H, 10-CH), 8.53 (d, ³J_{HH} = 1.8 Hz, 1H, 7-CH), 8.50 (dd, ³J_{HH} = 5.5, ³J_{HH} = 1.5 Hz, 1H, 1-CH), 8.24 (m, 2H, 4-CH and 3-CH), 7.66 (ddd, ³J_{HH} = 7.3, ³J_{HH} = 5.5, ⁴J_{HH} = 1.2 Hz, 1H, 2-CH), 7.28 (dd, ³J_{HH} = 5.8, ⁴J_{HH} = 1.8 Hz, 1H, 9-CH), 7.16 (d, ³J_{HH} = 8.7 Hz, 2H, 21,21'-CH), 6.98 (d, ³J_{HH} = 8.7 Hz, 2H, 22,22'-CH), 6.87 (d, ³J_{HH} = 8.8 Hz, 16,16'-CH), 6.70 (d, ³J_{HH} = 8.7 Hz, 2H, 17,17'-CH), 3.85 (br s, 2H, CH_{cluster}), 3.78 (s, 3H, 24-CH₃), 3.62 (s, 3H, 19-CH₃), 2.61 (q, ³J_{HH} = 7.3 Hz, 2H, 12-CH₂), 0.93 (t, ³J_{HH} = 7.4 Hz, 3H, 13-CH₃), 0.30–2.24 (m, 9H, BH_{cluster}). **Elemental analysis:** C₃₀H₃₇N₂O₂PdB₉ Calc. (%) C 54.49 H 5.64 N 4.24 Found (%) C 54.15 H 5.60 N 3.84; **UV-vis:** $\lambda_{max} = 240, 300, 420$ nm.

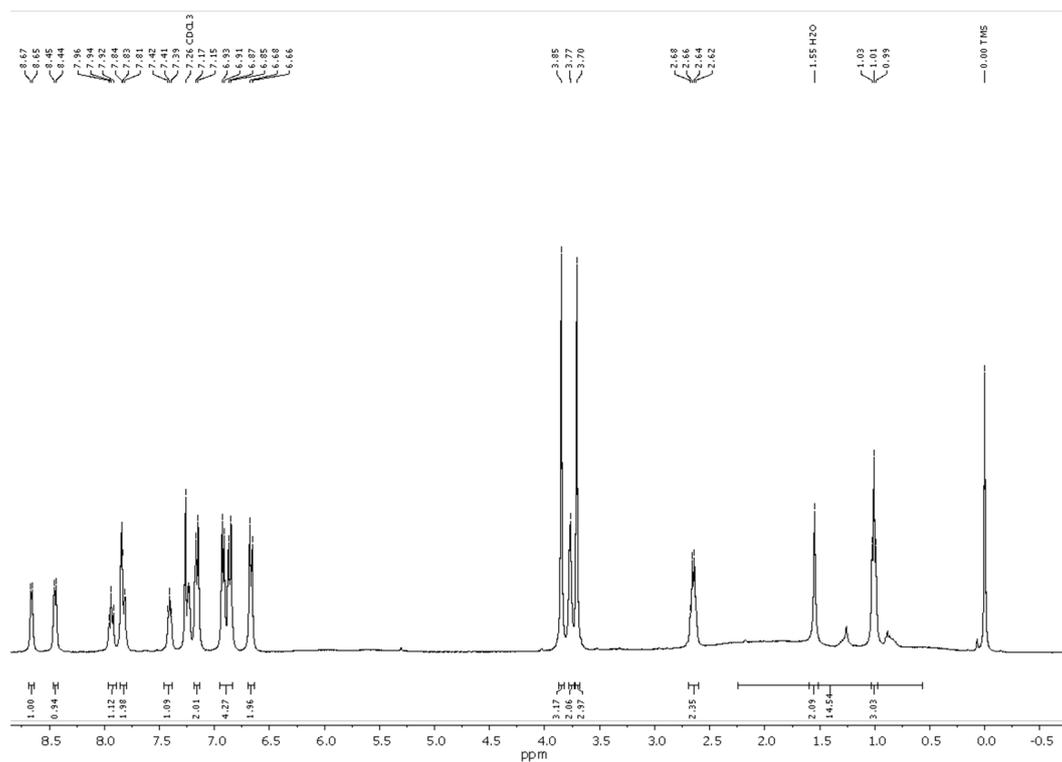


Figure S19. ¹H NMR spectrum of compound 8 in CDCl₃.

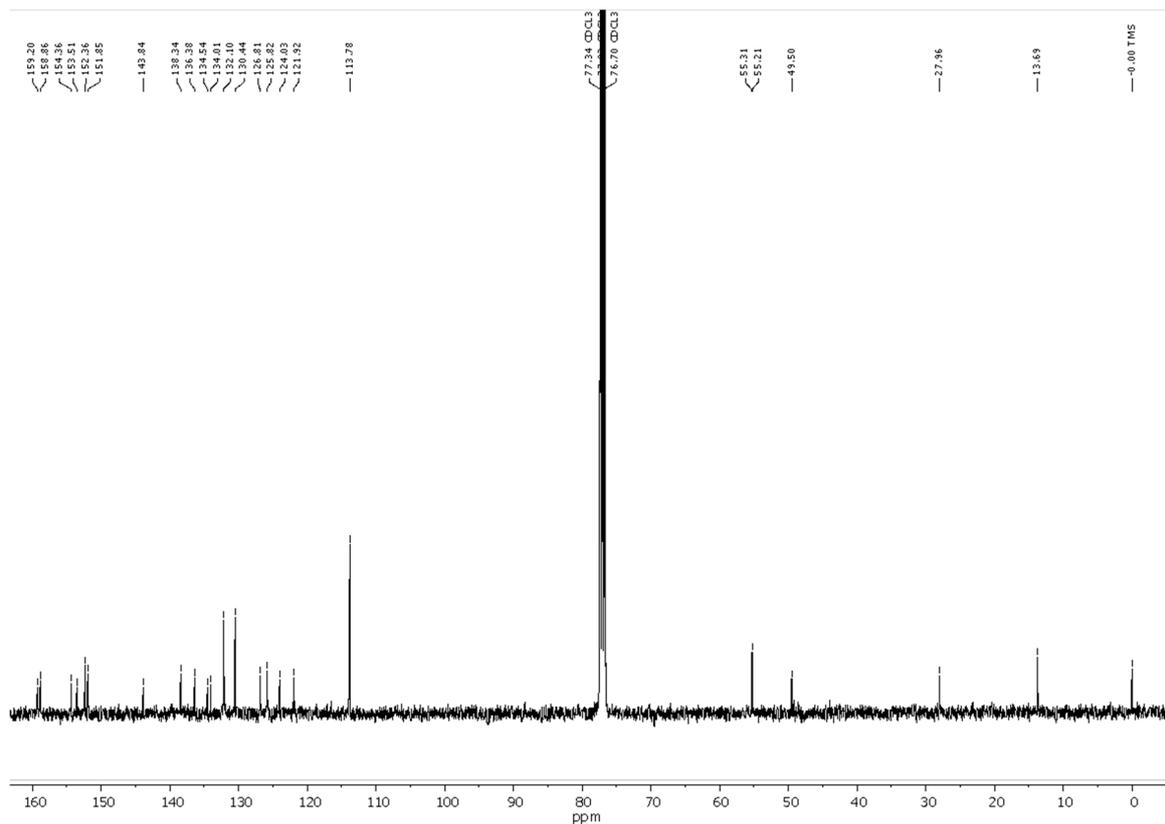


Figure S20. $^{13}\text{C}\{^1\text{H}\}$ NMR spectrum of compound **8** in CDCl_3 .

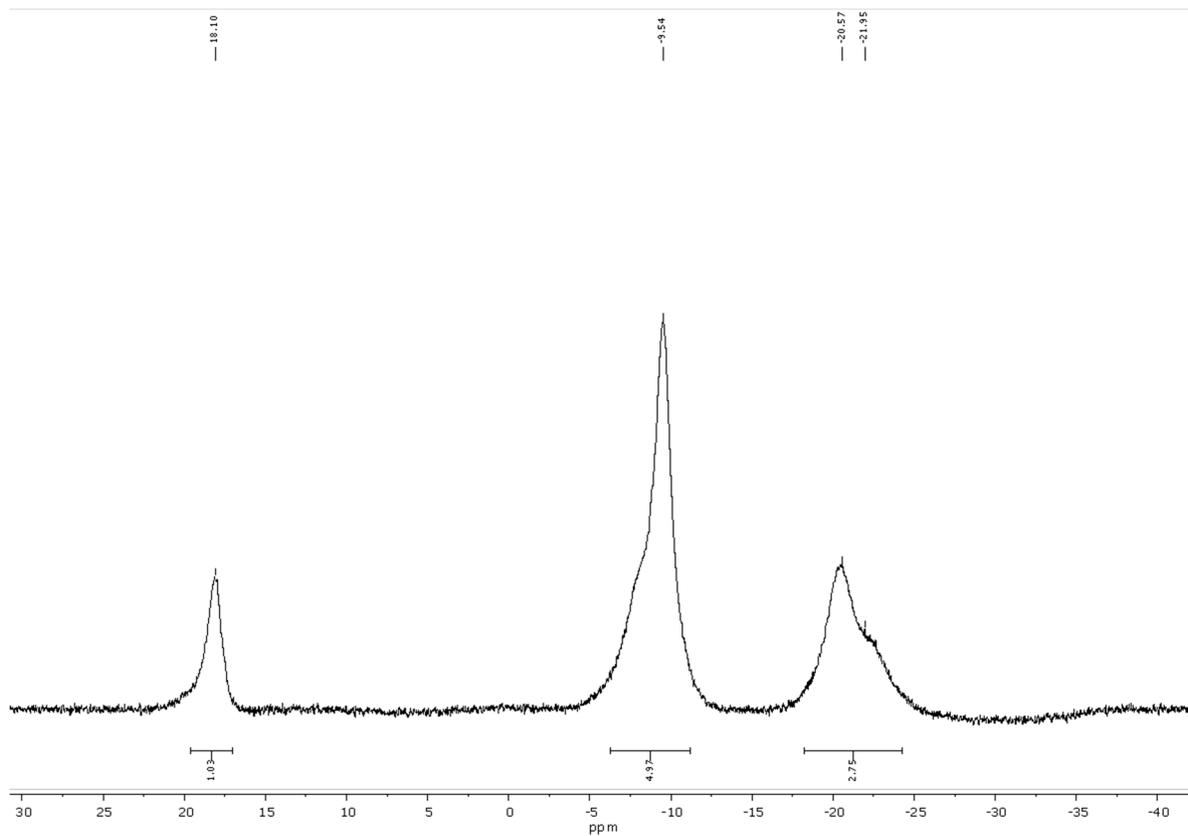


Figure S21. $^{11}\text{B}\{^1\text{H}\}$ NMR spectrum of compound **8** in CDCl_3 .

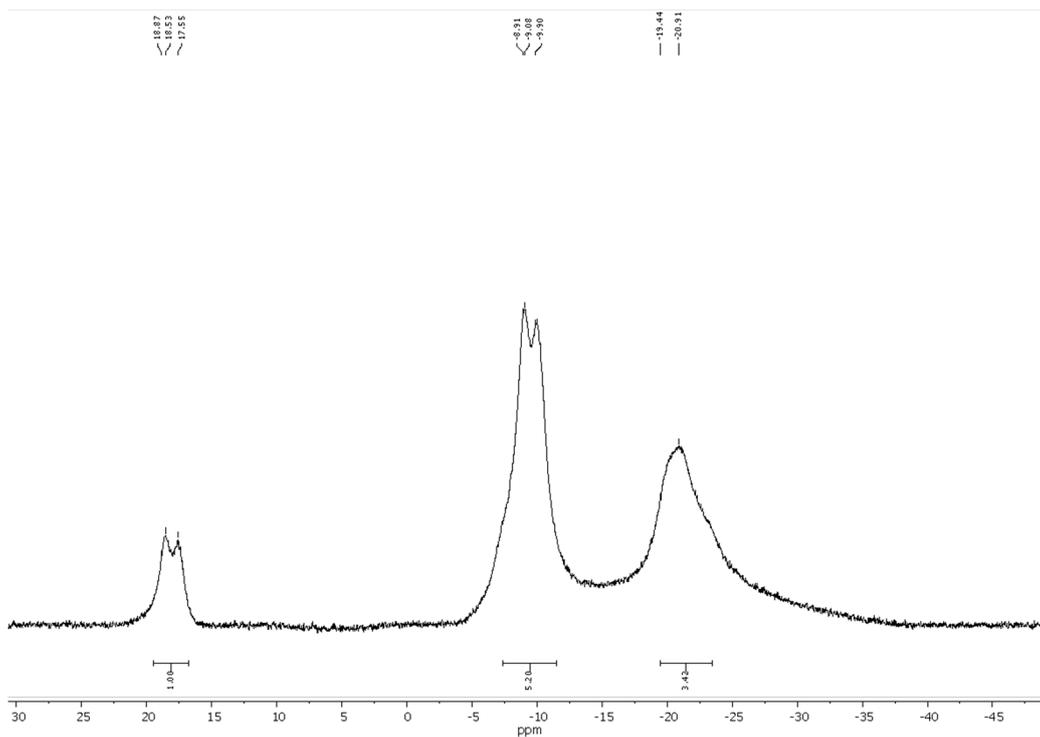


Figure S22. ^{11}B NMR spectrum of compound **8** in CDCl_3 .

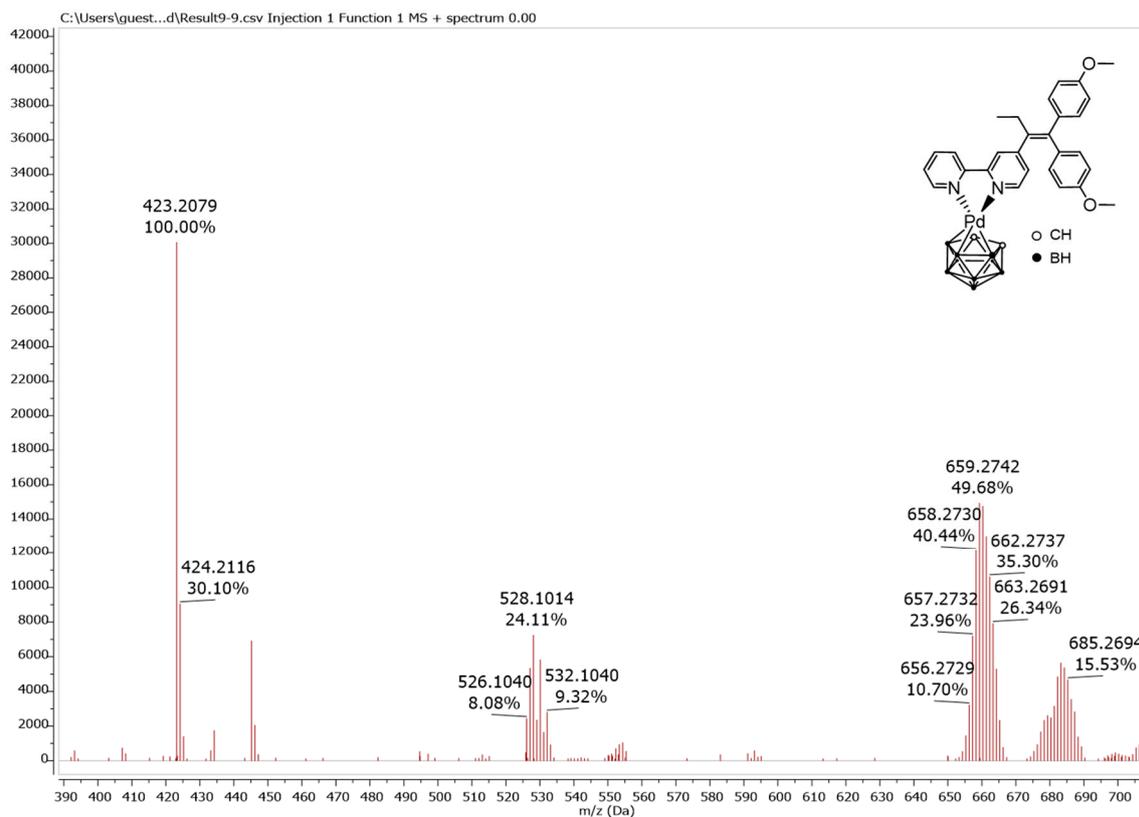


Figure S23. HR-ESI-MS (positive mode, acetonitrile) of **8**: m/z $[\text{M}+\text{Na}]^+$ = 685.2625 (calc.), 685.2694 (found); m/z $[\text{M}+\text{H}]^+$ = 659.2826 (calc.), 659.2742 (found); m/z $[\text{M}-\text{C}_2\text{B}_9\text{H}_{11}]^+$ = 528.1029 (calc.), 528.1014 (found), m/z $[\text{C}_{28}\text{H}_{26}\text{N}_2\text{O}_2+\text{H}]^+$ = 423.2073 (calc.), 423.2079 (found).

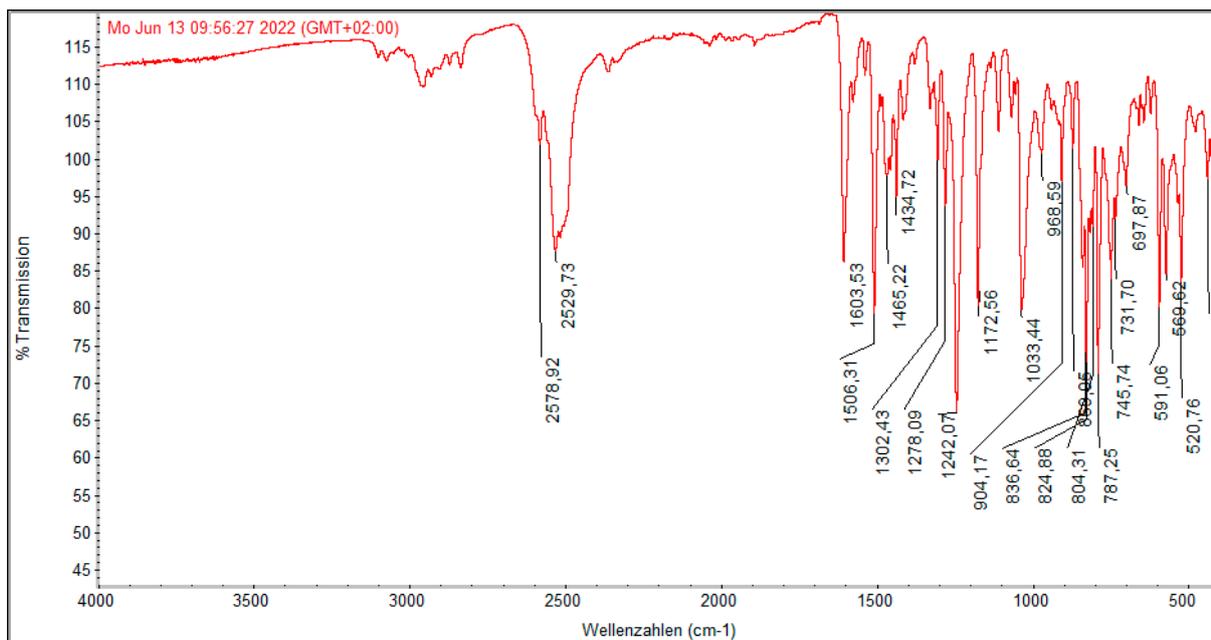


Figure S24. IR: 2961(w, C_{alk}-H), 2927 (w, C_{alk}-H), 2580-2500 (m, v (B-H)), 1603 (m, v(C=C)), 1506 (m, C_{arom}-H in-plane bending), 1465 (w, C_{arom}-H in-plane bending), 1278 (w), 1242 (s, C_{arom}-O), 1173 (m, C_{arom}-O) 1033 (m, C_{alk}-O), 850 (m, C_{arom}-H out-of-plane bending), 746 (m, C_{arom}-H out-of-plane bending) 591 (m), 521 (w).

2. X-ray crystallography

The data were collected on a Gemini diffractometer (Rigaku Oxford Diffraction) using Mo-K α radiation and ω -scan rotation. Data reduction was performed with CrysAlisPro [1] including the program SCALE3 ABSPACK for empirical absorption correction. All structures were solved by dual space methods with SHELXT [2] and the refinement was performed with SHELXL [3]. Non-hydrogen atoms were refined with anisotropic displacement parameters. Hydrogen atoms for **5**, **6** and **7** were calculated on idealised positions using the riding model, whereas for **8** a difference-density Fourier map was used to locate all hydrogen atoms. Structure figures were generated with Diamond [4].

The structures of **5** and **6**, as well as of **7** and **8**, are isotypic. For **7** and **8**, the carborane carbon atoms could be localised with a bond length and displacement parameter analysis.

CCDC deposition numbers given in the Tables below contain the supplementary crystallographic data for this paper. These data can be obtained free of charge via <https://summary.ccdc.cam.ac.uk/structure-summary-form> (or from the Cambridge Crystallographic Data Centre, 12 Union Road, Cambridge CB2 1EZ, UK; fax: (+44)1223-336-033; or deposit@ccdc.cam.ac.uk).

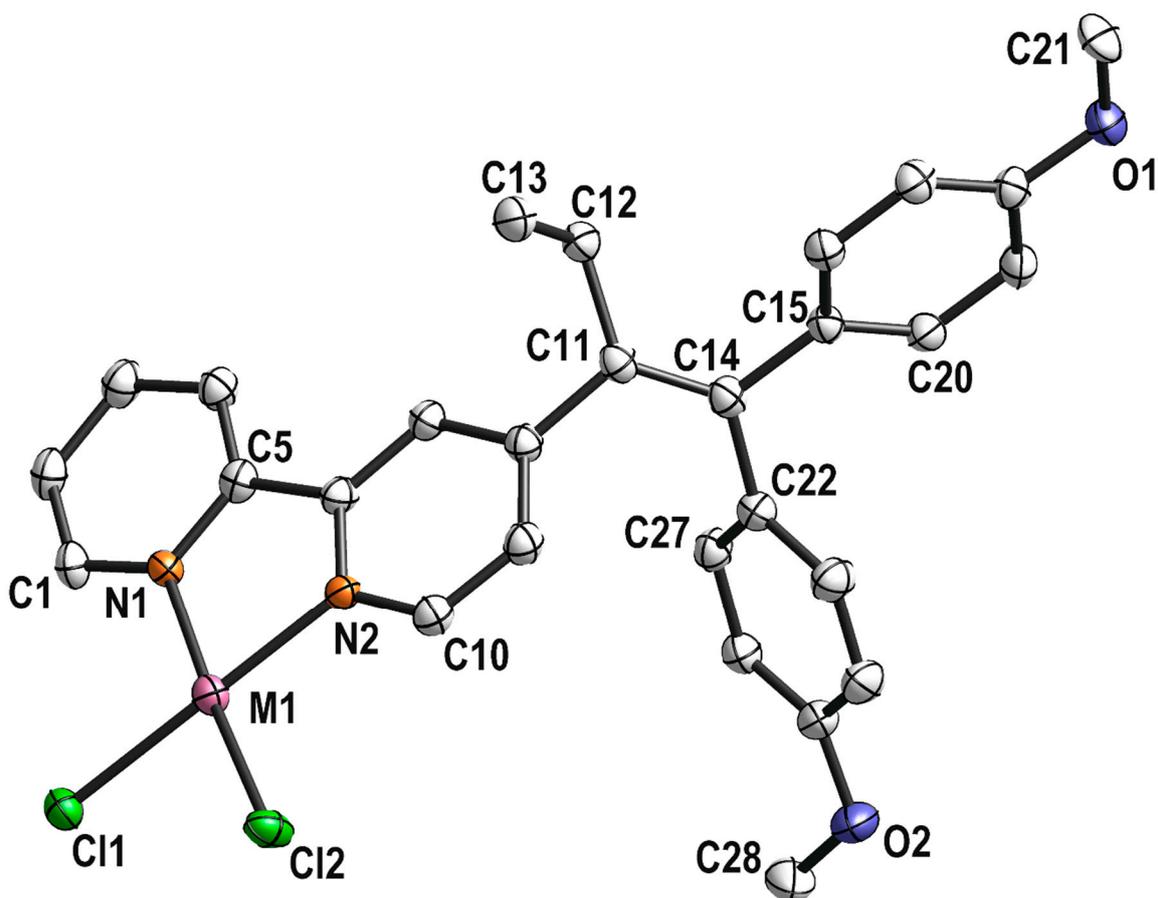


Figure S25. Molecular structures of **5** (M1 = Pt) and **6** (M1 = Pd) complexes in the solid state. Hydrogen atoms are omitted for clarity. Displacement ellipsoids are presented at 50% probability level. The calculated bond lengths and angles at PBE0 D3BJ/def2-TZVPP level of theory are in agreement with the crystallographic data. N1–Pt (2.010 calc., 2.019 exp.), N2–Pt (2.015 calc., 2.021 exp.), Cl1–Pt (2.296 calc., 2.305 exp.), Cl2–Pt (2.286 calc., 2.291 exp.); \angle N1–Pt–Cl1 (93.9° calc, 94.3° exp); \angle N2–Pt–Cl2 (95.1° calc, 94.8° exp); \angle Cl1–Pt–Cl2 (90.4° calc, 90.0° exp); \angle N1–Pt–N2 (80.6° calc, 81.1° exp). N1–Pd (2.010 calc., 2.019 exp.), N2–Pd (2.015 calc., 2.021 exp.), Cl1–Pd (2.296 calc., 2.305 exp.), Cl2–Pd (2.286 calc., 2.291 exp.); \angle N1–Pd–Cl1 (94.4° calc, 94.1° exp); \angle N2–Pd–Cl2 (94.2° calc, 94.3° exp); \angle Cl1–Pd–Cl2 (91.7° calc, 91.1° exp); \angle N1–Pd–N2 (79.7° calc, 80.6° exp).

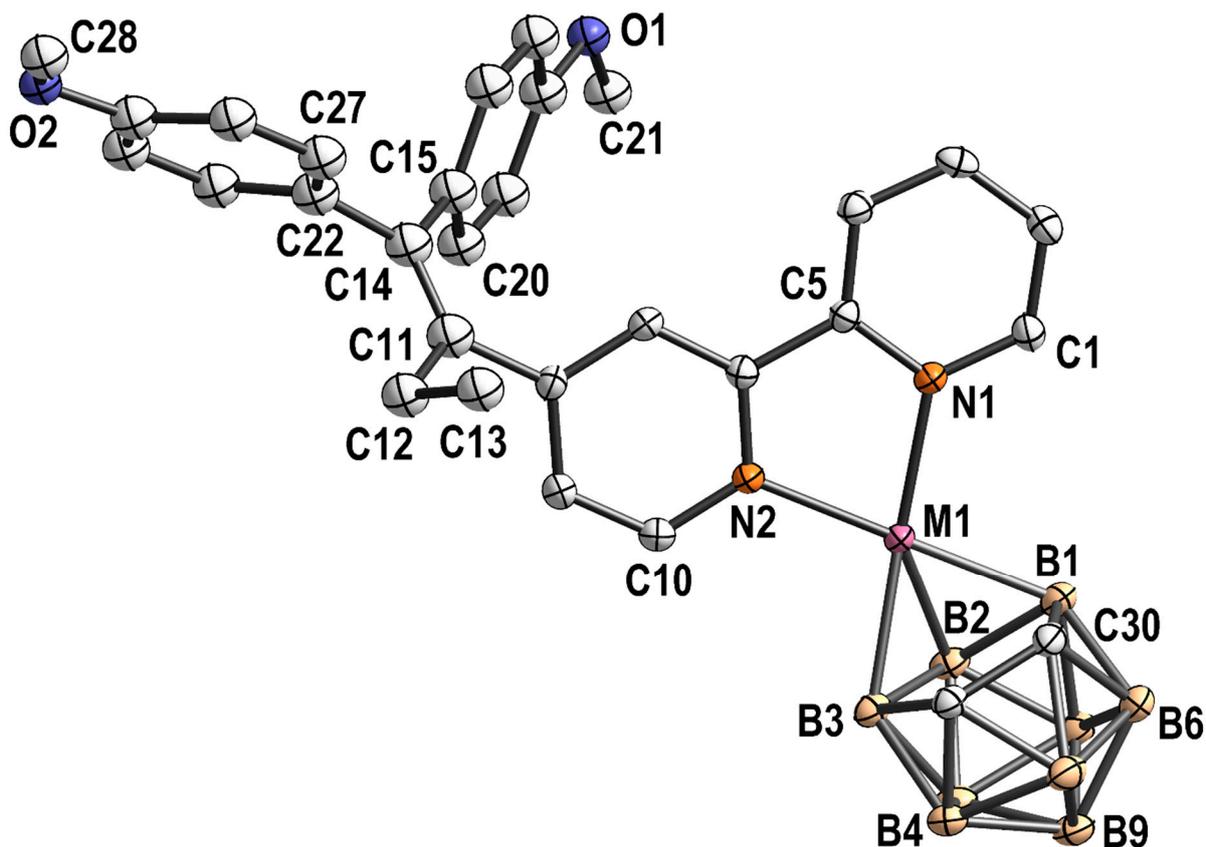


Figure S26. Molecular structure of **7** (M1 = Pt) and **8** (M1 = Pd) in the solid state. Hydrogen atoms are omitted for clarity. Displacement ellipsoids are presented at 50% probability level. The calculated bond lengths and angles at PBE0 D3BJ/def2-TZVPP level of theory are in agreement with the crystallographic data with the accuracy. N1–Pt (2.081 calc., 2.082 exp.), N2–Pt (2.074 calc., 2.072 exp.), B1–Pt (2.189 calc., 2.191 exp.), B2–Pt (2.172 calc., 2.183 exp.), B3–Pt (2.183 calc., 2.179 exp.); \angle B1–Pt–B3 (77.5° calc, 77.9° exp); \angle N1–Pt–N2 (77.1° calc, 77.1° exp); \angle N1–Pt–B1 (102.8° calc, 102.5° exp); \angle N2–Pt–B3 (102.6° calc, 102.5° exp), \angle B1–Pt–B2 (49.0° calc, 48.9° exp), \angle B2–Pt–B3 (49.2° calc, 49.0° exp). N1–Pd (2.113 calc., 2.110 exp.), N2–Pd (2.105 calc., 2.098 exp.), B1–Pd (2.172 calc., 2.178 exp.), B2–Pd (2.238 calc., 2.270 exp.), B3–Pd (2.167 calc., 2.190 exp.); \angle B1–Pd–B3 (76.7° calc, 77.2° exp); \angle N1–Pd–N2 (77.5° calc, 77.8° exp); \angle N1–Pd–B1 (103.0° calc, 103.0° exp); \angle N2–Pd–B3 (102.8° calc, 102.9° exp), \angle B1–Pd–B2 (47.6° calc, 46.9° exp), \angle B2–Pd–B3 (47.8° calc, 47.4° exp).

Table S1. Summary of the structure refinement data of complexes **5** and **6**.

Empirical formula	C ₂₈ H ₂₆ Cl ₂ N ₂ O ₂ Pt (5)	C ₂₈ H ₂₆ Cl ₂ N ₂ O ₂ Pd (6)
Formula weight	688.50	599.81
Temperature	130(2) K	130(2) K
Wavelength	71.073 pm	71.073 pm
Crystal system	Monoclinic	Monoclinic
Space group	<i>P2₁/c</i>	<i>P2₁/c</i>
Unit cell dimensions	a = 1102.57(2) pm	1105.69(3) pm
	b = 1142.19(2) pm	1140.40(4) pm
	c = 2005.24(4) pm	1996.46(8) pm
	α = 90°	90°
	β = 102.364(2)°	102.343(3)°
	γ = 90°	90°
Volume	2.46672(8) nm ³	2.4592(2) nm ³
Z	4	4
Density (calculated)	1.854 mg m ⁻³	1.620 mg m ⁻³
Absorption coefficient	5.935 mm ⁻¹	1.002 mm ⁻¹
<i>F</i> (000)	1344	1216
Crystal size	0.40 x 0.25 x 0.10 mm ³	0.15 x 0.03 x 0.02 mm ³
Theta range for data collection	1.891 to 32.515°	1.885 to 28.193°
Index ranges	-16 ≤ <i>h</i> ≤ 15, -17 ≤ <i>k</i> ≤ 16, -29 ≤ <i>l</i> ≤ 29	-13 ≤ <i>h</i> ≤ 10, -13 ≤ <i>k</i> ≤ 14, -24 ≤ <i>l</i> ≤ 24
Reflections collected	32170	17624
Independent reflections	8303 [R(int) = 0.0365]	5257 [R(int) = 0.0806]
Completeness (theta)	100.0 % (30.51°)	100.0 % (25.35°)
Absorption correction	Semi-empirical from equivalents	Semi-empirical from equivalents
Max. and min. transmission	1.00000 and 0.49996	1.00000 and 0.86051
Refinement method	Full-matrix least-squares on <i>F</i> ²	Full-matrix least-squares on <i>F</i> ²
Data / restraints / parameters	8303 / 0 / 319	5257 / 0 / 319
Goodness-of-fit on <i>F</i> ²	1.190	1.017
Final R indices [I > 2σ(I)]	R1 = 0.0391, wR2 = 0.0712	R1 = 0.0495, wR2 = 0.0862
R indices (all data)	R1 = 0.0496, wR2 = 0.0753	R1 = 0.0998, wR2 = 0.1035
Residual electron density	2.288 and -1.766 e·Å ⁻³	0.727 and -0.778 e·Å ⁻³
CCDC Number	2233556	2233557

Table S2. Summary of the structure refinement data of complexes **7** and **8**.

Empirical formula	C ₃₀ H ₃₇ B ₉ N ₂ O ₂ Pt (7)	C ₃₀ H ₃₇ B ₉ N ₂ O ₂ Pd (8)
Formula weight	749.99	661.30
Temperature	130(2) K	130(2) K
Wavelength	71.073 pm	71.073 pm
Crystal system	Monoclinic	Monoclinic
Space group	<i>P</i> 2 ₁ / <i>c</i>	<i>P</i> 2 ₁ / <i>c</i>
Unit cell dimensions	a = 1516.72(2) pm	1517.09(1) pm
	b = 1122.53(1) pm	1126.62(1) pm
	c = 1908.55(2) pm	1906.74(2) pm
	α = 90°	90°
	β = 108.172(1)°	108.367(1)°
	γ = 90°	90°
Volume	3.08736(6) nm ³	3.09295(5) nm ³
Z	4	4
Density (calculated)	1.614 mg/m ³	1.420 mg/m ³
Absorption coefficient	4.578 mm ⁻¹	0.633 mm ⁻¹
<i>F</i> (000)	1480	1352
Crystal size	0.25 × 0.10 × 0.03 mm ³	0.40 × 0.30 × 0.15 mm ³
Theta range for data collection	2.134 to 32.609°	2.129 to 34.785°
Index ranges	-22 ≤ h ≤ 22, -16 ≤ k ≤ 16, -28 ≤ l ≤ 28	-24 ≤ h ≤ 24, -18 ≤ k ≤ 17, -30 ≤ l ≤ 30
Reflections collected	58096	88279
Independent reflections	10663 [R(int) = 0.0467]	12812 [R(int) = 0.0342]
Completeness (theta)	100.0 % (30.51°)	100.0 % (33.14°)
Absorption correction	Semi-empirical from equivalents	Semi-empirical from equivalents
Max. and min. transmission	1.00000 and 0.66474	1.00000 and 0.98289
Refinement method	Full-matrix least-squares on F ²	Full-matrix least-squares on F ²
Data / restraints / parameters	10663 / 0 / 400	12812 / 0 / 545
Goodness-of-fit on F ²	1.022	1.049
Final R indices [I > 2σ(I)]	R1 = 0.0292, wR2 = 0.0592	R1 = 0.0269, wR2 = 0.0610
R indices (all data)	R1 = 0.0397, wR2 = 0.0633	R1 = 0.0332, wR2 = 0.0640
Residual electron density	1.382 and 1.740 e·Å ⁻³	1.079 and -0.711 e·Å ⁻³
CCDC Number	2233558	2233559

3. NMR study

The comparison of the chemical shifts in ^1H NMR spectra of the ligand **4** and the resulting complexes **5** and **6** after complexation with the respective metal complex precursors evidenced the formation of the complexes. The signals 1 and 10 of compounds **5** and **6** are the most affected ones. They are shifted downfield. When the chloride ligands are exchanged with *nido*-carborate the signals of the ligand are shifted upfield. The coordination of the *nido*-carborate dianion results in the formation of two isomers for **7** and **8** which are not distinguishable with NMR spectroscopy.

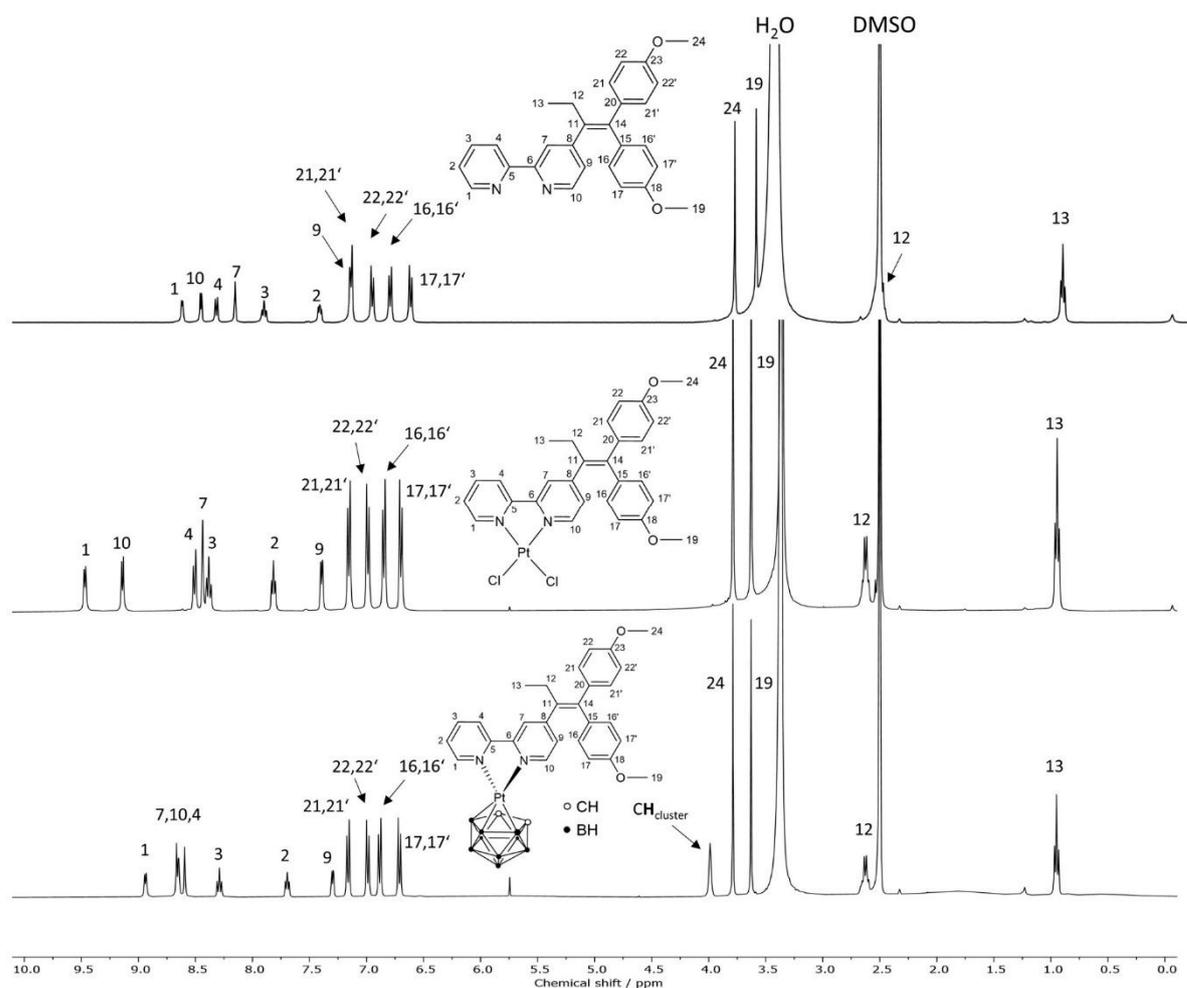


Figure S27. ^1H NMR spectrum of ligand **4** (top), platinum dichloride complex **5** (middle) and platinacarborane complex **7** (bottom) in DMSO- d_6 . The hydrogen atoms 1, 2, 3, 4, 7, 9 and 10 in **5** are shifted downfield compared to the ligand **4**. On incorporation of *nido*-carborate, the signals 1, 2, 9 and 10 in **7** are shifted upfield and 7 is shifted downfield compared to **5**. Only small shifts of protons 12 in **5** and **7** are observed compared to ligand **4**.

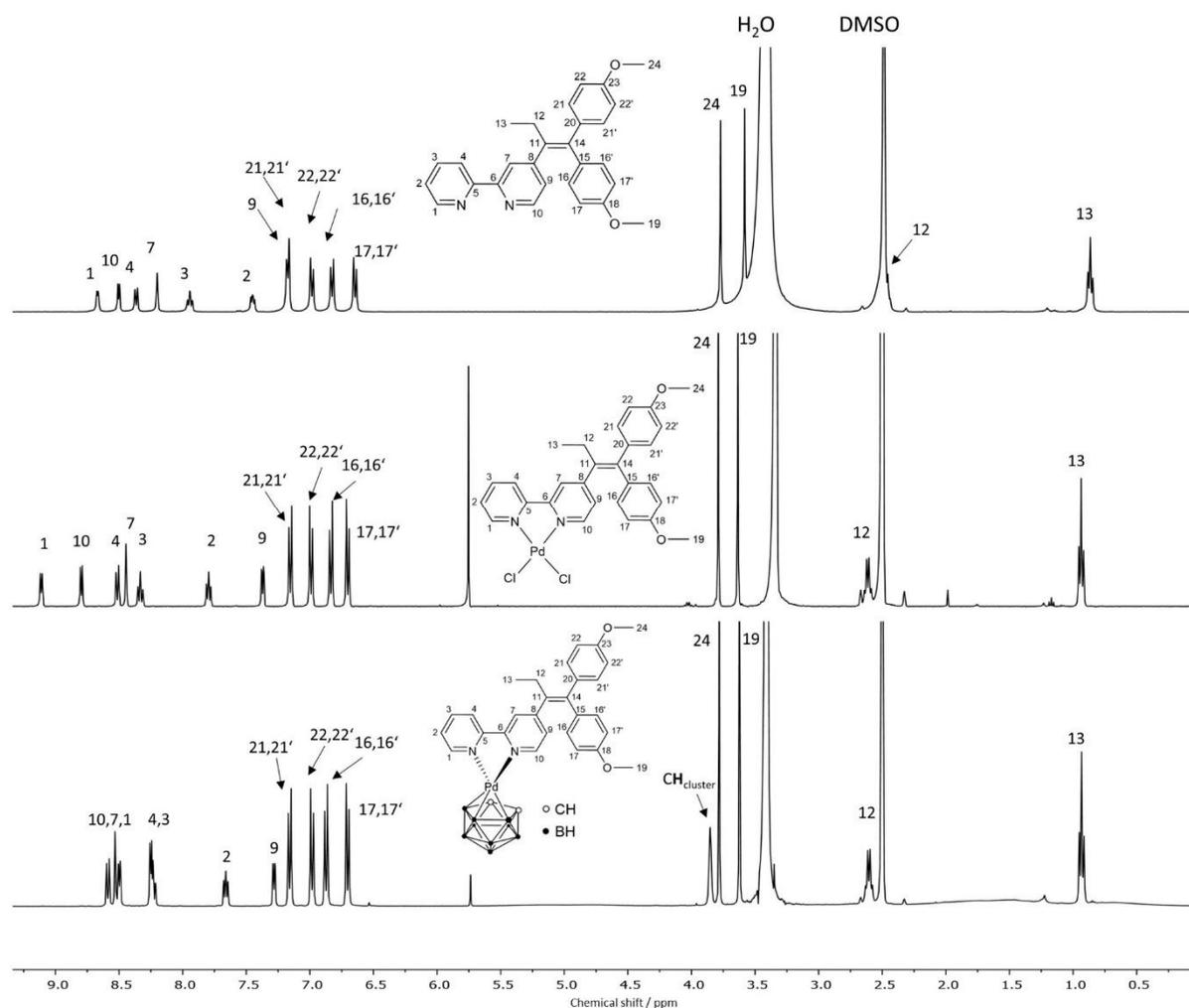


Figure S28. ^1H NMR spectra of ligand **4** (top), palladium dichloride complex **6** (middle) and palladacarborane complex **8** (bottom) in DMSO-d_6 . The hydrogen atoms 1, 2, 3, 4, 7, 9 and 10 in **6** are shifted downfield compared to the ligand **4**. On incorporation of *nido*-carborate the signals 1, 2, 3, 4 and 10 in **8** are shifted upfield and 7 is shifted downfield compared to **6**. Only small shifts of protons 12 in **6** and **8** are observed compared to ligand **4**.

4. Stability

For the *in vitro* tests, stock solutions of the compounds in DMSO were prepared and stored at +4 °C. To assure the stability of the compounds, ^1H NMR spectra were recorded in water-containing DMSO-d_6 in air. The ligand **4** is stable in DMSO solution for one year, as was already shown by Schwarze et al. [5]. Complexes **5–8** were analysed over 30 days exposed to air using ^1H NMR spectroscopy; additionally, $^{11}\text{B}\{^1\text{H}\}$ NMR spectra were recorded for **7** and **8**. The stability tests revealed that complexes **5** and **6** can be stored for at least 30 days without decomposition. Platinacarborane complex **7** exhibited minor changes related to aggregation of this compound after 14 days in DMSO. The palladacarborane complex **8** when stored for up to 3 days in DMSO solution demonstrated minor dissociation with the release of $\text{C}_2\text{B}_9\text{H}_{11}^{2-}$.

4.1. Complex 5

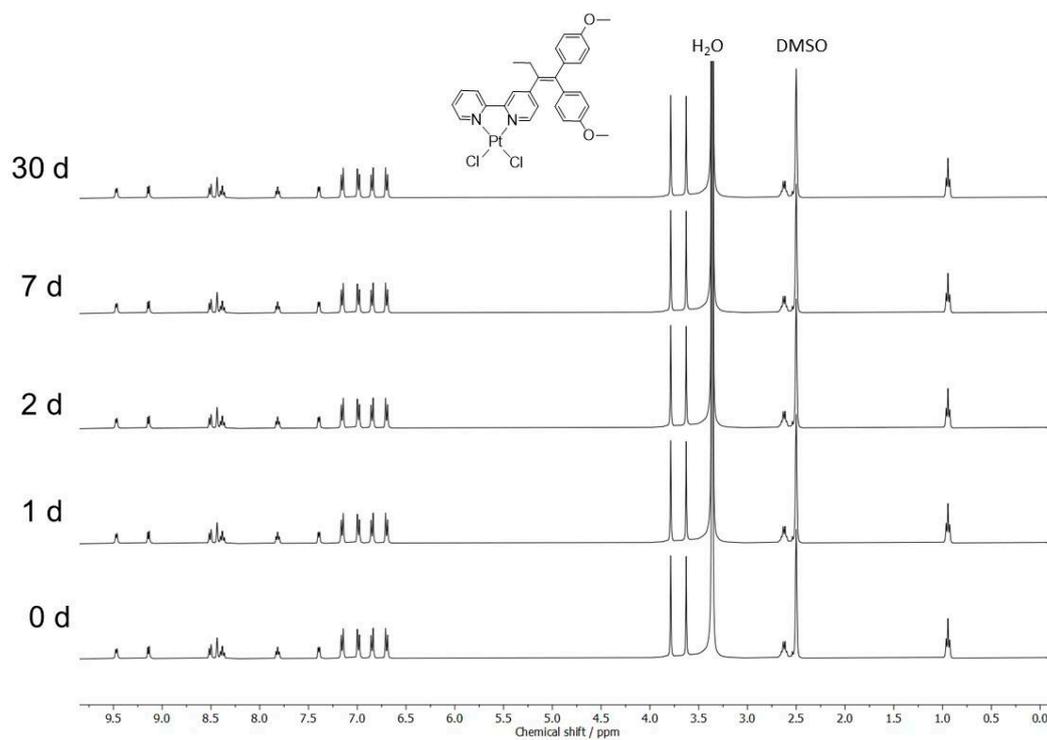


Figure S29. ¹H NMR spectra of **5** in DMSO-d₆ recorded over a month at rt, but stored between the measurements at +4 °C. No changes can be observed.

4.2. Complex 6

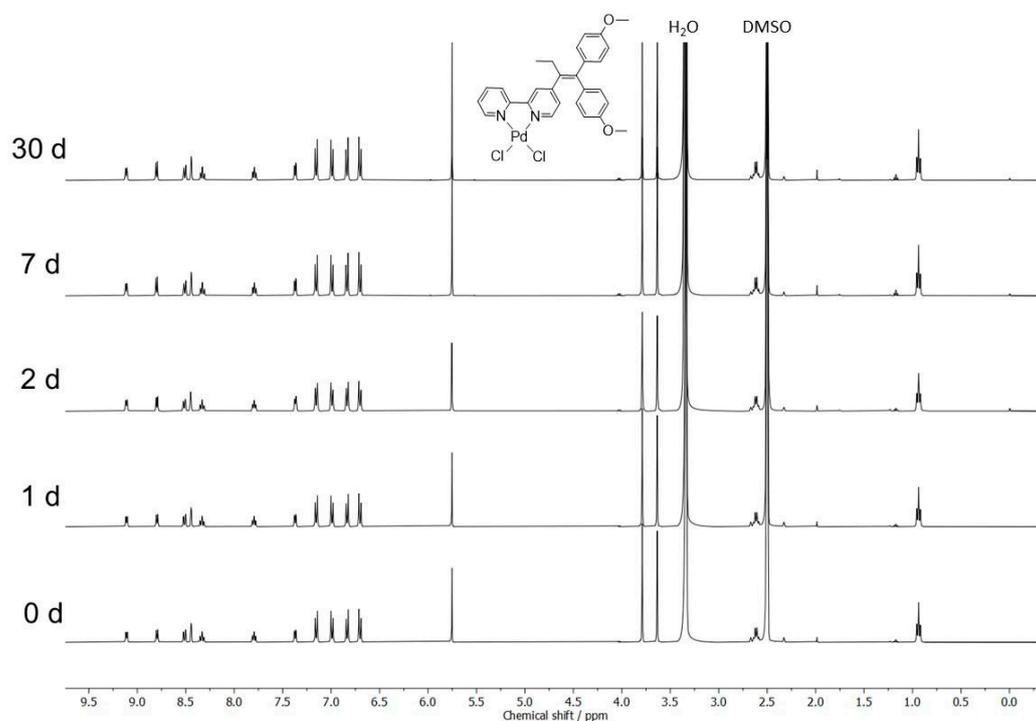


Figure S30. ¹H NMR spectra of **6** in DMSO-d₆ recorded over a month at rt, but stored between the measurements at +4 °C. No changes can be observed.

4.3. Complex 7

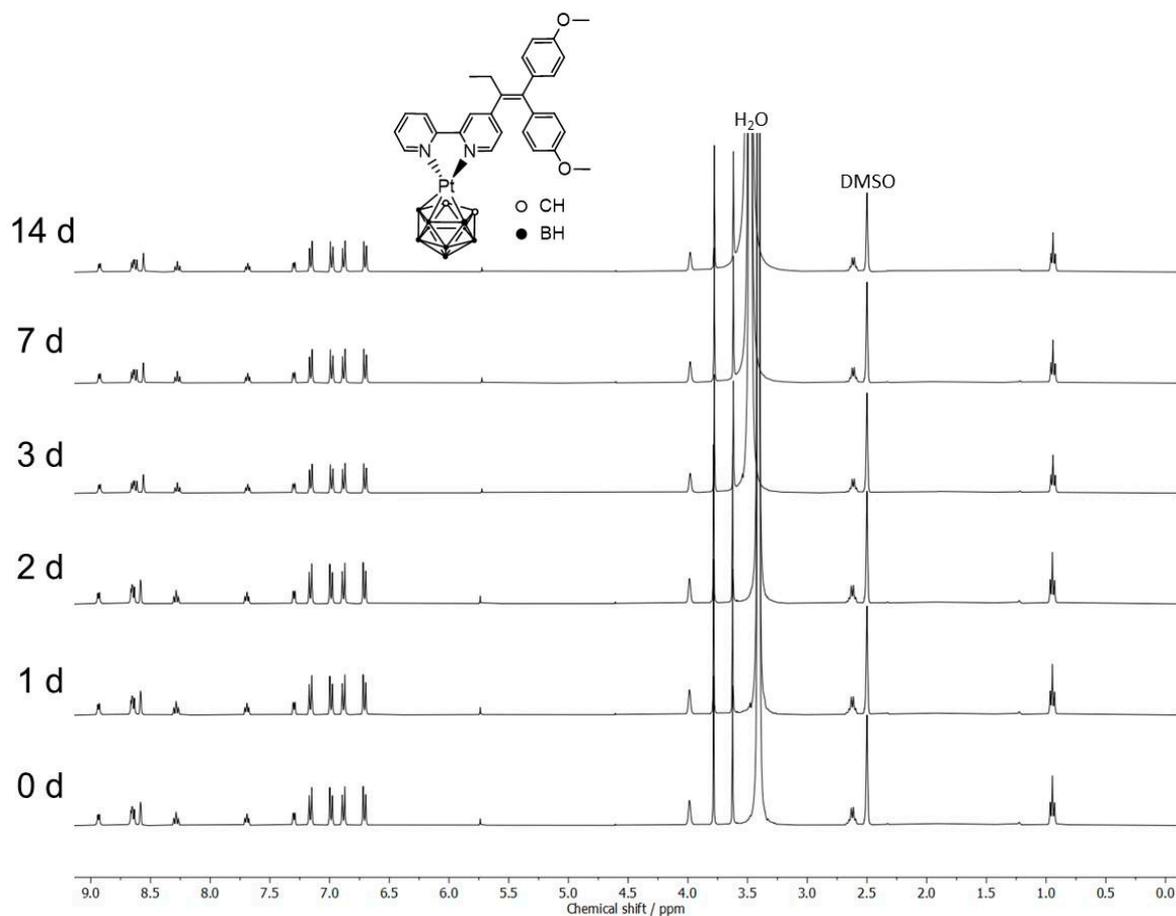


Figure S31. ^1H NMR spectra of 7 in DMSO-d_6 over 14 days at rt, but stored between the measurements at $+4^\circ\text{C}$. No changes can be observed.

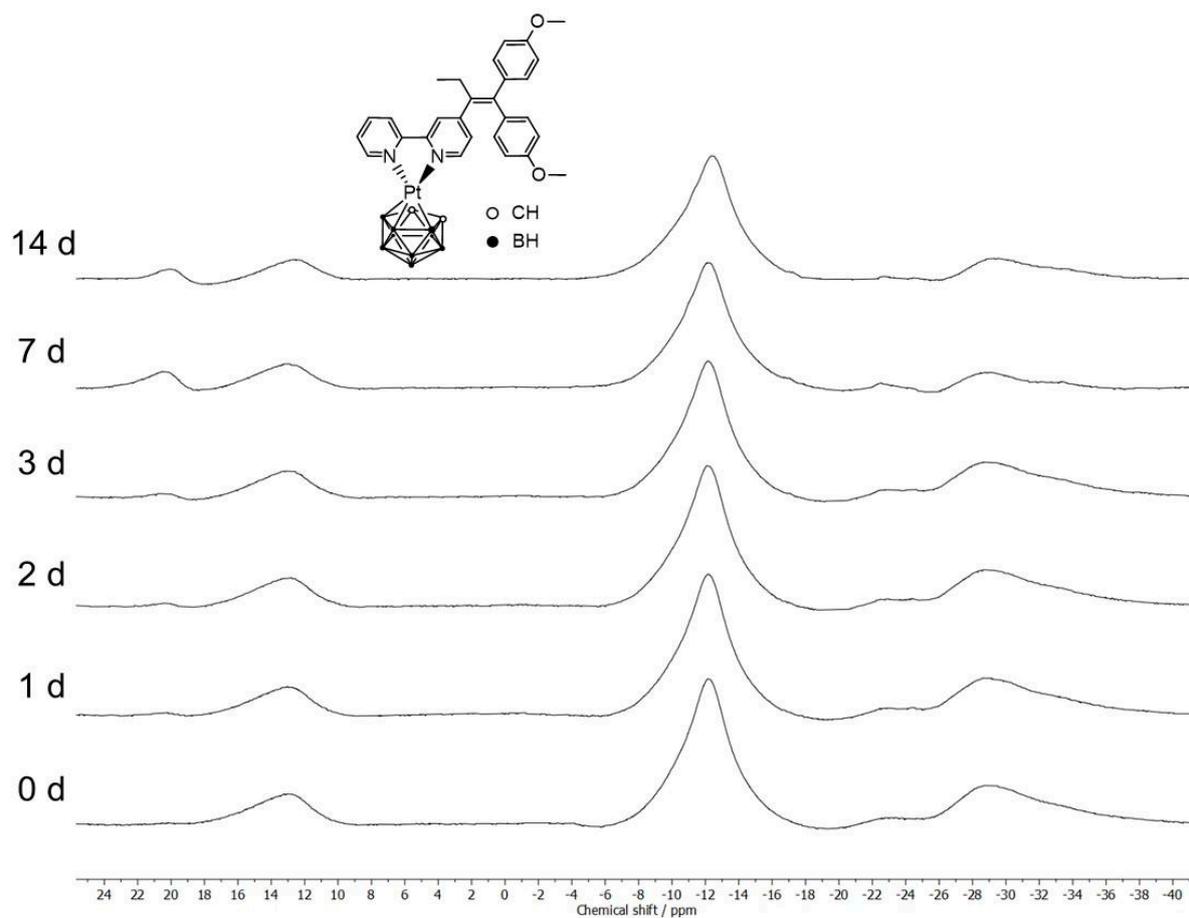


Figure S32. $^{11}\text{B}\{^1\text{H}\}$ NMR of **7** in DMSO-d_6 over 14 days at rt, but stored between the measurements at $+4\text{ }^\circ\text{C}$. Minor changes are observed after 7 days, most likely related to self-assembly.

4.4. Complex 8

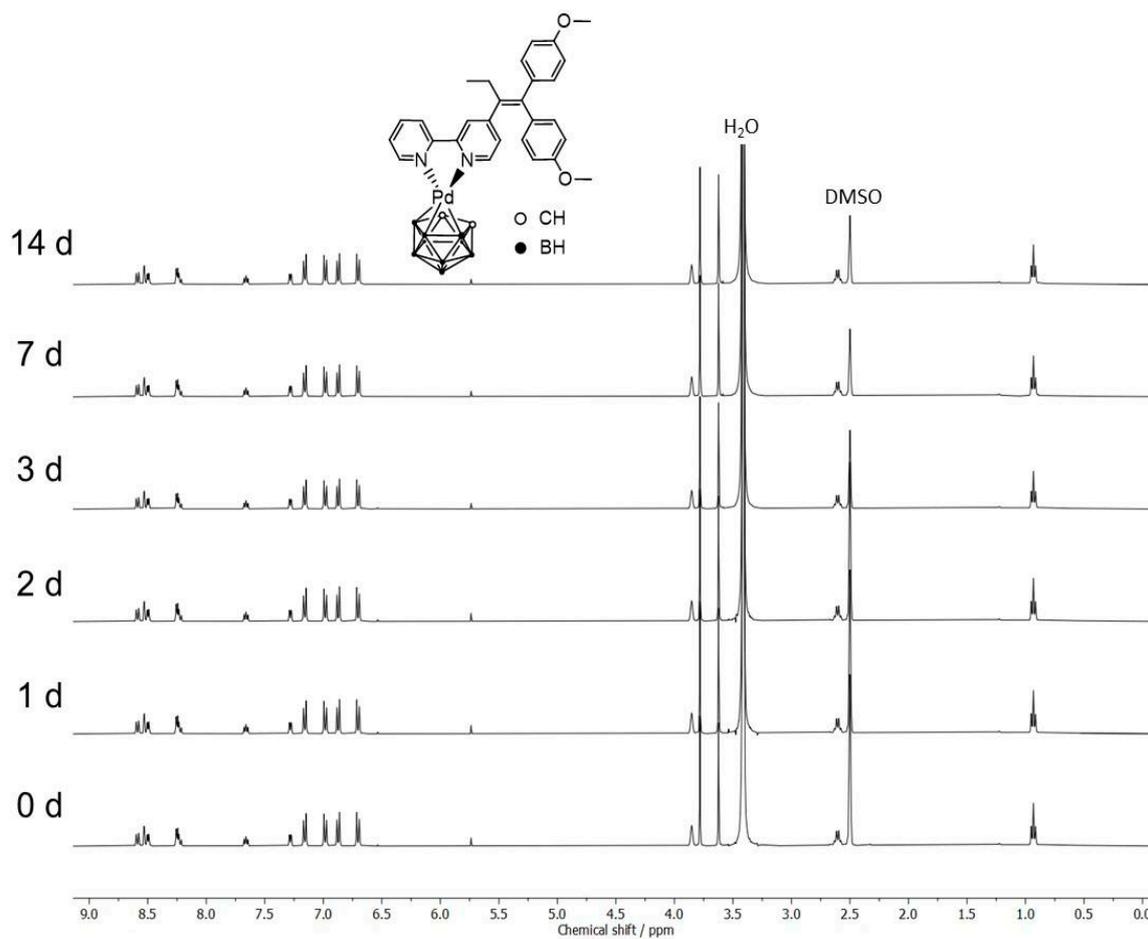


Figure S33. ^1H NMR spectra of **8** in DMSO-d_6 over 14 days at rt, but stored between the measurements at +4 °C. No changes can be observed.

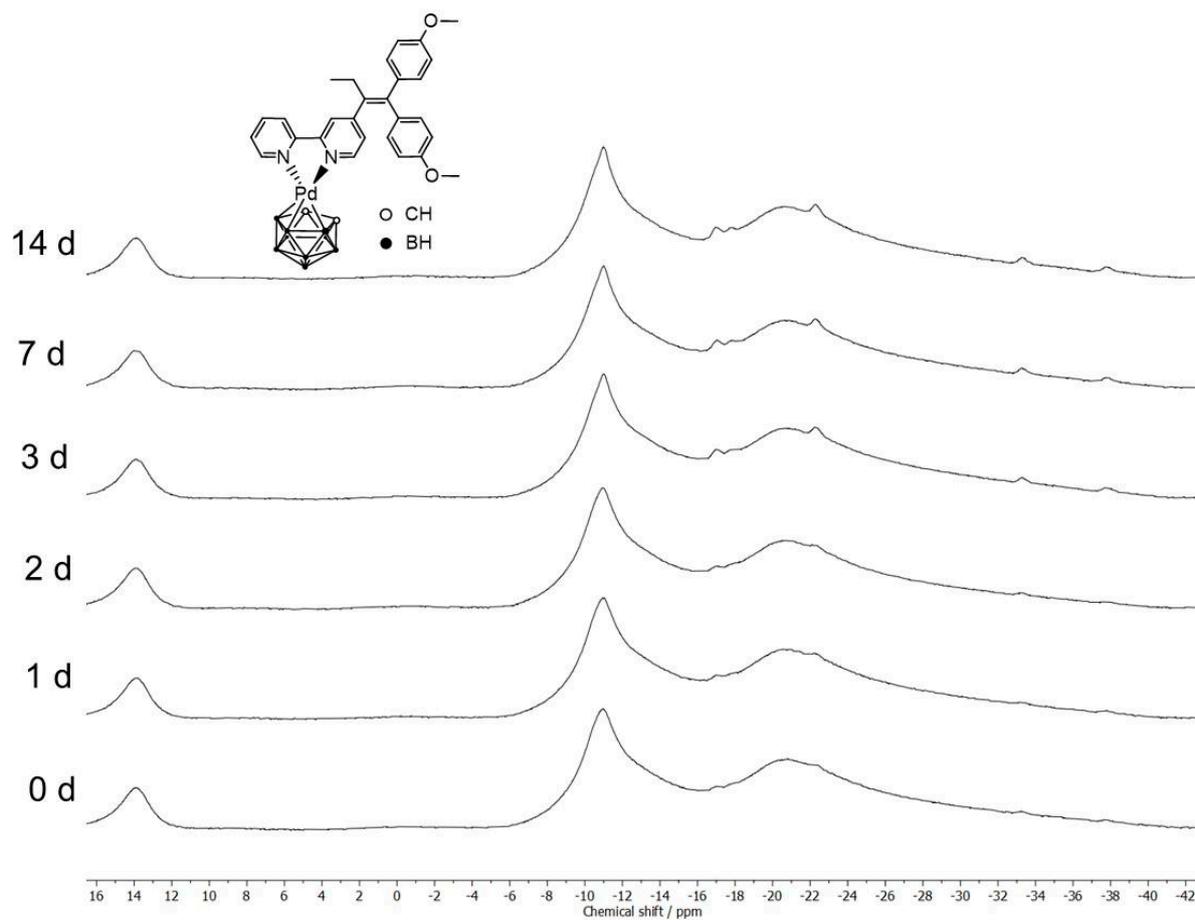


Figure S34. $^{11}\text{B}\{^1\text{H}\}$ NMR spectra of **8** in DMSO-d_6 over 14 days. Minor changes are observed after 3 days revealed by characteristic signals for *nido*-carborate.

of the final step were very low. Considering the multiple synthetic steps for the starting materials **13** and **14**, this approach seemed inefficient.

Table S3. Summary of attempts to prepare metallocarborane complexes **7** and **8** starting from precursor complexes **12**, **13** or **14** and **4**. The reactions were carried out in inert atmosphere using Schlenk technique. The solvents used in the reactions were dried according to the literature [9]. 1 Eq. of ligand **4** was reacted with 1 eq. of **12**, **13** or **14**.

Starting material	Solvent	Conditions	Changes
12 and 4	THF	RT (12 h)	No changes, only starting materials were observed via NMR spectroscopy
	THF	Refluxing (12 h)	No changes, only starting materials were observed via NMR spectroscopy
	DMF	Refluxing (3 h)	Pink, unreacted ligand 4 , decomposition of 12 : black particles (probably of Pt ⁰), release of <i>nido</i> -carborane and boric acid
	Benzene	Refluxing (12 h)	Blue, unreacted ligand 4 , decomposition of 12 : release of <i>nido</i> -carborane
	Toluene	Refluxing (5 h)	Purple, unreacted ligand 4 , decomposition of 12 : formation of black particles and release of <i>nido</i> -carborane
	Ether	RT (12h)	No changes, only starting materials were observed via NMR spectroscopy
	THF	Refluxing with Zn powder (12 h)	No changes, only starting materials were observed via NMR spectroscopy
	DCM	LED lamp, RT (12 h)	Colour changed slightly, 12 decomposed slightly, release of <i>nido</i> -carborane
13 and 4	THF	HBF ₄ ·Et ₂ O, -80 °C	Colour changed from brown to deep brown, yield of 7 up to 5%
14 and 4	THF	HBF ₄ ·Et ₂ O, -80 °C	Colour changed from brown to orange-brown, yield of 8 up to 5%

6. Computational Chemistry

6.1. Geometry optimisation

Before docking and further studies with QTAIM, the structures of compounds 4–8 were fully optimised. The geometry optimisation was done based on X-ray crystal structures using density functional theory (DFT) [10] and performed with the ORCA 4.2 package [11]. The functional PBE0 has been chosen based on the results of benchmark studies for a set of the transition metal complexes [12,13]. In order to improve the stability of the optimised geometry Becke-Johnson dispersion correction of third order (D3BJ) [14] were included in the functional. We applied the density fitting technique resolution-of-identity approximation (RI-J) [15] and chain-of-sphere approximation (COSX) [15] in the geometry optimisation to speed up the calculations. The polarised basis set def2-TZVPP [16] for all non-transition metals and with the additional zeroth order regular approximation (ZORA) [17] for the transition metals was applied for better convergence of the energy. The local minimum of the optimised geometry was verified with numerical frequencies analysis, where no imaginary frequencies were observed.

6.2. UV-vis spectroscopy and transitions

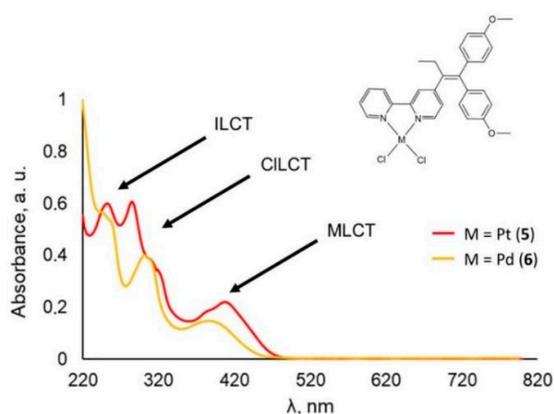


Figure S35. UV-vis spectra of compounds 5 and 6 recorded in acetonitrile.

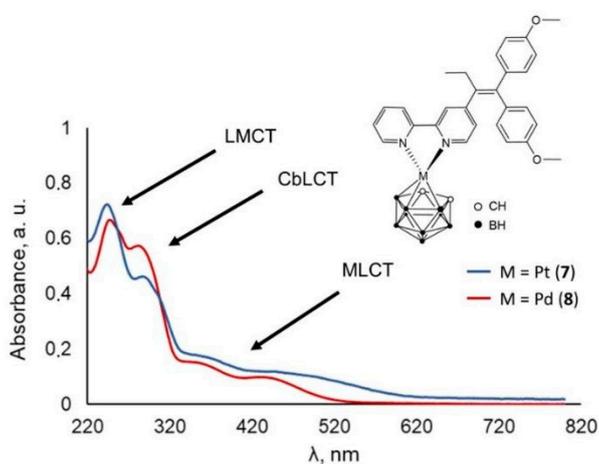


Figure S36. UV-vis spectra of compounds 7 and 8 recorded in acetonitrile.

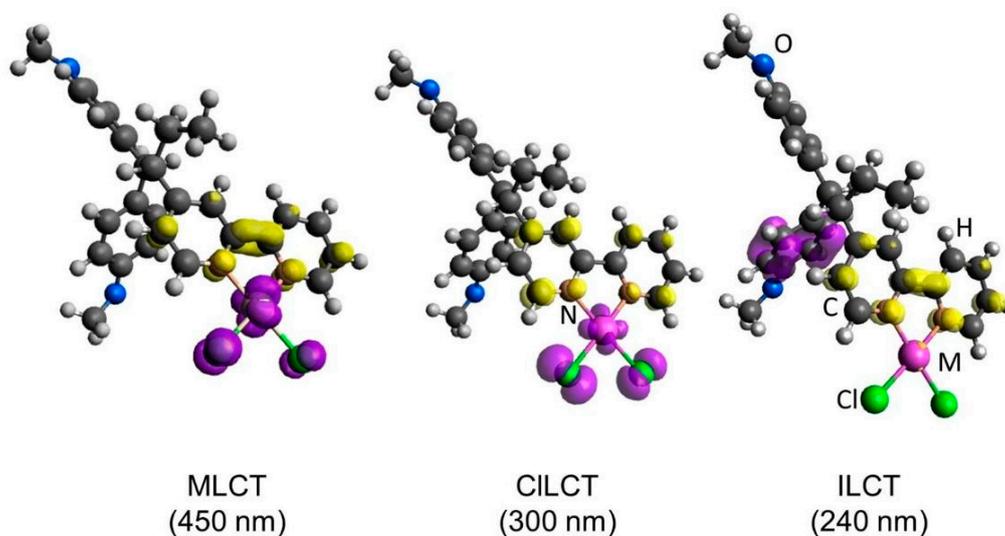


Figure S37. Difference densities of **5** and **6** visualising the charge transfer at certain wavelengths proceeding from purple to yellow iso-surfaces (L = ligand **4**, iso-value = 0.004). MLCT = metal-to-ligand **4** charge transfer, CILCT = chloride-to-ligand **4** charge transfer, ILCT = intraligand **4** charge transfer.

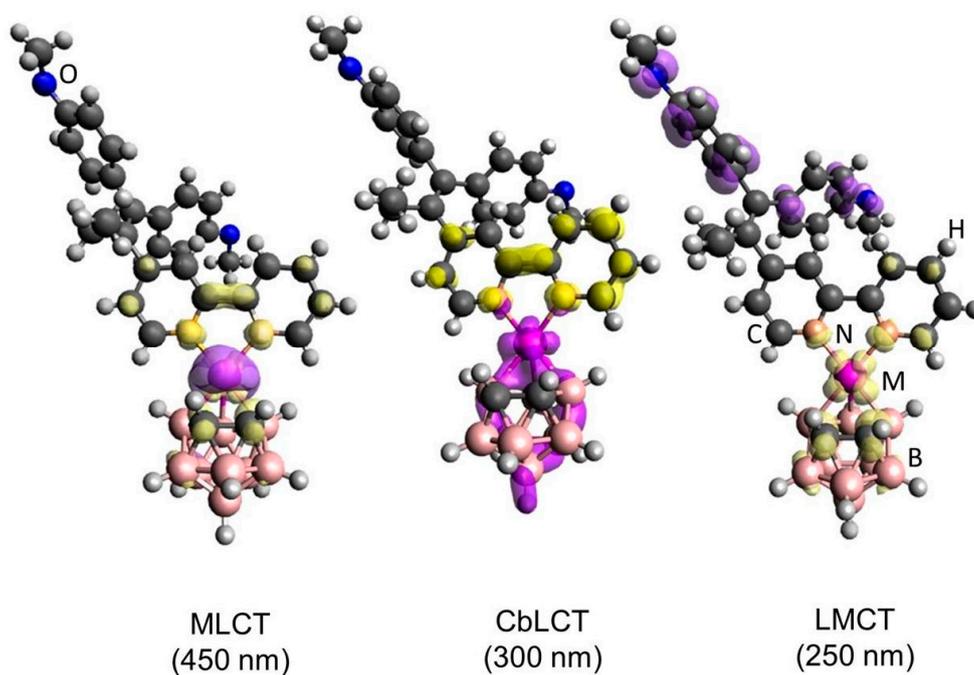


Figure S38. Difference densities of **7** and **8** visualising the charge transfer at certain wavelengths proceeding from purple to yellow iso-surfaces (L = ligand **4**, Cb = $[\text{C}_2\text{B}_9\text{H}_{11}]^{2-}$, iso-value = 0.004). MLCT = metal-to-ligand **4** charge transfer, CbLCT = $[\text{C}_2\text{B}_9\text{H}_{11}]^{2-}$ -to-ligand **4** charge transfer, LMCT = ligand **4**-to-metal charge transfer.

6.3. QTAIM: bonding interactions

In order to estimate the bonding interactions in the transition metal complexes Quantum Theory of Atoms in Molecules (QTAIM) [18] was applied. According to this theory, the distribution of the electron density of the molecular system is structured in critical points (CPs) [18, 19], which fully characterise this system. The first ($\nabla \rho(r)$, gradient) and the second derivatives ($H\rho(r)$, Hessian matrix (1)) [18, 19] of the electron density $\rho(\mathbf{r})$ reveal four types of CPs, which are the maxima, minima or saddle points of the electron density (ED).

$$H\rho(r) = \begin{bmatrix} \frac{\partial^2 \rho}{\partial x^2} & \frac{\partial^2 \rho}{\partial x \partial y} & \frac{\partial^2 \rho}{\partial x \partial z} \\ \frac{\partial^2 \rho}{\partial y \partial x} & \frac{\partial^2 \rho}{\partial y^2} & \frac{\partial^2 \rho}{\partial y \partial z} \\ \frac{\partial^2 \rho}{\partial z \partial x} & \frac{\partial^2 \rho}{\partial z \partial y} & \frac{\partial^2 \rho}{\partial z^2} \end{bmatrix} \quad (1)$$

In the CP, the gradient of the ED is equal to zero. When the molecule or molecular system is in the equilibrium state, matrix (1) can be diagonalised (all non-diagonal elements are zeros). The sum of the main diagonal elements of the matrix (1) is the Laplacian of ED (eq. (2)).

$$\nabla^2 \rho = \lambda_1 + \lambda_2 + \lambda_3 \quad (2)$$

According to the sum of the signs of λ , the four types of stable CPs can be distinguished: (3,-3): maximum, at the atomic nucleus the density reaches the maximum and all λ are negative; (3, -1): saddle point, the interatomic region between two atoms, where λ_1 and λ_2 are negative and λ_3 is positive (bonding CP or BCP); (3, +1): saddle point, the interatomic region between several atoms which form a ring where λ_1 and λ_2 are positive and λ_3 is negative; (3, +3): minimum, interatomic region where all λ are positive (molecular cage) [18,20].

In our description, we considered the topological parameters in the bonding critical points (BCPs, determine the interaction between two atoms) and in the ring critical points (RCPs). The RCPs indicate the interaction of several atoms, which form the ring. In order to classify these interactions, we considered such parameters as bond length, electron density (ED, ρ_{cp}), Laplacian of electron density ($\nabla^2 \rho_{cp}$), density of potential energy (V_{cp}), density of the full electron energy (H_{cp}) and the ratio $|V_{cp}|/G_{cp}$ in CPs. Positive values of the Laplacian in BCPs indicate the depletion of the electron charge, while negative values demonstrate its concentration. The ratio of the potential and kinetic energy densities $|V_{cp}|/G_{cp}$ allows to classify the interactions into three groups: shared (covalent and polar-covalent with $|V_{cp}|/G_{cp} > 2$, $\nabla^2 \rho_{cp} < 0$, $H_{cp} < 0$), transit (intermediate with $1 < |V_{cp}|/G_{cp} < 2$, $\nabla^2 \rho_{cp} > 0$, $H_{cp} < 0$), and closed-shell interactions ($|V_{cp}|/G_{cp} < 1$, $\nabla^2 \rho_{cp} > 0$, $H_{cp} > 0$) [19,21]. The greater the values of $|V_{cp}|$ compared to G_{cp} (and the greater the ratio $|V_{cp}|/G_{cp}$), the higher is the covalent character of the interaction. In general, in spite of the depletion of ED, at the BCPs of coordination bonds ($\nabla^2 \rho_{cp} > 0$) these bonds are stabilised by the negative values of the full electron energy density ($H_{cp} < 0$). The more negative H_{cp} the higher the stability of the interactions (Table S5).

Table S5. Topological properties of the electron density in the bond and ring critical points of complexes 5–8 and [3-(1',2':5',6'- η^4 -COD)-*closo*-3,1,2-PtC₂B₉H₁₁] (12)

Complex	Bond or ring	Bond length, Å	ρ_{cp} , a.u.	$\nabla^2 \rho_{cp}$, a.u.	V_{cp} , a.u.	G_{cp} , a.u.	H_{cp} , a.u.	$ V_{cp} /G_{cp}$
5	N1-Pt	2.019	0.131	0.465	-0.219	0.168	-0.051	1.3
	N2-Pt	2.021	0.133	0.462	-0.220	0.168	-0.052	1.3
	Cl1-Pt	2.305	0.108	0.210	-0.142	0.097	-0.045	1.5
	Cl2-Pt	2.291	0.110	0.210	-0.143	0.097	-0.046	1.5
	Pt-N1-C5-C6-N2	-	0.322	-0.969	-0.457	0.101	-0.356	4.3
	Pt-Cl1-H1-C1-N1	-	0.109	0.220	-0.144	0.100	-0.044	1.4
	Pt-Cl2-H10-C10-N2	-	0.014	0.061	-0.011	0.013	0.002	0.8
6	N1-Pd	2.017	0.113	0.452	-0.173	0.141	-0.032	1.2
	N2-Pd	2.020	0.111	0.448	-0.178	0.146	-0.032	1.2
	Cl1-Pd	2.276	0.098	0.236	-0.121	0.090	-0.031	1.5
	Cl2-Pd	2.286	0.099	0.235	-0.122	0.090	-0.032	1.5
	Pd-N1-C5-C6-N2	-	0.326	-0.991	-0.464	0.108	-0.356	4.3
	Pd-Cl1-H1-C1-N1	-	0.096	0.236	-0.122	0.090	-0.317	1.4
	Pd-Cl2-H10-C10-N2	-	0.013	0.057	-0.010	0.012	0.002	0.8
7	B4-Pt	2.191	0.099	-0.012	-0.088	0.043	-0.045	2.1
	B8-Pt	2.183	0.095	0.035	-0.095	0.052	-0.043	1.8
	B7-Pt	2.179	0.100	-0.014	-0.089	0.043	-0.046	2.1
	N1-Pt	2.077	0.112	0.398	-0.175	0.137	-0.038	1.3
	N2-Pt	2.082	0.114	0.401	-0.179	0.140	-0.039	1.3
	Pt-(B4, B7) (Pt-cage)	-	0.042	0.144	-0.042	0.039	-0.003	1.1
	Pt-N1-C5-C6-N2	-	0.121	-0.121	-0.108	0.039	-0.069	2.8
8	B4-Pd	2.178	0.089	0.031	-0.081	0.044	-0.037	1.8
	B8-Pd	2.270	0.078	0.109	-0.079	0.053	-0.026	1.5
	B7-Pd	2.190	0.090	0.029	-0.082	0.045	-0.037	1.8
	N1-Pd	2.111	0.096	0.400	-0.144	0.122	-0.022	1.2
	N2-Pd	2.098	0.094	0.395	-0.140	0.119	-0.021	1.2
	Pd-(B4, B7) (Pd-cage)	-	0.042	0.156	-0.044	0.041	-0.003	1.0
	Pd-N1-C5-C6-N2	-	0.120	-0.120	-0.107	0.038	-0.069	2.8
12	C1'-Pt	2.189	0.120	0.105	-0.141	0.083	-0.058	1.7
	C2'-Pt	2.201	0.101	0.142	-0.118	0.077	-0.041	1.5
	C5'-Pt	2.178	0.098	0.154	-0.115	0.077	-0.038	1.5
	C6'-Pt	2.135	0.120	0.108	-0.142	0.084	-0.058	1.7
	C1-Pt	2.337	0.093	0.179	-0.113	0.079	-0.034	1.4
	B4-Pt	2.235	0.082	0.052	-0.078	0.045	-0.033	1.7
	B8-Pt	2.274	0.099	-0.047	-0.091	0.040	-0.051	2.3
	B7-Pt	2.191	0.081	0.038	-0.073	0.041	-0.032	1.8

Short bond length, high ρ_{cp} values and lower H_{cp} of N–M compared to Cl–M evidence stronger bonding between the nitrogen atoms and the metal cations than between Cl and M. Despite slightly higher covalency of the Cl–M bond ($|V_{cp}|/G_{cp}$ of the N–M bond is insignificantly lower than this ratio of Cl–M), the parameters in RCP demonstrate that the ED concentrates in the M–N1–C5–C6–N2–M ring with great stabilising negative values of $\nabla^2 \rho_{cp}$

and H_{cp} . The interaction between Pt^{2+} or Pd^{2+} and ligand **4** is stronger than between the metal cations and the chlorides. Chloride ligands in complexes **5** and **6** form H bonds and, thus, close the rings $M-Cl-H-C-N-M$. The topological features of the RCPs, however, demonstrate that the formed rings are less stable than those containing the bidentate 2,2'-bpy ligand.

Several interesting features were observed for BCPs when considering the pair atomic interactions between polyhedral cluster and metal cations. For instance, even though the distances between the carbon atoms of the *nido*-carborate and the metal cations are less than the sum of the Van der Waals radii, the bond paths $C_{cluster}-M$ are not formed, while BCPs and bond paths exist between M and B atoms. However, in comparison to each other, B-Pt and B-Pd bonds show different character. For example, the platinum metal in compound **7** binds to B4 and B7 with a higher covalency compared to B8. This is evidenced by the negative values of the Laplacian, the lower stabilising H_{cp} and the ratio $|V_{bcp}|/G_{bcp}$ being slightly greater than 2. In palladacarborane complex **8**, B7-Pd and B4-Pd bonds are shorter and exhibit more covalent character than B8-Pd (lower $\nabla^2\rho_{cp}$ values, the total energy density is more negative and the ratio $|V_{cp}|/G_{cp}$ is closer to 2 in the BCPs for B7-Pd and B4-Pd than for B8-Pd). However, when we compare the binding nature of the sum of B-Pt and B-Pd bonds, we observe that the B-Pt bonds are of greater covalent character than the B-Pd bonds. The lower ρ_{cp} values and only positive $\nabla^2\rho_{cp}$ at the BCPs of B-Pd indicate the depletion of the electron charge for these bonds. In other terms, the bonding interactions between $[C_2B_9H_{11}]^{2-}$ and Pt^{2+} are stronger than between $[C_2B_9H_{11}]^{2-}$ and Pd^{2+} . This may reveal different biological properties of the compounds **7** and **8**, depending on the nature of the metal, when stability of these bonds is a key property.

The same analysis was performed for **12**. Although the bonding character of B-Pt exhibits equal or even higher extent of covalency, the ED values in the BCPs of B-Pt are lower than those observed for Pt-COD. The sum of H_{cp} for Pt-COD bonds are lower than for the (*nido*-carborate)-Pt bonds showing the stronger COD-Pt interaction. This could be a reason why we observed release of *nido*-carborate in some of the trials to replace the COD ligand.

The molecular graph (Figure S30) also demonstrates that one RCP in Pt-(B7,B8) ring is very close to the BCP of B8-Pt. Under certain conditions, these two points overlap with each other which leads to a merging of both the BCP and RCP. This situation is called catastrophe situation [22]. It also indicated the less stable interaction of Pt^{2+} with the *nido* cluster. This additionally explains the higher possibility to release $[C_2B_9H_{11}]^{2-}$ instead of COD (Table S3).

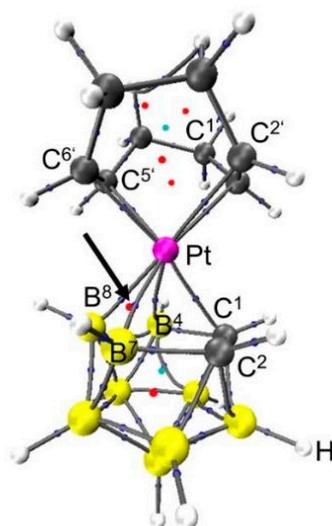


Figure S39. QTAIM analysis: molecular graph of [3-(1',2':5',6'- η^4 -COD)-*closo*-3,1,2-PtC₂B₉H₁₁] (12). The blue dots are BCPs, the red dots are RCPs and the grey lines connecting the nuclei and BCPs are the bond paths. The arrow shows the catastrophe, where the close position of the BCP and RCP may lead to the merge of the bond path.

6.4. Protein structure preparation

The crystal structure of the oestrogen receptor α was prepared for docking. The X-ray structure of the ER α is available from the protein data bank (PDB) under the code 3ERT [23] and contains the dimer with 4-hydroxytamoxifen as ligand in the ligand-binding pocket. One identical monomer was removed from the file and the other with the initial ligand was protonated using the Reduce program [24]. The force field CHARMM [25] was assigned in NAMD [26] for the atoms of the receptor, the ligand and water in the initial structure (3ERT). The molecular dynamic minimisation (Langevin dynamics) [27] with the ligand in the binding pocket of the receptor has been applied in order to avoid the contraction of the binding pocket. The dynamic was run for 500 ps with Langevin thermostat at 298 K. The integration time step is 2 fs/step and the size of the water cell is 70 x 64 x 82 Å³ with the centre placed at 4.55, 2.82, 33.22 of *x,y,z*-coordinates. The rigid model was applied for the water molecules (TIP3P). For the long-range electrostatic interaction, the particle-meshed Edwald method (PME) was used [28]. The coordinates of the minimised energy structure have been taken and 4-hydroxytamoxifen has been removed from the binding pocket for further docking.

6.5. Docking

The molecular docking was performed with the AutoDockTools4 software [29] using the Lamarckian Genetic Algorithm [30]. The force-field parameters for the transition elements (Pt and Pd) and boron atoms were added manually in the parameter file of the AutoDockTools4 library. The water molecules were eliminated and the non-polar hydrogen atoms were merged. The docking area was limited by the constructed grid box of the size 58 x 64 x 88 centred at 32.336, -1.613, 26.672 of *x,y,z*-coordinates (based on the position of the LBD). The following parameters were used in the docking: number of hybrid GA-LS runs: 500; population size: 150; maximum number of energy evaluations: 25,000,000, maximum number of top individuals to survive to next generation: 1; rate of gene mutation: 0.02; rate of crossover:

0.8; Mean of Cauchy distribution for gene mutation: 0.0; variance of Cauchy distribution for gene mutation: 1.0.

All the agonists such as the endogenous 17β -estradiol as well as some of the known synthetic antagonists such as ICI-164384 [31] exclusively bind to the C-terminal ligand-binding domain (LBD) [23]. Tamoxifen as a selective oestrogen receptor modulator (SERM) is an antagonist [23, 32]. Its metabolite 4-hydroxytamoxifen binds non-covalently to the LBD and induces the conformational change of helix 12, which is essential for activation function 2 (AF-2) activity [23]. In order to obtain information about the putative binding modes of compounds 2–9 to the ER α , docking simulations with 3–9 into the LBD of ER α (PDB code: 3ERT; [23]) were conducted using AutoDockTools4 [29]. The best ranked positions are shown in Figure S31 relative to the position of the metabolites 4,4'-dihydroxytamoxifen (3) and 4-hydroxytamoxifen (2).

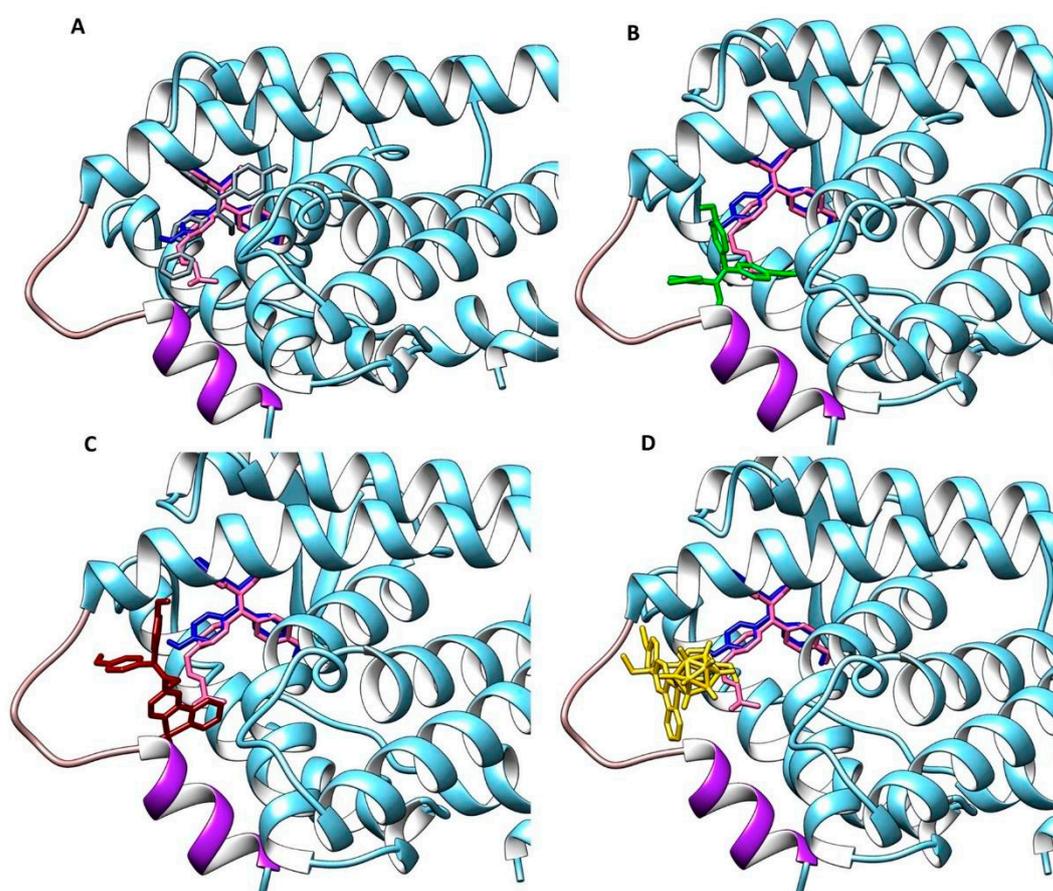


Figure S40. *In silico* investigation of the binding modes of compounds 2–9 based on docking. The highest ranked docked positions of compounds 9 (grey, A), 4 (green, B), 5 or 6 (magenta, C) and 7 or 8 (yellow, D) are shown compared to the position of the 2 (pink) and 3 (dark blue). Helix 12 is shown in purple, and the loop of the receptor is shown in light brown.

First, the binding energy of 3 was estimated by docking the molecule into the X-ray structure of the ER α (PDB code: 3ERT). We observed that the majority of conformations of 3 occupy a similar position as the original metabolite 2 with a scored binding energy value of

$-10.73 \text{ kcal mol}^{-1}$ (the binding energy is $-10.80 \text{ kcal mol}^{-1}$ for **2**, Figure S40). Both metabolites do not bind directly to helix 12 and probably induce the conformational changes of this helix indirectly [23]. Incorporation of 2,2'-bpy in the tamoxifen structure decreased the energy of the interaction with the LBD, with a binding energy of $-8.44 \text{ kcal mol}^{-1}$ (ligand **4**) and $-8.76 \text{ kcal mol}^{-1}$ (ligand **9**). However, the best ranked position of **9** suggests that this compound occupies the same positions as the metabolites **2** and **3**, while the predicted poses show that the best model position of **4** is located at the receptor loop, suggesting direct interaction with the loop and helix 12. Once the 2,2'-bpy unit was combined with the PtCl_2 or PdCl_2 moieties (**5** and **6**, respectively) the binding energies slightly improved (-9.08 and $-8.99 \text{ kcal mol}^{-1}$, respectively). Further, the incorporation of the bulky carborane appears to have a destabilising impact on the binding energy of compound **7** ($-8.14 \text{ kcal mol}^{-1}$). However, the palladium complex **8** showed comparable affinity and orientation, in comparison to compounds **5** and **6**, but with a higher binding energy of $-9.71 \text{ kcal mol}^{-1}$.

7. Bioanalytical measurements

7.1. PI staining on chamber slides

MCF-7 cells were seeded overnight at a density of 1.5×10^3 cells/well followed by treatment with an IC_{50} dose of the compounds **4–6** for 60 h. After incubation, the cells were washed with phosphate-buffered saline (PBS) and fixed with 4% paraformaldehyde (PFA) for 15 min at rt. Thereafter, the cells were washed with PBS and stained with a solution of propidium iodide (PI) in a concentration of $50 \mu\text{g mL}^{-1}$ with 0.1 mM ethylenediaminetetraacetic acid (EDTA) at pH 8.0, 0.1% Triton X-100 and RNase ($85 \mu\text{g mL}^{-1}$) in PBS for 2 min. Afterwards, for covering the slides, fluorescent mounting medium was used. Finally, the slides were analysed with a Zeiss AxioObserver Z1 inverted fluorescence microscope (Carl Zeiss AG, Oberkochen, Germany) at 200x magnification.

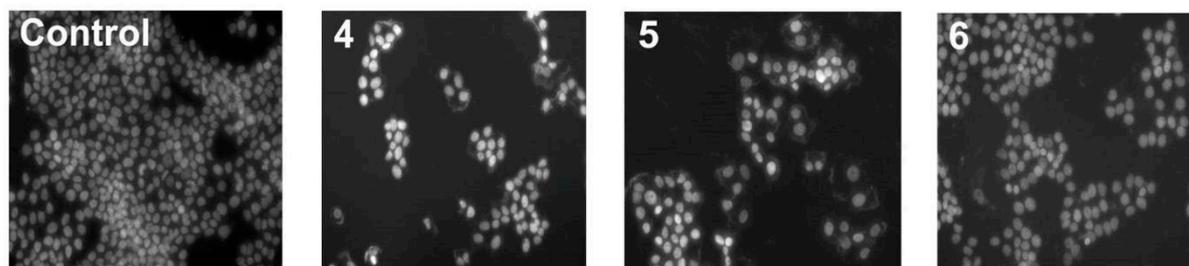


Figure S41. The potential of compounds **4–6** to induce apoptotic cell death. MCF-7 cells were exposed to IC_{50} doses of compounds **4**, **5** and **6**, and after 60 h the cells were analysed by fluorescent microscopy.

7.2. Cell viability

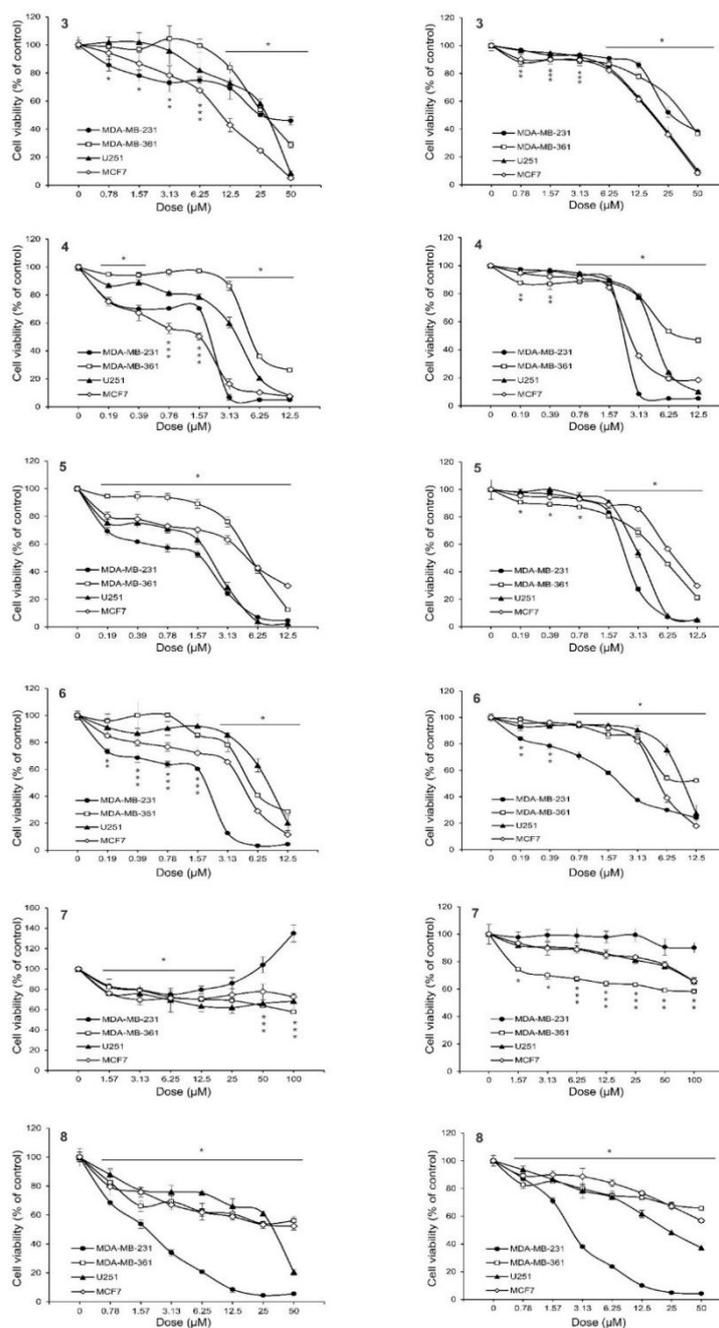


Figure S42. The effect of compounds 3–8 on tumour cell viability. Human breast cancer cell lines MCF-7, MDA-MB-361, MDA-MB-231 and the human glioblastoma cell line U251 were treated with 3–8 for 72 h and subjected to 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT, left) and crystal violet (CV, right) viability assays. The data is expressed as percentage of viability of untreated control (set as 100%) from one representative out of three independent experiments and presented as mean \pm SD of triplicate cultures (* $p < 0.05$ in comparison to untreated cells).

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