



Supplementary material: An efficient transition-metal-free route to oligo- α -pyridylamines via fluoroarenes

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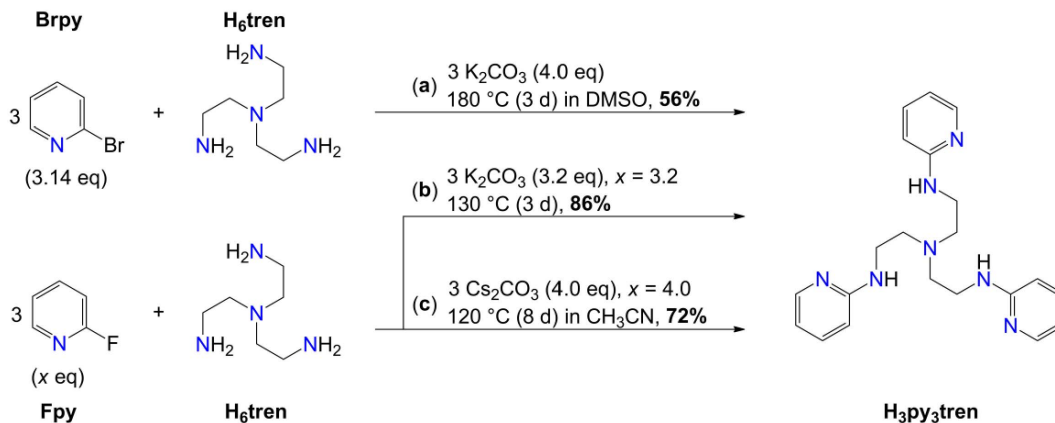
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Scheme S1: Synthesis of H₃py₃tren.

- (1) Synthesis of HBrdpa.
- (2) ¹H NMR spectrum of HBrdpa (Figure S1).
- (3) 2D NMR spectra of HFdpa (Figures S2–S4).
- (4) 2D NMR spectra of H₆tren(dpa)₃ (Figures S5–S8).
- (5) Details on ¹H NMR and ESI-MS detection of **SPa** (Figures S9 and S10, Scheme S2).

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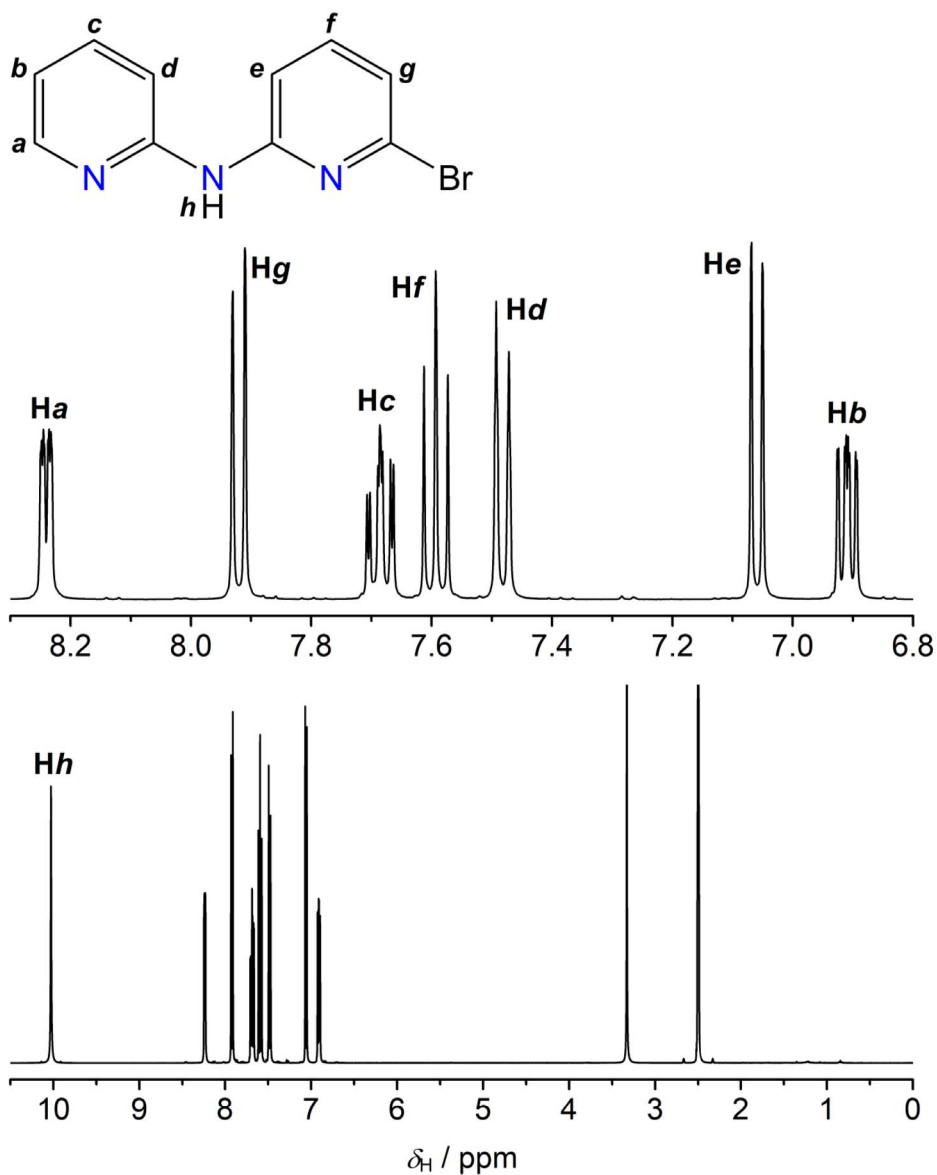
Scheme S1. Synthesis of H₃py₃tren as described in Ref. [1] (a), Ref. [2] (b), and Ref. [3] (c), with quoted isolated yields.

(1) Synthesis of HBRdpa

In a three neck round bottom flask (1 L) equipped with a mechanical overhead stirrer and a condenser, NH₂py (5.2510 g, 55.796 mmol) and Br₂py (13.2155 g, 55.787 mmol) were dissolved in anhydrous toluene (500 mL) under nitrogen atmosphere. ^tBuOK (10.2110 g, 90.999 mmol) was carefully added to the pale-yellow solution, which progressively turned to dark blue/black. The reaction mixture was heated to 80 °C and vigorously stirred for three days. It is worth to note that a mechanical stirrer is essential as a very dense solid mass is formed at a temperature of about 70 °C during the initial heating. This solid is progressively dispersed into the liquid phase during the first 24 h of reaction time. The reaction was monitored through thin-layer chromatography (TLC) on silica gel plates (one drop of reaction mixture in 0.5 mL of CH₂Cl₂; eluent Et₂O:*n*-hexane, 1:1 v/v; R_f(NH₂py) = 0.02; R_f(HBRdpa) = 0.31; R_f(Br₂py) = 0.55) and ¹H NMR spectroscopy in deuterated dimethyl sulfoxide (DMSO-*d*₆). The suspension was allowed to cool to room temperature, and the solvent was evaporated under reduced pressure. The light-

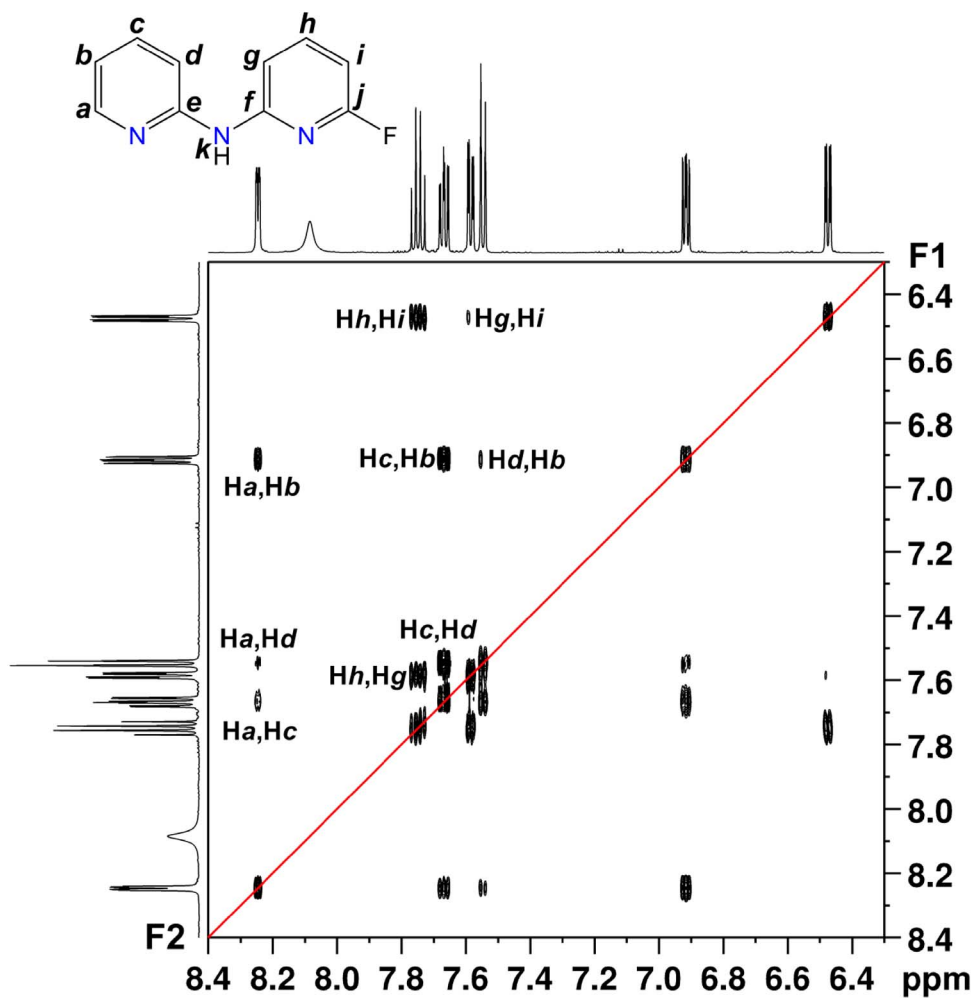
pink solid obtained was dissolved in CH₂Cl₂ (300 mL) and washed with water (150 mL + 2 × 50 mL). The inorganic phases were extracted with CH₂Cl₂ (2 × 50 mL). The organic phases were combined and dried over MgSO₄ (90 min). The desiccant was filtered off and the red organic solution was evaporated to dryness, to give a dark orange oil which slowly crystallized under vacuum (~14.0 g). This crude product was purified through gradient-elution FC (for ~7 g of crude material: SiO₂; eluent petroleum ether:Et₂O, from 1:0 to 1:1 v/v; *t*_{gradient} = 25 min, flow rate = 40 mL/min, column diameter = 40 mm, column length = 150 mm), which gave HBRdpa as a pale-yellow solid (9.01 g, 36.0 mmol, 64.6% isolated yield).

Mp 87.7–88.4 °C. ¹H NMR (DMSO-*d*₆, 298 K, 400.13 MHz): δ_H (ppm) = 10.03 (1H, s, H_h), 8.24 (1H, ddd, ³J(*a*, *b*) = 5.0, ⁴J(*a*, *c*) = 1.9, ⁵J(*a*, *d*) = 0.8, H_a), 7.92 (1H, dd, ³J(*g*, *f*) = 8.3, ⁴J(*g*, *e*) = 0.4, H_g), 7.69 (1H, ddd, ³J(*c*, *d*) = 8.4, ³J(*c*, *b*) = 7.2, ⁴J(*c*, *a*) = 2.0, H_c), 7.59 (1H, pt, ³J(*f*, *e*) ~ ³J(*f*, *g*) ~ 7.9, H_f), 7.48 (1H, dpt, ³J(*d*, *c*) = 8.4, ⁴J(*d*, *b*) ~ ⁵J(*d*, *a*) ~ 0.9, H_d), 7.06 (1H, dd, ³J(*e*, *f*) = 7.5, ⁴J(*e*, *g*) = 0.5, H_e), 6.91 (1H, ddd, ³J(*b*, *c*) = 7.2, ³J(*b*, *a*) = 5.0, ⁴J(*b*, *d*) = 1.0, H_b).

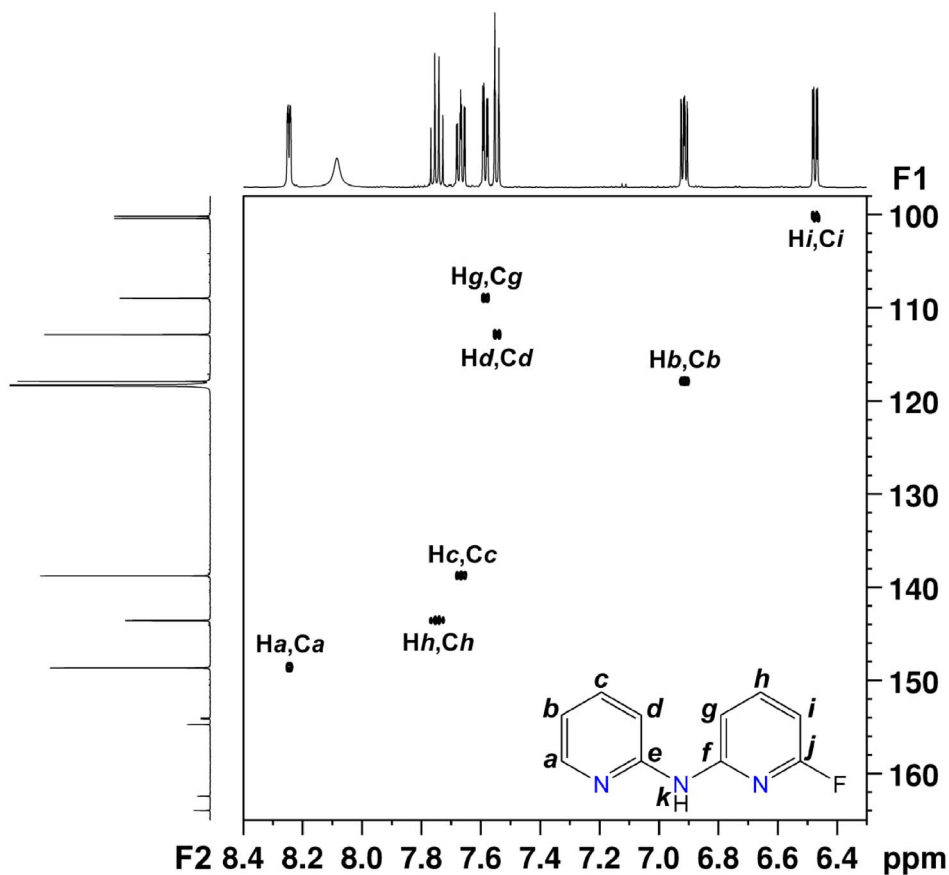
(2) ^1H NMR spectrum of HBrdpa

Supplementary Figure S1. Bottom: ^1H NMR spectrum of HBrdpa in DMSO- d_6 (298 K, 400.13 MHz). Top: atom-labelled structure of HBrdpa and magnification of the spectral region between 8.30 and 6.80 ppm. Processing parameters (TopSpin 4.0.6 [4]): SI = TD, LB = 0.30 Hz. δ_{H} (ppm) = 2.50 (quintet, residual protons in DMSO- d_6), 3.33 (s, water, OH).

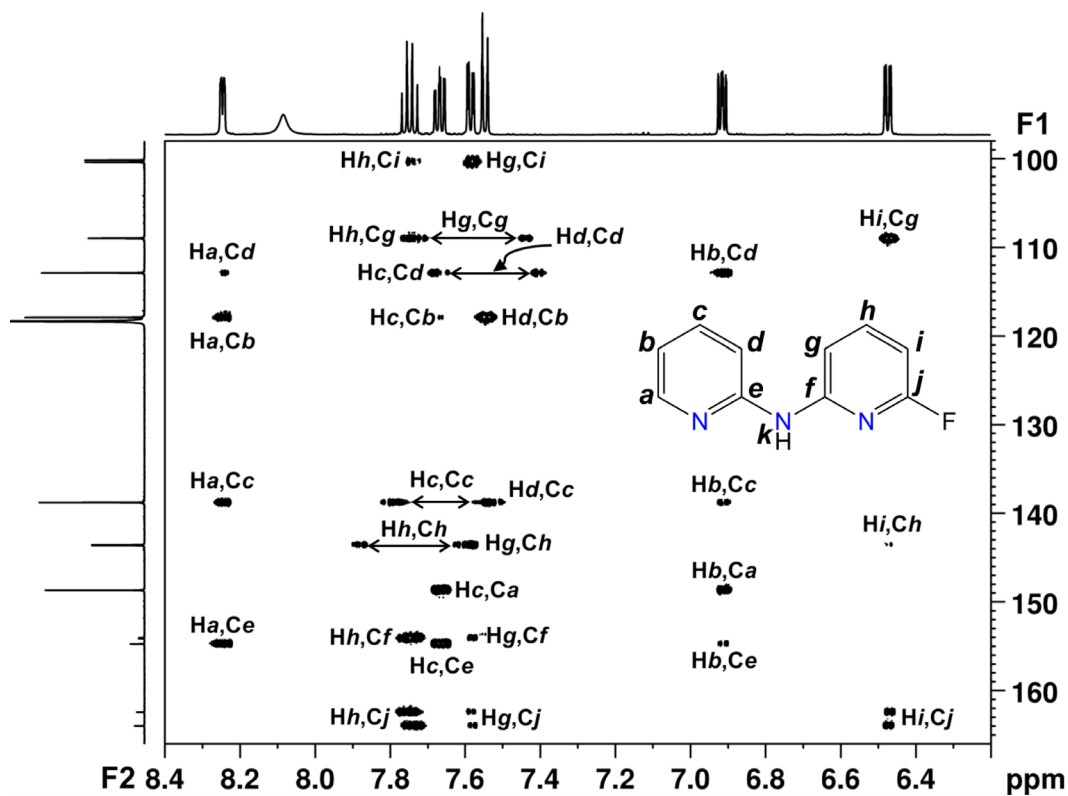
(3) 2D NMR spectra of HFdpa



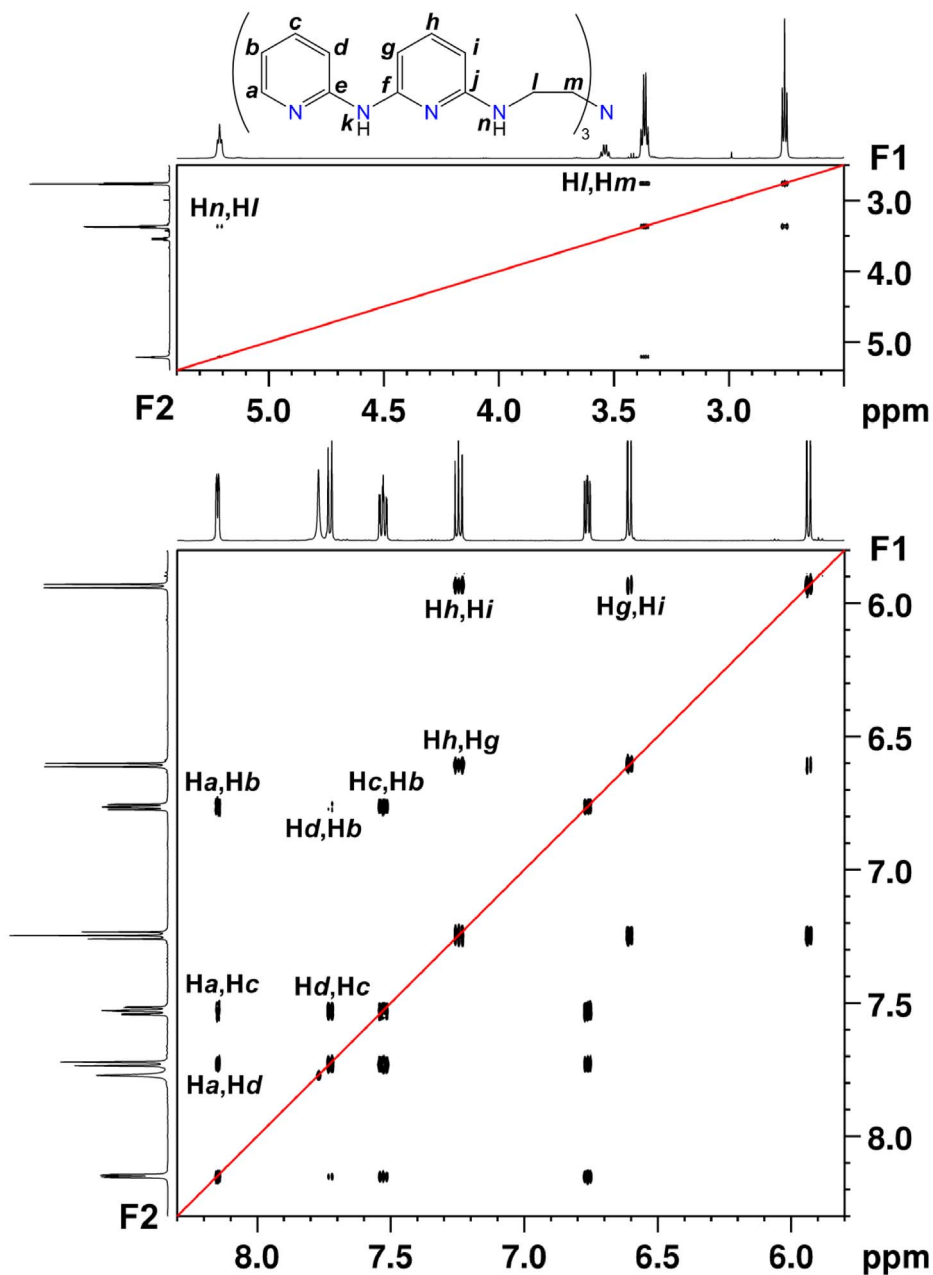
Supplementary Figure S2. Atom-labelled structure and ^1H - ^1H COSY spectrum of HFdpa in CD_3CN between 8.40 and 6.30 ppm (298 K, 600.13 MHz). The labelling of the cross-peaks indicates the ^1H - ^1H coupling (F2, F1). The red line highlights the diagonal peaks. Processing parameters (TopSpin 4.0.6 [4]) for F2 (x axis): SI = TD, LB = 1.00 Hz. Processing parameters for F1 (y axis): SI = 2·TD, LB = 0.30 Hz.



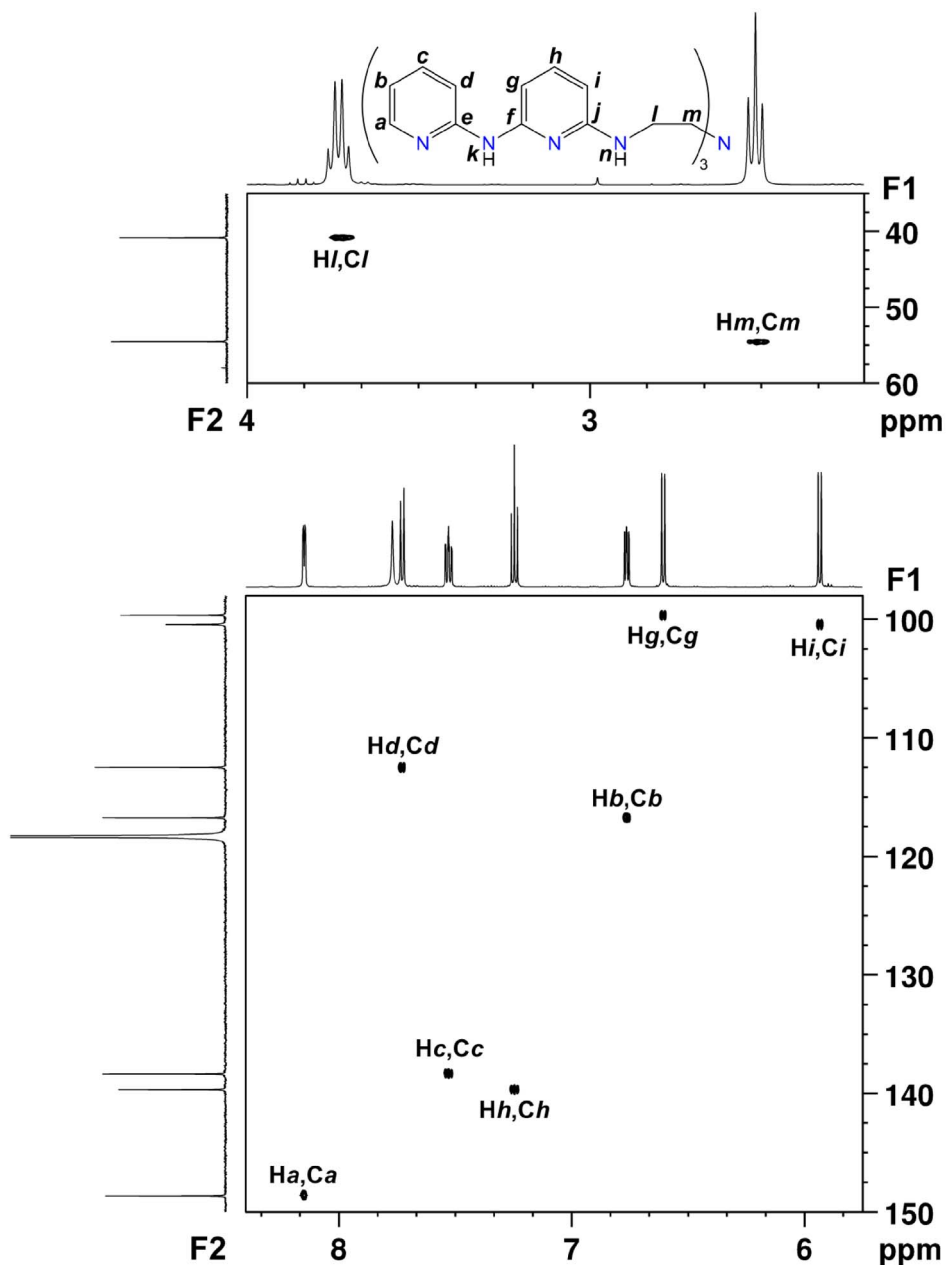
Supplementary Figure S3. Atom-labelled structure and ^1H - ^{13}C HSQC spectrum of HFdpa in CD_3CN for $\delta_{\text{H}} = 8.40\text{--}6.30$ ppm and $\delta_{\text{C}} = 165\text{--}98$ ppm (298 K, 600.13 MHz for F2, 150.90 MHz for F1). The labelling of the cross-peaks indicates the ^1H - ^{13}C coupling (F2, F1). Processing parameters (TopSpin 4.0.6 [4]) for F2 (x axis): SI = 2·TD, LB = 1.00 Hz. Processing parameters for F1 (y axis): SI = 3·TD, LB = 0.30 Hz.



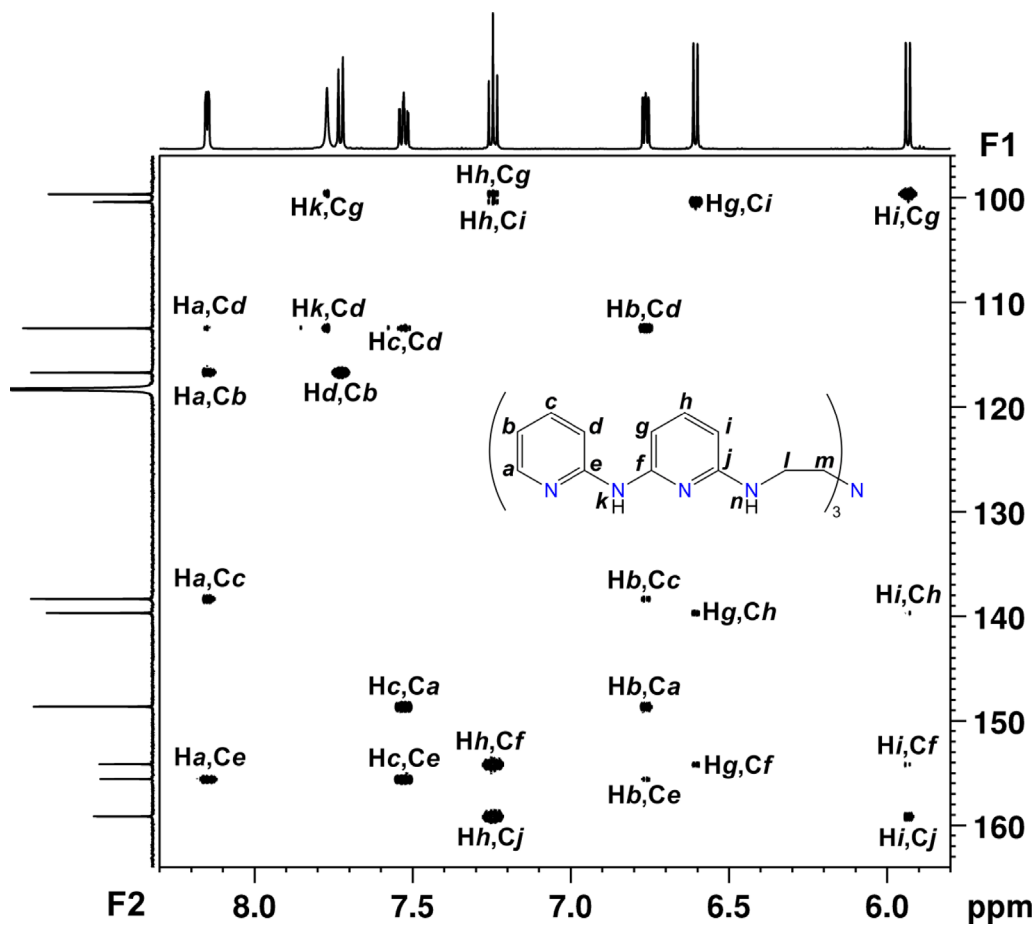
Supplementary Figure S4. Atom-labelled structure and ^1H - ^{13}C HMBC spectrum of HFdpa in CD_3CN for $\delta_{\text{H}} = 8.40\text{--}6.30$ ppm and $\delta_{\text{C}} = 166\text{--}98$ ppm (298 K, 600.13 MHz for F2, 150.90 MHz for F1). The labelling of the cross-peaks indicates the ^1H - ^{13}C coupling (F2, F1). Although a low-pass filter was employed for a better 1J suppression (pulse program: hmbcetgpl3nd), the spectrum exhibits the remanence of 1J couplings for sites *c*, *d*, *g*, and *h*. Processing parameters (TopSpin 4.0.6 [4]) for F2 (x axis): SI = TD, LB = 1.00 Hz. Processing parameters for F1 (y axis): SI = 3·TD, LB = 0.30 Hz.

(4) 2D NMR spectra of $\text{H}_6\text{tren}(\text{dpa})_3$ 

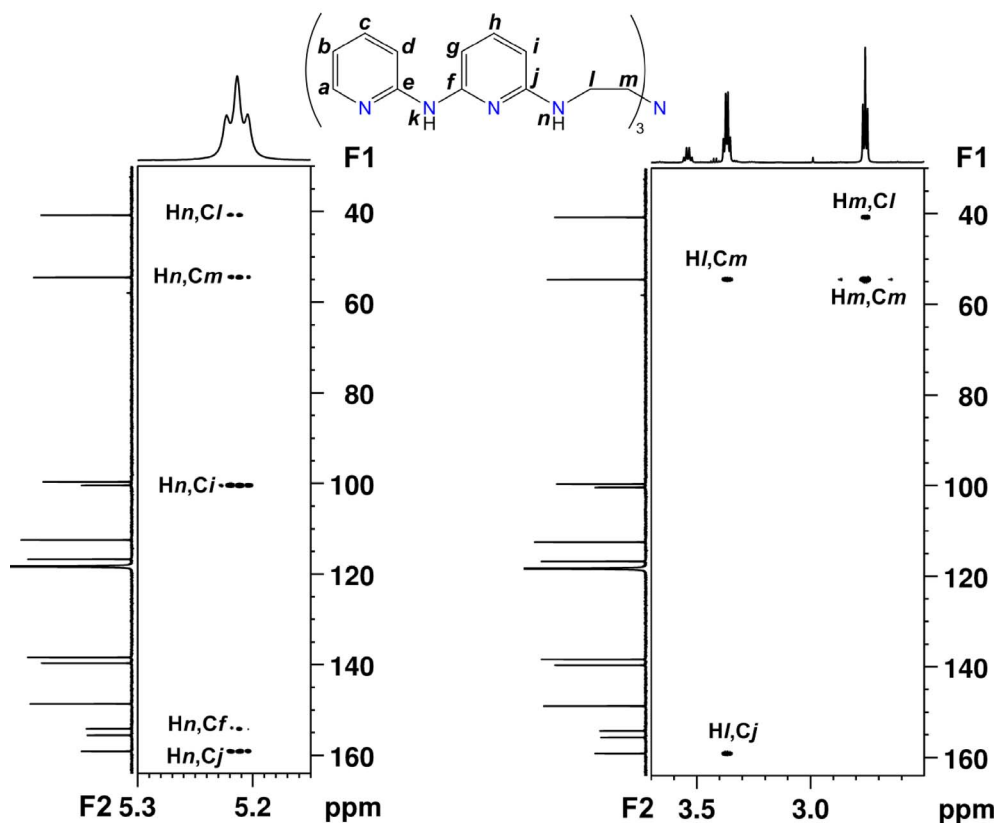
Supplementary Figure S5. Atom-labelled structure and ^1H - ^1H COSY spectrum of $\text{H}_6\text{tren}(\text{dpa})_3$ in CD_3CN for (bottom) $\delta_{\text{H}} = 8.30$ – 5.80 ppm and (top) $\delta_{\text{H}} = 5.40$ – 2.50 ppm (298 K, 600.13 MHz). The labelling of the cross-peaks indicates the ^1H - ^1H coupling (F2, F1). The red line highlights the diagonal peaks. Processing parameters (TopSpin 4.0.6 [4]) for F2 (x axis): SI = TD, LB = 1.00 Hz. Processing parameters for F1 (y axis): SI = 2·TD, LB = 0.30 Hz.



Supplementary Figure S6. Atom-labelled structure and ^1H - ^{13}C HSQC spectrum of $\text{H}_6\text{tren}(\text{dpa})_3$ in CD_3CN for (bottom) $\delta_{\text{H}} = 8.40$ – 5.75 ppm, $\delta_{\text{C}} = 150$ – 98 ppm, and (top) $\delta_{\text{H}} = 4.00$ – 2.20 ppm, $\delta_{\text{C}} = 60$ – 35 ppm (298 K, 600.13 MHz for F2, 150.90 MHz for F1). The labelling of the cross-peaks indicates the ^1H - ^{13}C coupling (F2, F1). Processing parameters (TopSpin 4.0.6 [4]) for F2 (x axis): SI = 2·TD, LB = 1.00 Hz. Processing parameters for F1 (y axis): SI = 3·TD, LB = 0.30 Hz.



Supplementary Figure S7. Atom-labelled structure and ^1H - ^{13}C HMBC spectrum of $\text{H}_6\text{tren}(\text{dpa})_3$ in CD_3CN for $\delta_{\text{H}} = 8.30\text{--}5.80$ ppm and $\delta_{\text{C}} = 164\text{--}96$ ppm (298 K, 600.13 MHz for F2, 150.90 MHz for F1). The labelling of the cross-peaks indicates the ^1H - ^{13}C coupling (F2, F1). Processing parameters (TopSpin 4.0.6 [4]) for F2 (x axis): SI = TD, LB = 1.00 Hz. Processing parameters for F1 (y axis): SI = 3·TD, LB = 0.30 Hz.

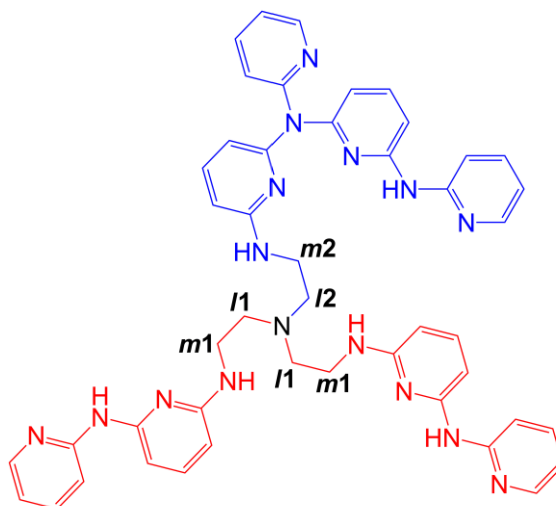


Supplementary Figure S8. Atom-labelled structure and ^1H - ^{13}C HMBC spectrum of $\text{H}_6\text{tren}(\text{dpa})_3$ in CD_3CN for (left) $\delta_{\text{H}} = 5.30$ – 5.15 ppm, $\delta_{\text{C}} = 165$ – 30 ppm, and (right) $\delta_{\text{H}} = 3.70$ – 2.50 ppm, $\delta_{\text{C}} = 165$ – 30 ppm (298 K, 600.13 MHz for F2, 150.90 MHz for F1). The labelling of the cross-peaks indicates the ^1H - ^{13}C coupling (F2, F1). Although a low-pass filter was employed for a better 1J suppression (pulse program: `hmbcetgpl3nd`), the spectrum exhibits the remanence of 1J coupling for site *m*. Processing parameters (TopSpin 4.0.6 [4]) for F2 (*x* axis): SI = TD, LB = 1.00 Hz. Processing parameters for F1 (*y* axis): SI = 3·TD, LB = 0.30 Hz.

(5) Details on ^1H NMR and ESI-MS detection of **SPa**

After the reaction described in Scheme 3b, $\text{H}_6\text{tren}(\text{dpa})_3$ was found admixed in a 1:0.26 MR with side-product **SPa**, which co-eluted during the FC purification. **SPa**, whose structure is represented in Scheme S2, arises from a further nucleophilic substitution reaction between $\text{H}_6\text{tren}(\text{dpa})_3$ and BrHdpa . Due to its low symmetry, **SPa** introduces many extra peaks in the ^1H NMR spectrum of the chromatographed mixture (Figure S9). In particular, the spectrum shows a new set of eleven signals, indicated in Figure S9 by red circles, which are similar to those found in $\text{H}_6\text{tren}(\text{dpa})_3$ in terms of hyperfine splitting and relative areas, and arise from the two “unmodified” branches of **SPa** (also colored in red in Scheme S2). The chemical shifts of seven of these signals are significantly different from those of $\text{H}_6\text{tren}(\text{dpa})_3$, and they were easily identified in the ^1H NMR spectrum. On the other hand, the remaining four are hidden by the corresponding peaks of $\text{H}_6\text{tren}(\text{dpa})_3$ (*Ha*, *Hb*, *Hc* and *Hh*). However, their presence was clearly demonstrated by 2D NMR spectroscopy analysis (^1H - ^1H COSY). The branch of **SPa** undergoing the “extra” arylation (colored in blue in Scheme S2) should be responsible for a set of eighteen peaks. In Figure S9 seventeen over the eighteen expected peaks were detected (blue circles), and their presence was confirmed by ^1H - ^1H COSY spectroscopy.

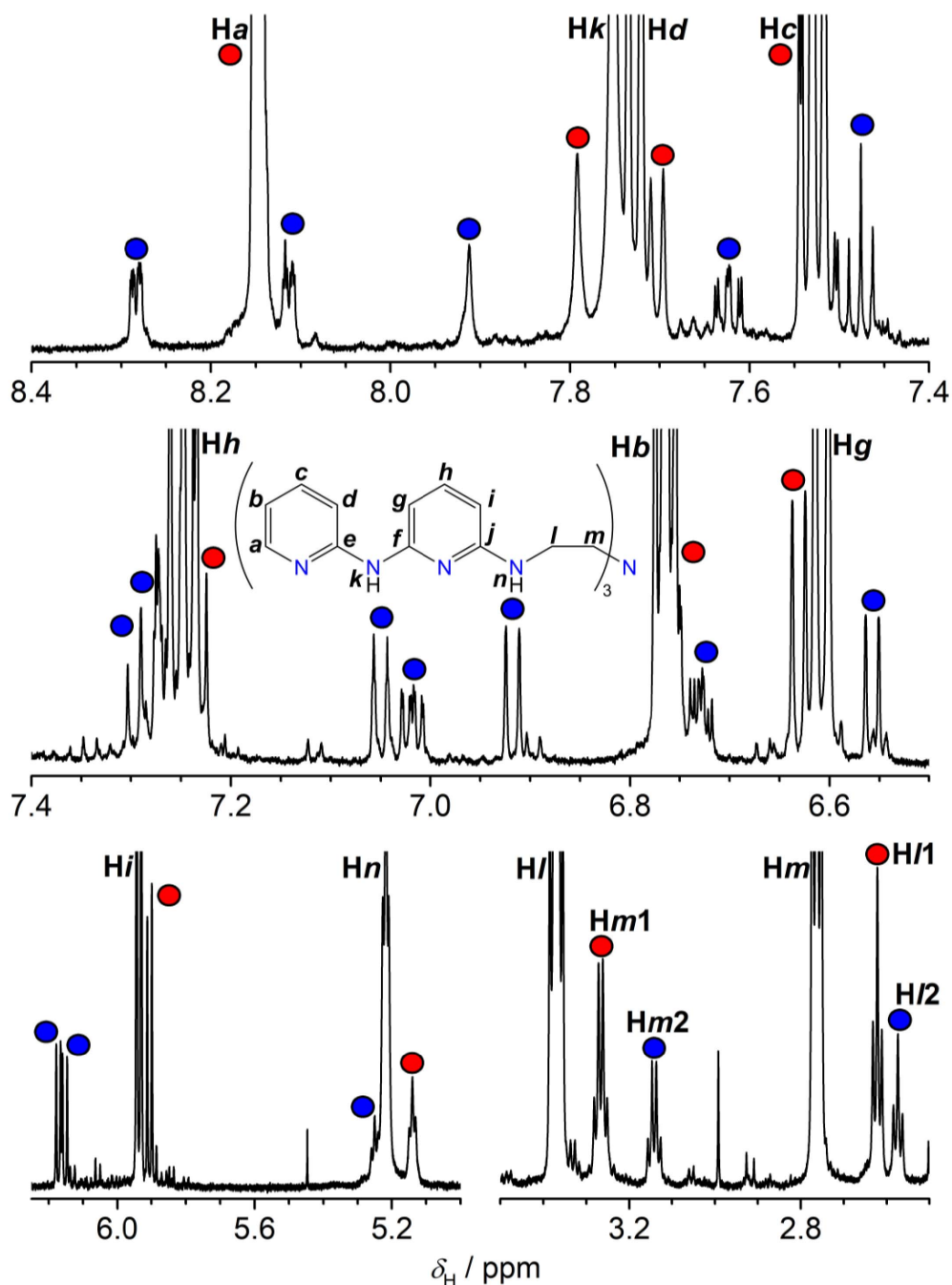
Unfortunately, due to the complexity of the spectra, the remaining peak was not spotted. This hidden signal is part of a set of sixteen peaks with identical areas, which arise from fourteen pyridyl and two amino protons. The fifteen spotted peaks are in the correct 1:2 integrated intensity ratio with the seven aromatic CH and two NH signals marked with red circles. The two remaining peaks correspond to the aliphatic H atoms and their integrated intensity is in fact in the expected 2:1 and 1:2 ratio with respect to the fifteen aromatic peaks (sixteen expected) and the aliphatic signals marked with red circles, respectively. All the aliphatic protons of **SPa** were assigned with good confidence, following the same scheme adopted for $\text{H}_6\text{tren}(\text{dpa})_3$, while the assignment of the aro-



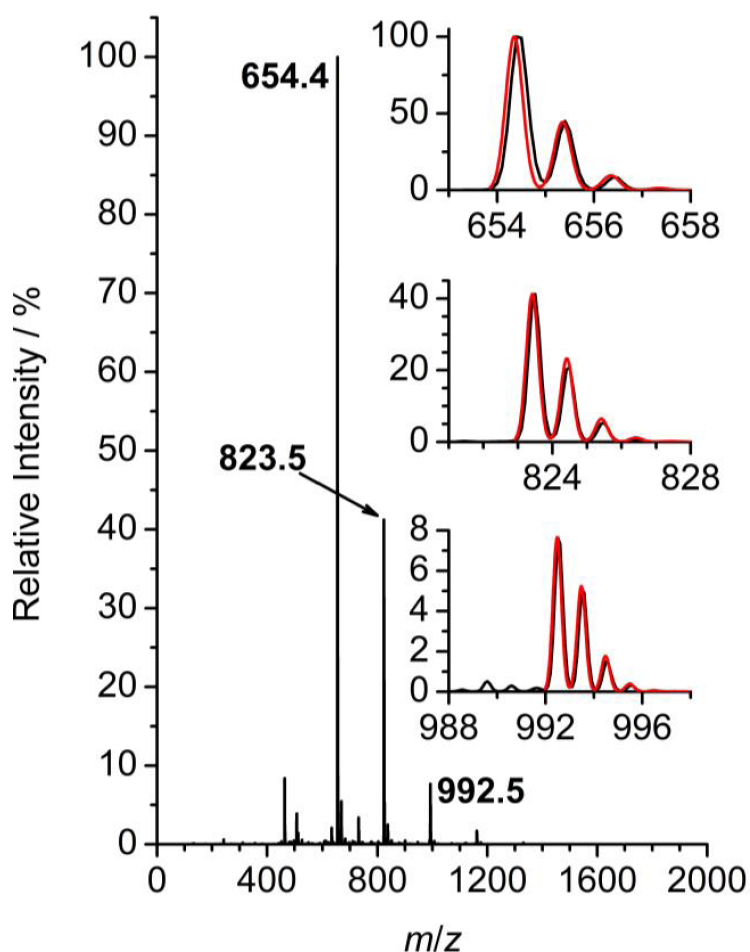
Scheme S2. Structure of **SPa** and assignment of the aliphatic H atoms. The red (blue) region of **SPa** generates the ^1H NMR signals marked with red (blue) circles in Figure S9 (below).

matic resonances is not straightforward due to many overlapping peaks. Therefore, the high-field region is the most informative one to understand which species are present in a sample, and to estimate their MR.

These NMR spectroscopy findings are in accordance with ESI-MS spectrometry measurements. The ESI-MS spectrum of the chromatographed mixture containing $\text{H}_6\text{tren}(\text{dpa})_3$ and **SPa** is shown in Figure S10. Two intense and well resolved peaks are present at $m/z = 654.4$ (100%) and 823.5 (41%), which are well simulated by the ionic species $[\text{H}_6\text{tren}(\text{dpa})_3 + \text{H}]^+$ and $[\text{H}_5\text{tren}(\text{dpa})_4 + \text{H}]^+$ (**SPa** + H) $^+$. The intensity ratio between these two signals ($\sim 1:0.4$) is lower than the MR observed with ^1H NMR spectroscopy (1:0.26). A minority peak well simulated by $[\text{H}_4\text{tren}(\text{dpa})_5 + \text{H}]^+$ is clearly visible at $m/z = 992.5$ (8%), even if the species $\text{H}_4\text{tren}(\text{dpa})_5$ was not detected in the ^1H NMR spectrum of the same mixture. Since NMR is a better quantitative technique than ESI-MS spectrometry, these findings suggest that the signals related to $[\text{H}_5\text{tren}(\text{dpa})_4 + \text{H}]^+$ and $[\text{H}_4\text{tren}(\text{dpa})_5 + \text{H}]^+$ are enhanced here, probably due to ionic collision phenomena.



Supplementary Figure S9. ^1H NMR spectrum of the chromatographed mixture containing $\text{H}_6\text{tren}(\text{dpa})_3$ and SPa , in CD_3CN , for (bottom right) $\delta_{\text{H}} = 3.50\text{--}2.50$ ppm, (bottom left) $\delta_{\text{H}} = 6.25\text{--}5.00$ ppm, (middle) $\delta_{\text{H}} = 7.40\text{--}6.50$ ppm, and (top) $\delta_{\text{H}} = 8.40\text{--}7.40$ ppm (298 K, 600.13 MHz). The labels show the assignments for $\text{H}_6\text{tren}(\text{dpa})_3$ (all protons) and SPa (aliphatic protons only). The red and blue circles indicate the signals originating from the different branches of SPa , which are marked with the same colors in Scheme S2. Processing parameters (TopSpin 4.0.6 [4]): SI = TD, LB = 0.30 Hz.



Supplementary Figure S10. ESI-MS spectrum of the chromatographed mixture containing $\text{H}_6\text{tren}(\text{dpa})_3$ and **SPa** (direct infusion, CH_2Cl_2 , positive ion mode). The insets show the experimental (black line) and simulated (red line) isotopic patterns of the peaks at $m/z = 654.4$ ($[\text{H}_6\text{tren}(\text{dpa})_3 + \text{H}]^+$), 823.5 ($[\text{H}_5\text{tren}(\text{dpa})_4 + \text{H}]^+$) and 992.5 ($[\text{H}_4\text{tren}(\text{dpa})_5 + \text{H}]^+$). The peak at $m/z = 464.3$ (8%) was not assigned.

References

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