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Supporting Information

Inhibition of [FeFe]-Hydrogenase by Formaldehyde: Proposed Mechanism and Reactivity of FeFe Alkyl Complexes

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Figure S1. IR spectra of $[Fe_2(SCH_2)_2NCH_2(CO)_4(PMe_3)_2]BF_4$ ([2]BF₄) and $[Fe_2(SCH_2)_2NCH_2(CO)_4(PMe_3)_2]BAr^{F_4}$ ([2]BAr^{F_4}) in CH₂Cl₂ solution.



Figure S2. HR-MS (ESI) spectrum of $[Fe_2[(SCH_2)_2NCH_2](CO)_4(PMe_3)_2]BF_4$ ([2]BF4) in CH₂Cl₂ solution.



Figure S3. ESI-MS spectrum of ¹³C labeled $[Fe_2[(SCH_2)_2NCH_2](CO)_4(PMe_3)_2]BF_4$ ([2]BF₄) in CH₂Cl₂ solution.



Figure S4. ¹H NMR spectrum of $[Fe_2[(SCH_2)_2NCH_2](CO)_4(PMe_3)_2]BF_4$ ([2]BF₄) in CD₂Cl₂ at room temperature.



Figure S5. ³¹P NMR spectrum of $[Fe_2[(SCH_2)_2NCH_2](CO)_4(PMe_3)_2]BF_4$ ([2]BF₄) in CD₂Cl₂ at room temperature.



Figure S6. ¹H NMR spectrum of $[Fe_2[(SCH_2)_2NCH_2](CO)_4(PMe_3)_2]BAr^F_4$ ([**2**]BAr^F₄) in CD_2CI_2 at room temperature.



Figure S7. ³¹P NMR spectrum of $[Fe_2[(SCH_2)_2NCH_2](CO)_4(PMe_3)_2]BAr^F_4$ ([**2**]BAr^F₄) in CD₂Cl₂ at room temperature.

Three isomers were detected and assigned as follows:

δ 22.59 (d, J_{PP} = 7.3 Hz), 9.45 (d, J_{PP} = 7.4 Hz), *trans*-dibasal; 34.46 (d, J_{PP} = 7.7 Hz), 24.10 (d, J_{PP} = 7.6 Hz), *cis*-dibasal; 18.41 (s), 9.21 (s), apical-basal.



Figure S8. ¹³C NMR spectrum of $[Fe_2[(SCH_2)_2NCH_2](CO)_4(PMe_3)_2]BAr^F_4$ ([**2**]BAr^F₄) in CD₂Cl₂ at room temperature.





Figure S10. 2D-HSQC (¹H-¹³C) NMR spectrum (CH₂ region) of $[Fe_2[(SCH_2)_2NCH_2](CO)_4(PMe_3)_2]BAr^{F_4}$ ([**2**]BAr^{F_4}) in CD₂Cl₂ at room temperature.



Figure S11. ¹³C NMR spectrum of ¹³C labeled [Fe₂[(SCH₂)₂NCH₂](CO)₄(PMe₃)₂]BAr^F₄ ([**2**]BAr^F₄) in CD₂Cl₂ at room temperature.



Figure S12. ¹H NMR spectrum (C*H*₂ region) of ¹³C labeled (top) and unlabeled (bottom) [Fe₂[(SCH₂)₂NCH₂](CO)₄(PMe₃)₂]BAr^F₄ ([**2**]BAr^F₄) in CD₂Cl₂ at room temperature.

Exchange Process in [2]BAr^{F4} Determined by Spin Saturation Transfer

1D Spin-Saturation Transfer (SST) experiment (both ³¹P and ¹H NMR) was used to monitor the slow exchange process in [**2**]BAr^F₄ in CD₂Cl₂ at 298 K. One signal is selectively irradiated, and the signal intensity of the other peak connected with the irradiated signal via chemical exchange is monitored.

For exchange process of

$$A \xrightarrow{k_{A}} B$$

Selectively irradiation of peak A, the intensity of peak B changes following the equation below, and the exchange rate can be extracted:

$$\frac{I(t)}{I(0)} = \frac{\tau'}{\tau} \exp\left(-\frac{t}{\tau'}\right) + \frac{\tau'}{T_1}$$
$$\tau' = \frac{1}{\frac{1}{T_1} + \frac{1}{\tau}}$$

Where t is the irradiation time applied to peak A, T₁ is the longitudinal (spin-lattice) relaxation time of peak B (measured by inversion-recovery method here), τ is the life time of nuclei stays at site B, and the exchange rate $k_{\rm B} = 1/\tau$.



Figure S13. Stacked ¹H NMR spectra of [**2**]BAr^F₄ (29 mM) with different relaxation time in CD₂Cl₂ at 298 K (top); Plot of peak intensity at δ 5.52 (red), 4.79 (blue), 4.17 (green) and 3.84 (orange) *vs* relaxation time (bottom).

Fitted with three parameter exponential equation: $I_t = A + B^* \exp(-t/T_1)$.

Results:

δ 5.52, $T_1 = 1.29$ s; δ 4.79, $T_1 = 1.31$ s; δ 4.17, $T_1 = 1.17$ s; δ 3.84, $T_1 = 1.17$ s.



Figure S14. ¹H NMR spectra of [**2**]BAr^F₄ (29 mM) at 298 K in CD₂Cl₂ showing the change in intensity of the proton signal at δ 5.52, 4.17, 3.84 and 3.25 upon irradiation at δ 4.79 (top), and the plot of intensity *vs* irradiation time at δ 4.79 (bottom).



Figure S15. ¹H NMR spectra of [**2**]BAr^F₄ (29 mM) at 298 K in CD₂Cl₂ showing the change in intensity of the proton signal at δ 5.52, 4.79, 4.17 and 3.25 upon irradiation at δ 3.84 (top), and the plot of intensity vs irradiation time at δ 3.84 (bottom).



Figure S16. Spin saturation transfer spectra of [**2**]BAr^F₄ at 298 K in CD₂Cl₂, showing the decline in intensity of the phosphine signal at δ 9.45, 24.10, 18.41 and 9.21 upon irradiation at δ 22.59.



Figure S17. ¹H NMR spectrum of $[Fe_2[(SCH_2)_2NH_2](CO)_4(PMe_3)_2]BF_4$ ([1H]BF₄) in CD₂Cl₂ at room temperature.



Figure S18. ³¹P NMR spectrum of $[Fe_2[(SCH_2)_2NH_2](CO)_4(PMe_3)_2]BF_4$ ([1H]BF₄) in CD₂Cl₂ at room temperature.

-26.26



Figure S19. IR spectra of $Fe_2[(SCH_2)_2NH](CO)_4(PMe_3)_2$ (1) [$Fe_2[(SCH_2)_2NH_2](CO)_4(PMe_3)_2$]BF₄ ([1H]BF₄) in CH₂Cl₂ solution.

and



Figure S20. IR spectrum of $[HFe_2[(SCH_2)_2NH](CO)_4(PMe_3)_2]BAr^{F_4}$ ([H1]BAr^{F_4}) in CH₂Cl₂ solution.



Figure S21. ¹H NMR spectrum of $[HFe_2[(SCH_2)_2NH](CO)_4(PMe_3)_2]BAr^{F_4}$ ([H1]BAr^{F_4}) in CD₂Cl₂ at room temperature.



Figure S22. ³¹P NMR spectrum of $[HFe_2[(SCH_2)_2NH](CO)_4(PMe_3)_2]BAr^{F_4}$ ($[H1]BAr^{F_4}$) in CD_2Cl_2 at room temperature.



Figure S23. IR spectrum for reaction of $[HFe_2[(SCH_2)_2NH](CO)_4(PMe_3)_2]BAr^{F_4}$ ([H1]BAr^{F_4}) with 3 equiv paraformaldehyde in CH₂Cl₂ solution.

The result indicates the reaction of hydride tautomer $[H1]^+$ with paraformaldehyde does not give $[2]^+$, which highlight the role of the ammonium center in the conversion of **1** to $[2]^+$.



Figure S24. IR spectrum of $Fe_2[(SCH_2)_2NH](CO)_4(PMe_3)_2$ (1) and 1 + CH₂O for 24 h in CH₂Cl₂ solution.



Figure S25. ¹³C NMR spectrum for the reaction of $[Fe_2(SCH_2)_2NH(CO)_4(PMe_3)_2]$ with 0.25 equiv ¹³CH₂O in CD₂Cl₂ at room temperature.

Assignments:

δ 80.11 (enriched with ¹³C) and 52.56 ([Fe₂[(SCH₂)₂N¹³CH₂OH](CO)₄(PMe₃)₂]); δ 45.91 ([Fe₂[(SCH₂)₂NH](CO)₄(PMe₃)₂]).

In order to exam whether there is exchange during the formation of N-CH₂-OH, the reaction of $[Fe_2[(SCH_2)_2NH](CO)_4(PMe_3)_2]$ with 0.25 equiv ¹³CH₂O in CD₂Cl₂ was checked at room temperature. According to the ¹³C NMR determined after 72 h, two new peaks appeared at δ 80.11 (enriched with ¹³C) and δ 52.56, which could be assigned to the 1 : 1 adduct of $[Fe_2[(SCH_2)_2N^{13}CH_2OH](CO)_4(PMe_3)_2]$. However, the peak at δ 45.91 corresponding to the SCH₂ of Fe₂[(SCH₂)₂NH](CO)₄(PMe₃)₂ has not been enriched, indicating no exchange during this process.



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Figure S26. ¹³C NMR spectrum for the reaction of $[Fe_2[(SCH_2)_2NH](CO)_4(PMe_3)_2]$ with 1 equiv CH₂O in CD₂Cl₂ at room temperature.

Assignments: δ 80.15 and 52.60 ([Fe₂[(SCH₂)₂NCH₂OH](CO)₄(PMe₃)₂]); δ 90.23, 89.22, 87.34, 86.94, 86.50, 82.67 for -(CH₂)_n-OH and 52.30, 51.68 for SCH₂ ([Fe₂[(SCH₂)₂N(CH₂)_nOH](CO)₄(PMe₃)₂]); δ 45.91 ([Fe₂[(SCH₂)₂NH](CO)₄(PMe₃)₂]).



Figure S27. IR spectra for reaction of $[Fe_2(pdt)(CO)_4(PMe_3)_2] + CH_2O + HBF_4$ in CH₂Cl₂ solution.

According to the IR spectra, no reaction is evident when $[Fe_2(pdt)(CO)_4(PMe_3)_2]$ was treated with CH₂O. In the presence of 1 equiv HBF₄, bridging hydride $[HFe_2(pdt)(CO)_4(PMe_3)_2]BF_4$ was formed as the protonation product.



Figure S28. IR spectra for reaction of $Fe_2[(SCH_2)_2NH](CO)_4(PMe_3)_2$ (1) + PhCHO + HBF₄·Et₂O in CH₂Cl₂ solution. The band at 1702 cm⁻¹ is assigned to C=O vibration of PhCHO.

According to the IR spectra, no reaction is evident when **1** was treated with PhCHO. In the presence of HBF₄, ([Fe₂[(SCH₂)₂NH₂](CO)₄(PMe₃)₂]BF₄) ([**1**H]BF₄) was formed as the protonation product.



Figure S29. ¹³C NMR spectrum (CH₂ region) for the reaction of $[Fe_2[(SCH_2)_2NH](CO)_4(PMe_3)_2]$ (47 mM) + ¹³CH₂O + HBAr^F₄ (1 : 1 : 1) over time in CD₂Cl₂ at 296 K (top); Plot of peak intensity *vs* time, showing the decline in intensity of the signal at δ 76.66, 75.77, 75.49 (brown), and the increase in intensity of the signal at δ 59.03, 58.46, 58.16, 57.09 (blue) (bottom). The exchange process reaches equilibrium after ~90 minutes.

$$A \xrightarrow[k_{-1}]{k_1} B$$

 k_1 is the rate coefficient for the reaction that consumes A; k_{-1} is the rate coefficient for the backwards reaction, then

$$\frac{k_1}{k_{-1}} = \frac{[\mathsf{B}]_{\mathsf{e}}}{[\mathsf{A}]_{\mathsf{e}}}$$

 $[A]_e$ and $[B]_e$ are the concentrations of A and B at equilibrium, respectively. $[A]_0$ is the initial concentration of reactant A, $[A]_t$ is the concentration of A at time t, then

$$\ln\left(\frac{I_0 - I_e}{I_t - I_e}\right) = \ln\left(\frac{[A]_0 - [A]_e}{[A]_t - [A]_e}\right) = (k_1 + k_{-1})t$$



Figure S30. Plot of $\ln[(I_0 - I_e)/(I_t - I_e)]$ vs reaction time ([**2**]⁺₀ = 47 mM). *Fitting equation*: $\ln[(I_0 - I_e)/(I_t - I_e)] = (k_1 + k_{-1})t$. *Results:* $k_1 + k_{-1} = 8.31 \times 10^{-4} \pm 2.01 \times 10^{-5} s^{-1}$. The linear plot indicates the kinetics of exchange are first order in the concentration of [**2**]⁺, which points to an intramolecular process.



Figure S31. IR spectrum of crude Et₄N[Fe₂[(SCH₂)₂NCH₂CN](CN)(CO)₅] (Et₄N[**4**]) in CH₃CN solution.



Figure S32. ESI-MS spectrum of $Et_4N[Fe_2[(SCH_2)_2NCH_2CN](CN)(CO)_5]$ ($Et_4N[4]$) in CH₃CN solution.



Figure S33. ESI-MS spectrum of ¹³C labeled Et₄N[Fe₂[(SCH₂)₂NCH₂CN](CN)(CO)₅] (Et₄N[**4**]) in CH₃CN solution.



Figure S34. ¹H NMR spectrum of Et₄N[Fe₂[(SCH₂)₂NCH₂CN](CN)(CO)₄PPh₃] (Et₄N[**5**]) in CD₂Cl₂ at room temperature.



Figure S35. ³¹P NMR spectrum of Et₄N[Fe₂[(SCH₂)₂NCH₂CN](CN)(CO)₄PPh₃] (Et₄N[**5**]) in CD₂Cl₂ at room temperature.



Figure S36. ¹³C NMR spectrum of Et₄N[Fe₂[(SCH₂)₂NCH₂CN](CN)(CO)₄(PPh₃)] (Et₄N[**5**]) in CD₂Cl₂ at room temperature.

Assignments: δ 66.04, 15.48 (diethyl ether)



Figure S37. IR spectrum of Et₄N[Fe₂[(SCH₂)₂NCH₂CN](CN)(CO)₄PPh₃] (Et₄N[**5**]) in CH₂Cl₂ solution.



Figure S38. ESI-MS spectrum of Et₄N[Fe₂[(SCH₂)₂NCH₂CN](CN)(CO)₄PPh₃] (Et₄N[**5**]) in CH₂Cl₂ solution.



[HPPh₃]BF₄ in CH₃CN solution.



Figure S40. IR spectra of $(Et_4N)[HFe_2[(SCH_2)_2NH](CN)_2(CO)_4]$ ($(Et_4N)[H3]$) + 2 equiv CH₂O in CH₃CN solution.



Figure S41. IR spectra for the reaction of $(Et_4N)_2[Fe_2(pdt)(CN)_2(CO)_4]$ with CH₂O and [HPPh₃]BF₄ in CH₃CN solution.



Figure S42. ¹H NMR spectrum of [Fe₂[(SCH₂)₂NCH₂PPh₃](CO)₄(PMe₃)₂]BF₄ ([**6**]BF₄) in CD₂Cl₂ at room temperature.

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Figure S43. ³¹P NMR spectrum of [Fe₂[(SCH₂)₂NCH₂PPh₃](CO)₄(PMe₃)₂]BF₄ ([**6**]BF₄) in CD₂Cl₂ at room temperature.



Figure S44. IR spectra of $[Fe_2[(SCH_2)_2NCH_2PPh_3](CO)_4(PMe_3)_2]BF_4$ ([6]BF₄) and [6]BAr^F₄ in CH₂Cl₂ solution.



Figure S45. ESI-MS spectrum of $[Fe_2[(SCH_2)_2NCH_2PPh_3](CO)_4(PMe_3)_2]BF_4$ ([6]BF₄) in CH_2Cl_2 solution.

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 $[Fe_2[(SCH_2)_2NCH_2PPh_3](CO)_4(PMe_3)_2]BArF_4$ ([6]BArF_4) in CD₂Cl₂ at room temperature.



FigureS47. ${}^{31}P$ NMRspectrumof ${}^{13}C$ labeled[Fe2[(SCH2)2NCH2PPh3](CO)4(PMe3)2]BArF4 ([6]BArF4) in CD2Cl2 at room temperature.



Figure S48. ¹H NMR spectrum of Fe₂[(SCH₂)₂NMe](CO)₄(PMe₃)₂ in CD₂Cl₂ at room temperature.



Figure S49. ³¹P NMR spectrum of Fe₂[(SCH₂)₂NMe](CO)₄(PMe₃)₂ in CD₂Cl₂ at room temperature.





Figure S51. ESI-MS spectrum of Fe₂[(SCH₂)₂NMe](CO)₄(PMe₃)₂ in CH₂Cl₂ solution.

S54

Identification code	ed79ks ([2]BAr ^F ₄)	ed04Ls ([1 H]BF ₄)
Empirical formula	C45 H36 B F24 Fe2 N O4 P2	C12.67 H25.33 B Cl1.33 F4
	S2	Fe2 N O4 P2 S2
Formula weight	1359.32	627.51
Temperature	100(2) K	100(2) K
Wavelength	0.71073 Å	0.71073 Å
Crystal system	Triclinic	Triclinic
Space group	P-1	P-1
Unit cell dimensions	a = 13.5275(3) Å	a = 14.0486(2) Å
	b = 14.3475(4) Å	b = 16.9932(3) Å
	c = 16.5656(4) Å	c = 17.3960(3) Å
Volume	2661.82(11) Å ³	3759.01(11) Å ³
Z	2	6
Density (calculated)	1.696 Mg/m ³	1.663 Mg/m ³
Absorption coefficient	0.811 mm ⁻¹	1.643 mm ⁻¹
F(000)	1360	1908
Crystal size	0.228 x 0.184 x 0.064 mm ³	0.191 x 0.153 x 0.107 mm ³
Theta range for data collection	2.290 to 28.301°.	2.149 to 28.307°.
Index ranges	-18<=h<=18, -19<=k<=19	-18<=h<=18, -22<=k<=22
	-22<=l<=22	-23<=l<=23
Reflections collected	190539	222825
Independent reflections	13211 [R(int) = 0.0312]	18706 [R(int) = 0.0328]
Completeness to theta = 25.242°	99.9 %	99.9 %
Absorption correction	Semi-empirical from	Semi-empirical from
	equivalents	equivalents
Max. and min. transmission	0.7457 and 0.7147	0.7457 and 0.7016
Refinement method	Full-matrix least-squares on F ²	Full-matrix least-squares on F ²
Data / restraints / parameters	13211 / 52 / 743	18706 / 548 / 995
Goodness-of-fit on F ²	1.023	1.050
Final R indices [I>2sigma(I)]	R1 = 0.0348, wR2 = 0.0840	R1 = 0.0213, wR2 = 0.0512
R indices (all data)	R1 = 0.0369, wR2 = 0.0856	R1 = 0.0241, wR2 = 0.0527
Extinction coefficient	n/a	n/a
Largest diff. peak and hole	1.246 and -0.990 e.Å ⁻³	0.970 and -0.765 e.Å ⁻³

Table S1. Crystal Data and Structure Refinements.

Table el Cijetal Bata alla		
Identification code	ed82Ls (Et4N[5])	ed15Ls ([6]BF ₄)
Empirical formula	C35 H41 Fe2 N4 O4 P S2	C34.79 H48.44 B Cl0.14 F4
		Fe2 N O4.93 P3 S2
Formula weight	788.51	920.04
Temperature	120(2) K	100(2) K
Wavelength	0.71073 Å	0.71073 Å
Crystal system	Monoclinic	Monoclinic
Space group	P2 ₁ /c	P2 ₁ /c
Unit cell dimensions	a = 16.8201(7) Å	a = 13.5415(3) Å
	b = 17.3607(7) Å	b = 17.0231(4) Å
	c = 13.5621(6) Å	c = 18.4766(4) Å
Volume	3719 3(3) Å3	4173 68(16) Å ³
7	1	4170.00(10)71
	+	7
Density (calculated)	1.408 Mg/m ³	1.464 Mg/m ³
Absorption coefficient	0.978 mm ⁻¹	0.977 mm ⁻¹
F(000)	1640	1904
Crystal size	0.468 x 0.305 x 0.054 mm ³	0.223 x 0.214 x 0.079 mm ³
Theta range for data collection	1.983 to 26.477°.	2.089 to 27.500°.
Index ranges	-21<=h<=21, -21<=k<=21	-17<=h<=17, -22<=k<=22
	-16<=l<=16	-23<=l<=23
Reflections collected	73436	60739
Independent reflections	7636 [R(int) = 0.0605]	9573 [R(int) = 0.0352]
Completeness to theta = 25.242°	99.9 %	99.9 %
Absorption correction	Semi-empirical from	Semi-empirical from
	equivalents	equivalents
Max. and min. transmission	0.7454 and 0.5933	0.7456 and 0.7005
Refinement method	Full-matrix least-squares on F ²	Full-matrix least-squares on F ²
Data / restraints / parameters	7636 / 332 / 542	9573 / 90 / 511
Goodness-of-fit on F ²	1.186	1.103
Final R indices [I>2sigma(I)]	R1 = 0.0600, wR2 = 0.1300	R1 = 0.0317, wR2 = 0.0718
R indices (all data)	R1 = 0.0690, wR2 = 0.1341	R1 = 0.0386, wR2 = 0.0750
Extinction coefficient	n/a	n/a
Largest diff. peak and hole	0.859 and -0.663 e.Å ⁻³	0.504 and -0.344 e.Å ⁻³

Table S2. Crystal Data and Structure Refinements.

Crystallography

Ed04Ls ([1H]BF₄) CCDC: 2104104

Intensity data were collected on a Bruker D8 Venture kappa diffractometer equipped with a Photon-II CPAD detector. An Iµs microfocus Mo source (λ = 0.71073 Å) coupled with a multi-layer mirror monochromator provided the incident beam. The sample was mounted on a 0.3 mm nylon loop with the minimal amount of Paratone-N oil. Data was collected as a series of φ and/or ω scans. Data was collected at 100 K using a cold stream of N_{2(g)}. The collection, cell refinement, and integration of intensity data was carried out with the APEX3 software.¹ A multi-scan absorption correction was performed with SADABS². The structure was phased with intrinsic phasing methods using SHELXT³ and refined with the full-matrix least-squares program SHELXL.⁴

A structural model consisting of three target molecules, three BF₄ anions, and two dichloromethane solvent molecules in the asymmetric unit was developed.

Two of the three BF₄ anions were modeled as disordered over two orientations. Within each anion, all 1,2 and 1,3 distances were restrained to be similar (esd 0.01, 0.02 Å). Rigid-bond restraints (esd 0.004) were imposed on displacement parameters for all disordered sites and similar displacement amplitudes (esd 0.005) were imposed on disordered sites overlapping by less than the sum of van der Waals radii. The site occupancy ratios were allowed to freely refine.

Both dichloromethane solvent molecules were modeled as disordered over two orientations. Within each solvent molecule, all 1,2 and 1,3 distances were restrained to be similar (esd 0.01, 0.02 Å). Similar displacement amplitudes (esd 0.005) were imposed on disordered sites overlapping by less than the sum of van der Waals radii. The site occupancy ratios were allowed to freely refine.

H atom treatment - Methyl H atom positions, R-CH₃, were optimized by rotation about R-C bonds with idealized C-H, R--H and H--H distances. All of the amine H atom positions were located in the difference map; their positions were allowed to freely refine. At convergence, all amine H atoms were in good H- bonding geometries. Remaining H atoms were included as riding idealized contributors. Methyl and amine H atom U's were assigned as 1.5 times U_{eq} of the carrier atom; remaining H atom U's were assigned as 1.2 times carrier U_{eq}.

The -3 5 0 reflection was omitted from the final refinement due to being partially obscured by the beam stop support in some orientations.

Ed15Ls ([6]BF4) CCDC: 2104105

Intensity data were collected on a Bruker D8 Venture kappa diffractometer equipped with a Photon-II CPAD detector. An Iµs microfocus Mo source (λ = 0.71073 Å) coupled with a multi-layer mirror monochromator provided the incident beam. The sample was mounted on a 0.3 mm nylon loop with the minimal amount of Paratone-N oil. Data was collected as a series of φ and/or ω scans. Data was collected at 100 K using a cold stream of N_{2(g)}. The collection, cell refinement, and integration of intensity data was carried out with the APEX3 software.¹ A multi-scan absorption correction was performed with SADABS². The structure was phased with intrinsic phasing methods using SHELXT³ and refined with the full-matrix least-squares program SHELXL.⁴

A structural model consisting of the target molecule, a BF₄ counter ion, and a disordered solvent molecule position in the asymmetric unit was developed.

The solvent molecule position was refined as occupational disorder of diethyl ether and dichloromethane. The dichloromethane occupancy was only approximately seven percent; to maintain a reasonable geometry with such a low occupancy it was refined as an idealized, rigid fragment.⁵ Similar displacement amplitudes (esd 0.01) were imposed on disordered sites overlapping by less than the sum of van der Waals radii. The site occupancy ratio was allowed to freely refine.

H atom treatment - Methyl H atom positions, R-CH₃, were optimized by rotation about R-C bonds with idealized C-H, R--H and H--H distances. Remaining H atoms were included as riding idealized contributors. Methyl H atom U's were assigned as 1.5 times U_{eq} of the carrier atom; remaining H atom U's were assigned as 1.2 times carrier U_{eq} .

The 1 1 0, 1 0 0, and 0 1 1 reflections were omitted from the final refinement due to being partially obscured by the beam stop in some orientations. The 6 6 12 reflection was omitted from the final refinement due to being partially obscured by the Cu beam stop in some orientations. The -8 1 3 and -9 1 2 reflections both showed large F_0^2 vs. F_c^2 deviations with F_0^2 being larger than F_c^2 . Inspection of individual frame images revealed that in several instances there was a hot pixel on the detector in close enough proximity to the reflection that it may have been included in the integration box. These reflections were omitted from the final refinement.

Ed79ks ([2]BAr^F₄) CCDC: 2104106

Intensity data were collected on a Bruker D8 Venture kappa diffractometer equipped with a Photon-II CPAD detector. An Iµs microfocus Mo source (λ = 0.71073 Å) coupled with a multi-layer mirror monochromator provided the incident beam. The sample was mounted on a 0.3 mm nylon loop with the minimal amount of Paratone-N oil. Data was collected as a series of φ and/or ω scans. Data was collected at 100 K using a cold stream of N_{2(g)}. The collection, cell refinement, and integration of intensity data was carried out with the APEX3 software.¹ A multi-scan absorption correction was performed with SADABS². The structure was phased with intrinsic phasing methods using SHELXT³ and refined with the full-matrix least-squares program SHELXL.⁴

A structural model consisting of the target molecule plus one BArF₂₄ anion in the asymmetric unit was developed.

Two of the CF₃ groups on the BArF₂₄ anion were modeled as disordered; the C20 group over two orientations and the C29 group over three orientations. Similarity restraints (esd 0.01 Å) were imposed on all disordered C---F bond distances. Within each disordered group, the fluorine displacement parameters were constrained to be the same. The C20 site occupancy ratio was allowed to freely refine. The site occupancies for the three C29 orientations were allowed to freely refine with in the restraint that the total occupancy was 1.000(1).

H atom treatment - Methyl H atom positions, R-CH₃, were optimized by rotation about R-C bonds with idealized C-H, R--H and H--H distances. Remaining H atoms were included as riding idealized contributors. Methyl H atom U's were assigned as 1.5 times U_{eq} of the carrier atom; remaining H atom U's were assigned as 1.2 times carrier U_{eq} .

The -1 1 1 reflection was omitted from the final refinement due to being partially obscured by the beam stop in some orientations.

Ed82Ls (Et4N[5]) CCDC: 2104107

Intensity data were collected on a Bruker D8 Venture kappa diffractometer equipped with a Photon-II CPAD detector. An Iµs microfocus Mo source (λ = 0.71073 Å) coupled with a multi-layer mirror monochromator provided the incident beam. The sample was mounted on a 0.3 mm nylon loop with the minimal amount of Paratone-N oil. Data was collected as a series of φ and/or ω scans. Data was collected at 120 K using a cold stream of N_{2(g)}. The collection, cell refinement, and integration of intensity data was carried out with the APEX3 software.¹ A multi-scan absorption correction was performed with SADABS². The structure was phased with intrinsic phasing methods using SHELXT³ and refined with the full-matrix least-squares program SHELXL.⁴

A structural model consisting of the target molecule plus one tetraethylammonium cation in the asymmetric unit was developed.

The nitrile substituent on the adt bridge was modeled as disordered over two orientations. Similarity restraints (esd 0.01 Å) were imposed on all chemically equivalent bond distances. To maintain a more linear geometry about the nitrile group for the minor orientation, similarity restraints (esd 0.02 Å) were imposed on the distances between the nitrile nitrogen atoms and the CH₂ H atoms. Similar displacement amplitudes were imposed on disordered sites overlapping by less than the sum of van der Waals radii. The site occupancy ratio was allowed to freely refine.

The tetraethylammonium cation was modeled as disordered over two orientations. All 1,2 and 1,3 distances of the cation were restrained to be similar (esd 0.01, 0.04 Å). Similar displacement amplitudes were imposed on disordered sites overlapping by less than the sum of van der Waals radii. The site occupancy ratio was allowed to freely refine.

H atom treatment - Methyl H atom positions, R-CH₃, were optimized by rotation about R-C bonds with idealized C-H, R--H and H--H distances. Remaining H atoms were included as riding idealized contributors. Methyl H atom U's were assigned as 1.5 times U_{eq} of the carrier atom; remaining H atom U's were assigned as 1.2 times carrier U_{eq} .

The -1 1 1 and 0 2 0 reflections were omitted from the final refinement due to being partially obscured by the beam stop in some orientations.

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(2) Krause, L., Herbst-Irmer, R., Sheldrick, G. M. and Stalke, D. *J. Appl. Cryst.*, **2015**, *48*, 3-10.

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- (5) Guzei, I. A. J. Appl. Cryst., **2014**, 47, 806-809.