

Relaxation Dispersion NMR to reveal fast Dynamics in Brønsted Acid Catalysis: Influence of Sterics and H-bond strength on conformations and substrate hopping

N. Lokesh, Johnny Hioe, Johannes Gramüller and Ruth M. Gschwind*

Institute for organic chemistry,
University of Regensburg, D-93053 Regensburg, Germany

E-mail : ruth.gschwind@chemie.uni-regensburg.de

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Experimental data

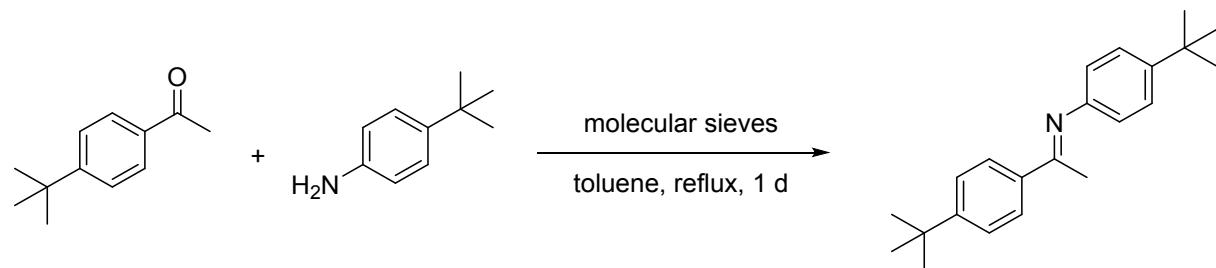
Deuterated solvents were purchased from Deutero or Sigma Aldrich. Where dry solvents were essential, CD_2Cl_2 was freshly distilled over CaH_2 and Toluene was refluxed over Na/Benzophenone under argon atmosphere. The catalysts were purchased from Sigma Aldrich or STREM chemicals.

High-resolution mass spectra were measured by the central analytics division in the Institute of Organic Chemistry. Gas chromatography coupled with a mass selective detector was performed on an Agilent 6890N Network GC-System.

Synthesis of Imine Substrates

The imines were prepared as described in the literature.¹⁻⁴ The toluene was used either in p.A. quality or was dried by refluxing over sodium. The ¹⁵N-enriched aniline (if used) was purchased from Euriso-top GmbH and Sigma Aldrich.

(E)-N,1-bis(4-(tert-butyl)phenyl)ethan-1-imine [5]



Molecular sieves 4 Å (4 g) was weighed into a 50 mL Schlenk flask and dried with a heat gun at 350 °C for 30 min under reduced pressure. Under Argon flow, 4-(*tert*-butyl)aniline (10.0 mmol, 1.49 g, 1.60 mL, 1.0 eq) and 1-(4-(*tert*-butyl)phenyl)ethan-1-one (10.0 mmol, 1.76 g, 1.83 mL, 1.0 eq.) were added and dissolved in 20 mL anhydrous toluene. Under Argon flow, a reflux condenser was added to the setup and flushed with argon for 3 min. A drying tube filled with CaCl₂ was added to the setup. The solution was refluxed for 24 h. Afterward, the heating bath was removed, and the reaction mixture allowed to cool down. The orange mixture was filtrated, and the solvent was removed under reduced pressure to give an orange-yellow solid. The crude product was washed with methanol to give the product (4.76 mmol, 1.46 g, 48%) as yellow needles predominantly as its *E*-isomer (>99% determined by ¹H NMR).

¹H-NMR (400.1 MHz, CD₂Cl₂) δ_H = 7.91 (m, 2H), 7.48 (m, 2H), 7.37 (m, 2H), 6.71 (m, 2H), 2.21 (s, 3H), 1.36 (s, 9H), 1.34 ppm (s, 9H).

¹³C-NMR {¹H} (100.6 MHz): δ_C = 164.8, 153.7, 149.2, 145.8, 137.0, 126.8, 125.7, 125.2, 119.0, 34.7, 34.1, 31.2, 31.0, 16.9 ppm.

HR-MS (ESI, *m/z*): found 307.2289 (M)⁺ (calculated for 307.2300 for C₂₂H₂₉N); Diff(ppm) = -1.92 ppm.

Sample preparation

Preparation of binary complexes in CD₂Cl₂

The catalyst was dried for 30 min at 150°C under reduced pressure. Ketimine and catalyst were directly weighed into a 5 mm NMR tube under an inert argon atmosphere. CD₂Cl₂ (0.6 ml; freshly distilled over CaH₂) and 1.0 ml of tetramethylsilane atmosphere were added to the tube. The sample was stored in an -80°C freezer. A 1:1 ratio of catalyst/ketimine was used for all samples unlike stated otherwise. A concentration of 0.25 – 100 mM was used, as indicated specifically.

Spectrometer data

NMR experiments were performed on Bruker Avance III HD 400 MHz spectrometer, equipped with 5 mm BBO BB-1H/D probe head with Z-Gradients and a Bruker Avance III HD 600 MHz spectrometer, equipped with a 5 mm CPPBBO BB-1H/19F. The temperature was controlled in the VT-experiments by BVT 3000 and BVTE 3900. For NMR measurements employing standard NMR solvents 5 mm NMR tubes were used, if not otherwise noted. NMR Data were processed, evaluated and plotted with TopSpin 3.2 software. Further plotting of the spectra was performed with Corel Draw X14 – X17 software. ^1H , ^{13}C chemical shifts were referenced to TMS or the respective solvent signals. The heteronuclei ^{19}F and ^{31}P were referenced, employing $\nu(\text{X}) = \nu(\text{TMS}) \times \Xi_{\text{reference}} / 100\%$ according to Harris et al.⁵ The following frequency ratios and reference compounds were used: $\Xi(^{19}\text{F}) = 94.094011 (\text{CCl}_3\text{F})$, $\Xi(^{31}\text{P}) = 40.480742 (\text{H}_3\text{PO}_4)$.

Pulse programs

All pulse programs used are standard Bruker NMR pulseprograms except for the $R_{1\alpha}$ measurements.

Acquisition Parameters

^1H NMR: Pulse program: zg; Relaxation delay = 2 – 3 s, Acquisition time = 2.48 s, SW = 22.0 ppm, TD = 64k, NS = 8 – 64; zg30; Relaxation delay = 2 s, Acquisition time = 2.48 s, SW = 22.0 ppm, TD = 64k, NS = 8 – 64;

^2D - $^1\text{H},^1\text{H}$ NOESY: Pulse program: noesygpph/noesygpphpp; Relaxation delay = 5 - 8 s, NS = 8-32, mixing time (D8) = 300.00 ms; TD = 4096; increments = 512 - 1k;

^2D - $^1\text{H},^1\text{H}$ COSY: Pulse program: cosygpqf; Relaxation delay = 5 - 8 s, NS = 8-32, TD = 4096; increments = 512 - 1k;

^{13}C NMR: Pulse program: zpg30; Relaxation delay = 2.00 s, Acquisition time = 0.80 s, SW = 270.0 ppm, TD = 64k, NS = 1k – 2k;

^2D - $^1\text{H},^{13}\text{C}$ HSQC: Pulse program: hsqcedetgpsisp2.3; Relaxation delay = 4 - 8 s, NS = 8-32, $^1J_{\text{CH}}$ = 145 Hz; TD = 4096; increments = 512 - 1k;

^2D - $^1\text{H},^{13}\text{C}$ HMBC: Pulse program: hmbcgplpndqf; Relaxation delay = 4 - 8 s, NS = 8-32, $^1J_{\text{CH}}$ = 145 Hz, J_{CH} (long range) = 10 Hz; TD = 4096; increments = 512 - 1k;

^2D - $^1\text{H},^{31}\text{P}$ HMBC: Pulse program: inv4gplrndqf; Relaxation delay = 4 - 8 s, NS = 8-32, TD = 4096; increments = 256 - 1k;

^{15}N NMR: Pulse program: zg; Relaxation delay = 10.00 s, Acquisition time = 0.54 s, SW = 502.8 ppm, TD = 32k, NS = 256 – 2048;

^2D - $^1\text{H},^{15}\text{N}$ HMBC: Pulse program: inv4gplrndqf; Relaxation delay = 5 - 8 s, NS = 16-32, delay for evolution of long range couplings (D6) = 20.00 ms; TD = 4096; increments = 128 - 512;

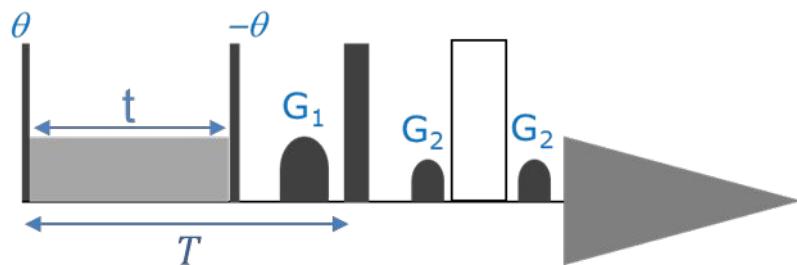
^{19}F -NMR: Pulse program: zg30; Relaxation delay = 2 – 3 s, Acquisition time = 11.60 s, SW = 10.0 ppm, TD = 128k, NS = 8 – 64;

2D-¹H,¹⁹F HOESY: Pulse program: hoesyph; Relaxation delay = 5 - 8 s, NS = 16-32, mixing time (D8) = 500.00 ms; TD = 4096; increments = 1k;

1D ^1H off-resonance $R_{1\rho}$

Pulse Sequence

The recent reported 1D ^1H off-resonance $R_{1\rho}$ pulse sequence is adopted with slight modification for our measurement.^{6,7}



$$R_{1\rho} = -\frac{1}{T} \ln \frac{I}{I_0}$$

Figure 1: Applied pulse sequence and mathematical relation to measure $R_{1\rho}$ experimentally. The applied angle (θ) is 35° . A spinlock period (t) of 100 ms is applied. The filled (black) and unfilled rectangular pulses respectively indicate 90° and 180° hard pulses. The gradient pulses of $G_1 = 70\%$ and $G_2 = 40\%$ of the maximum were applied. I_0 is the intensity without any spinlock period and I is the intensity with spinlock period measured over a multiple effective magnetic field.

Parameter Optimization

The spinlock period (t) was incremented in 10 ms steps with maximum spinlock power (≈ 20 kHz). At each step, the temperature was monitored (sample heating can occur due to application of high intensity spinlock field over an extended time). Around 120 ms, temperature oscillation was observed. To avoid any influence of temperature changes on the measurement, the spinlock period of 100 ms was applied in all our measurements.

The normal relaxation period ($T-t$) in the absence of the spinlock is tested for 3 delays (50 ms, 100 ms and 150 ms). For all three different relaxation periods ($T-t$), we found similar exchange rates within the experimental errors. To minimise signal loss due to relaxation, ($T-t$) of 50 ms was used in all our measurements.

Method testing

To test the applicability of the R_{1p} method, we applied the above described pulse sequence with optimized parameters on the known peak in the spectrum (*para*-methyl protons of the hydrolyzed imine ketone part), which does not experience any chemical exchange process (on a ms- μ s timescale) and on the same proton for the CPA/imine complex which experiences chemical exchange. The measured rate for the hydrolyzed ketone showed a nearly straight line (no offset-Lorentzian decay with increasing effective field ω_{eff}), which was expected for non exchanging protons. For the same proton in the binary CPA/imine complex, an offset-Lorentzian decay was observed, which demonstrates the presence of chemical exchange on a ms- μ s timescale.

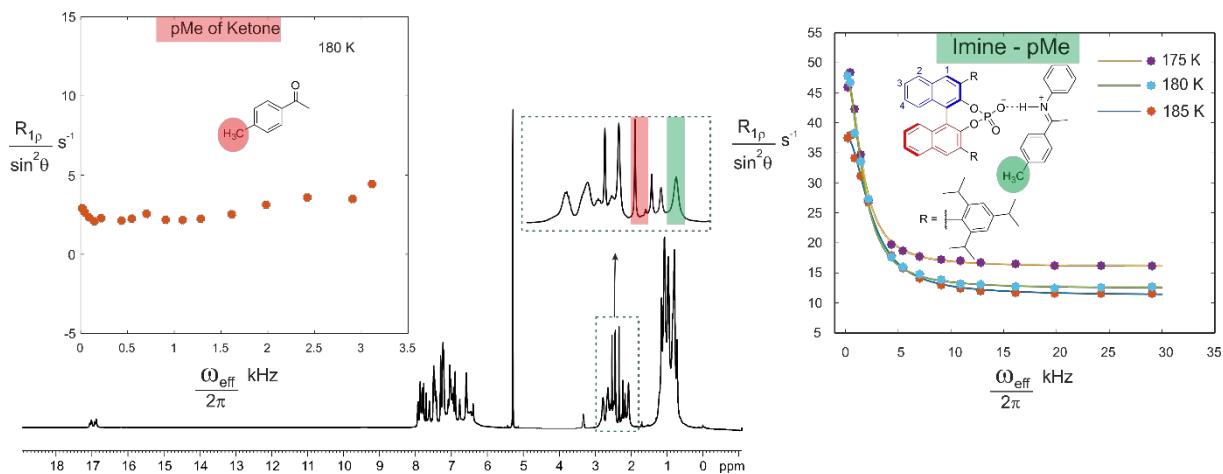


Figure 2: To test the method, we selected the proton peak of the *p*-methyl group in the hydrolyzed ketone moiety (red shaded) and the proton peak of the *p*-methyl imine in the TRIP/imine binary complex (green shaded) within the same sample (to maintain identical conditions). The hydrolyzed ketone does not undergo any chemical exchange on a ms- μ s time scale. The corresponding measured R_{1p} data at 180 K with incremental effective field showed a near horizontal line, proving the absence of chemical exchange (left side). On the other hand, the imine inside the catalyst undergoes $E_{\text{i}} \rightleftharpoons E_{\text{II}}$ exchange and thus the corresponding R_{1p} data at 185, 180 and 175 K with incremental effective field showed offset-Lorentzian decays for the *p*-methyl protons in the TRIP/imine complex. This confirms the presence of chemical exchange process and hence validates our method for chemical exchange detection in ms- μ s range.

Curve fitting

In order to analyze the R_{1p} measurements, the signal intensities (integrals) are determined for every effective field strength ω_{eff} (i.e. every different spinlock power) and normalized to the reference integral (spinlock power = 0 W). Initially, to extract k_{ex} , the measured experimental points are curve fitted to the following theoretically described equation by using MATLAB (see literature for further details⁸).

$$\frac{R_{eff}}{\sin^2\theta} = R_2 - R_1 + \frac{(\Delta\omega)^2 P_A P_B \tau_{ex}}{(1 + \tau_{ex}^2 \omega_{eff}^2)} \quad (Eq. 1)$$

Here R_2 is the transverse relaxation rate constant

R_1 is the longitudinal relaxation rate constant

$\Phi = (\Delta\omega)^2 P_A P_B \tau_{ex}$ encodes both chemical shift difference and population of the minor (P_B) and major (P_A) exchange species

Φ and the average isotropic chemical shift Ω (in ppm) are related by

$$\Phi = -\Omega^2 + a\Omega + b \quad (Eq. 2)$$

Here,

$$a = 2\Omega_A - \Delta\omega$$

$$b = \Omega_A(\Delta\omega - \Omega_A)$$

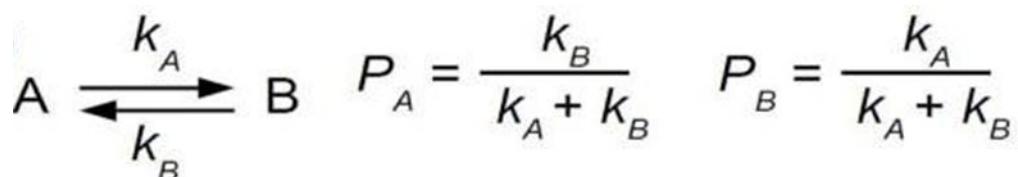
These can be rearranged as

$$\Delta\omega = (a^2 + 4b)^{1/2}, \text{ and } \Omega_{A,B} = [a \pm ; (a^2 + 4b)^{1/2}] / 2 \quad (Eq. 3)$$

Φ and Ω can be obtained experimentally at different temperatures. Assuming the chemicalshift difference $\Delta\omega$ between major (A) and minor (B) species is constant, one can calculate $\Delta\omega$. From $\Delta\omega$ value and Φ values, populations can be extractable.

Directionality

For the chemical exchange between A and B, the exchange rate constant k is: $k_{ex} = k_A + k_B$. By using populations and k_{ex} one can extract rate constants k_A and k_B on both sides.



If a temperature independence within the varied range of 10 K is assumed, then the coefficients a and b can be determined by least squares optimization. Subsequently, $\Delta\omega$ is calculated by $\Delta\omega = (a^2 + 4b)^{1/2}$, and $\Omega_{A,B} = [a \pm (a^2 + 4b)^{1/2}]/2$. Once $\Delta\omega$ is known, P_A , P_B , k_A and k_B are obtained from Φ_{ex} and k_{ex} at each temperature.

Hence, in order to separate the populations and chemical shift difference from the term ϕ , it requires additional experimental R_{1p} measurements at more than one temperature. Therefore we measured R_{1p} experiments for each sample at two or three temperatures (185 K, 180 K and 175 K).

Results of the R_{1p} measurements

TRIP/3

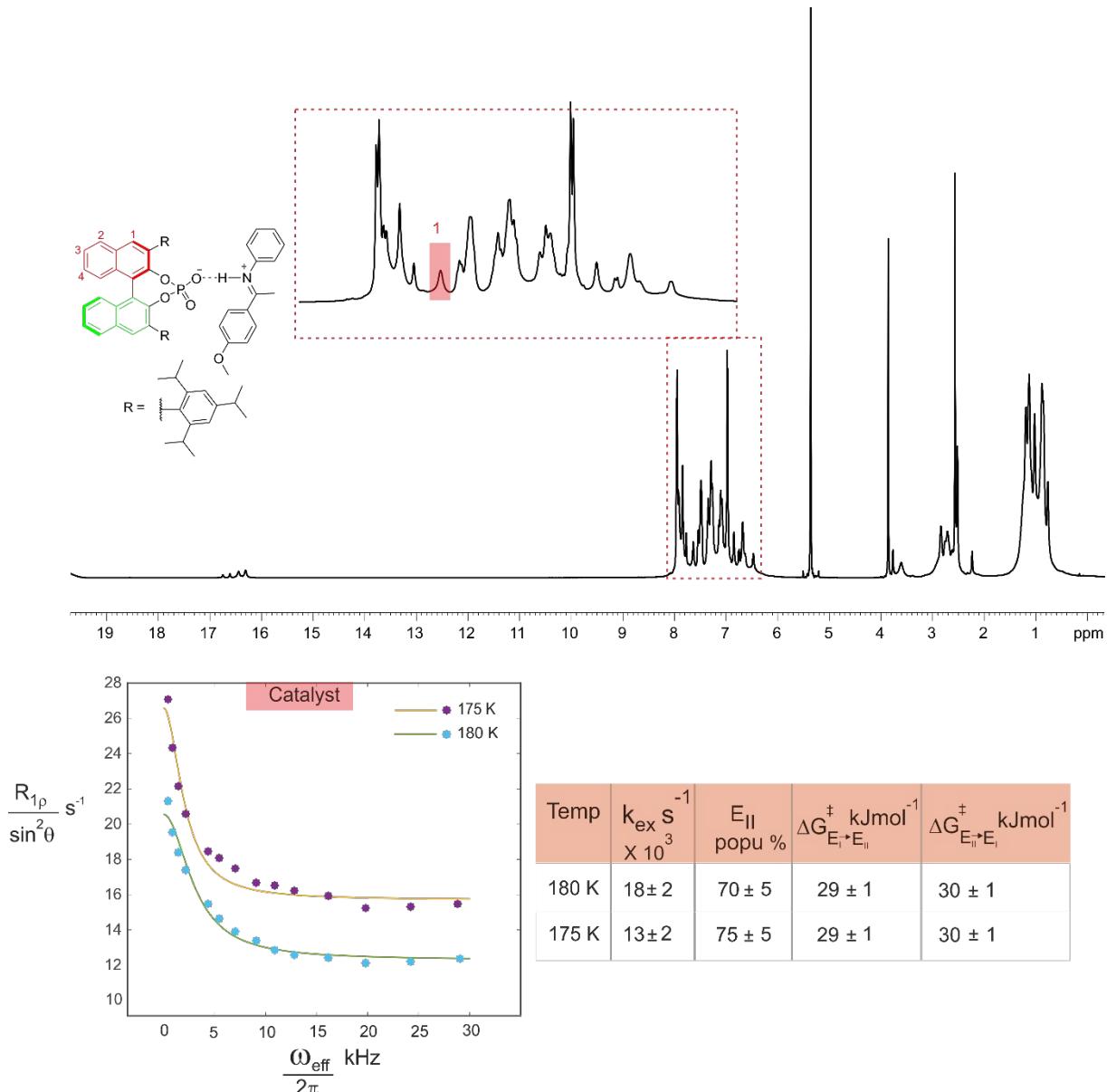


Figure 3: R_{1p} measurement for the TRIP/3 complex. The proton H_1 of the catalyst (marked in the spectrum, red shaded) was selected as probe. The offset-Lorentzian decay indicates the

presence of chemical exchange ($k_{\text{ex}} \approx 18000 \text{ s}^{-1}$ at 180 K). With experimental data obtained at two different temperatures, the populations and directionality were extracted.

TRIP/2

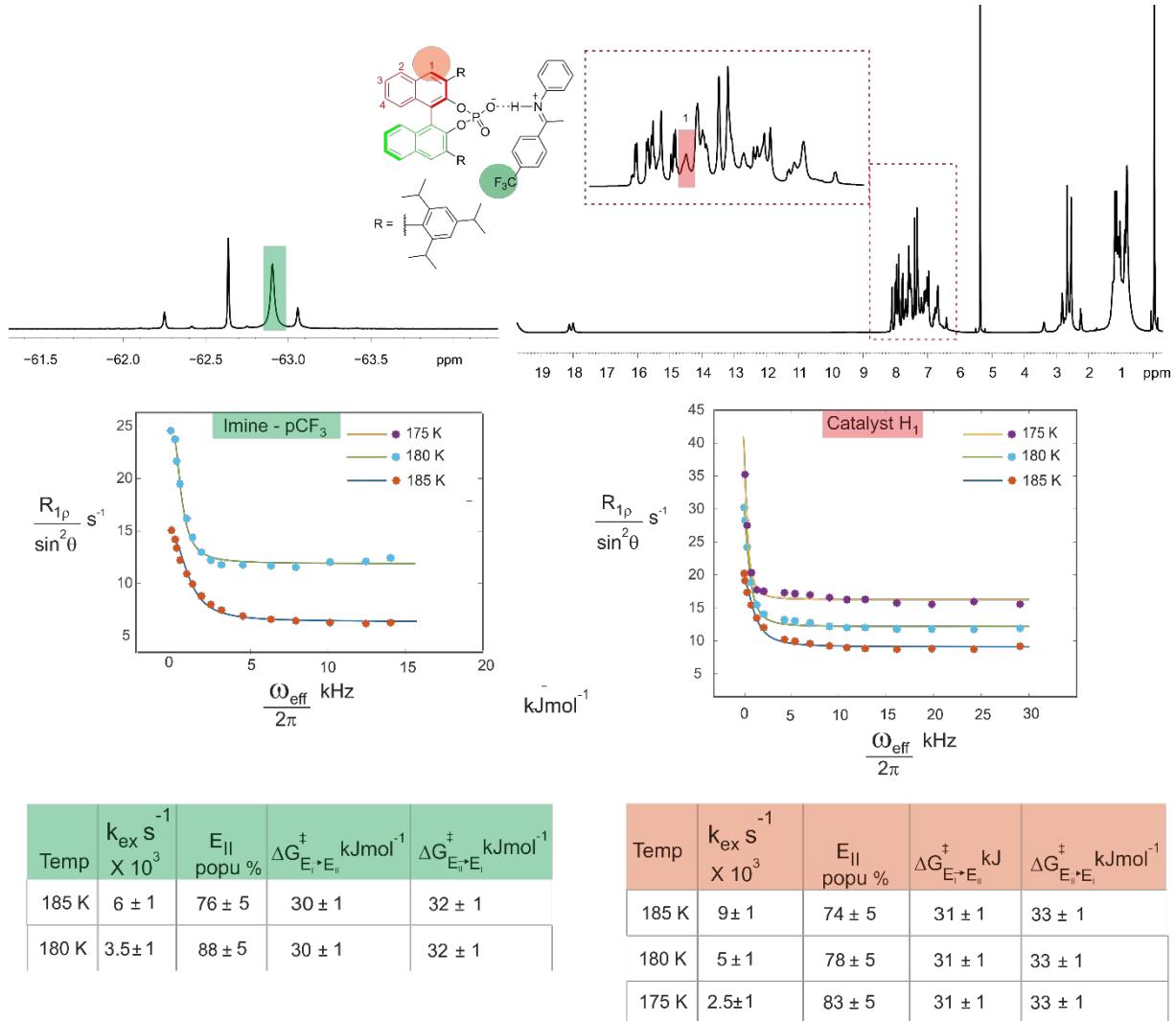


Figure 4: $R_{1\rho}$ measurement for TRIP/2 complex. The proton H₁ of the catalyst (shaded in red) and the fluorine signals of the imine-CF₃ group (shaded in green) were selected as probes. The offset-Lorentzian decay indicates presence of chemical exchange ($k_{\text{ex}} \approx 3-5 \cdot 10^3 \text{ s}^{-1}$ at 180 K). With experimental data obtained at three different temperatures, the populations and directionality were extracted.

TiPSY/1

Temp K	$k_{\text{ex}} \text{ s}^{-1} \times 10^3$	$E_{\text{ii}} \%$	$\Delta G_{E_i \rightarrow E_{\text{ii}}}^{\ddagger} \text{ kJ mol}^{-1}$	$\Delta G_{E_{\text{ii}} \rightarrow E_i}^{\ddagger} \text{ kJ mol}^{-1}$
185	3.0 ± 1	95 ± 5	32 ± 1	36 ± 1
180	2.5 ± 1	96 ± 4	32 ± 1	36 ± 1
175	2.5 ± 2	95 ± 5	31 ± 1	35 ± 1

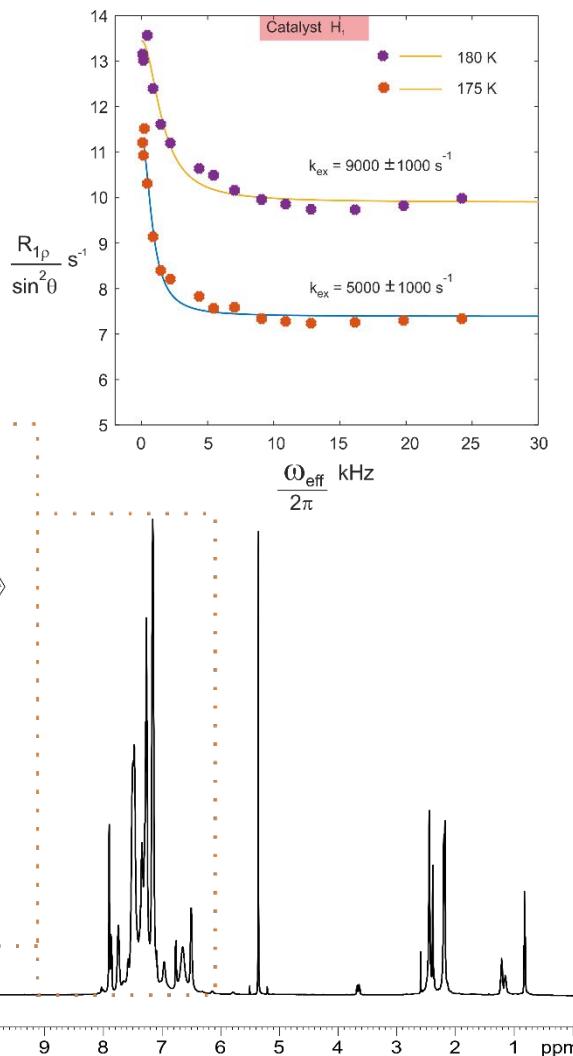


Figure 5: R_{1p} measurement for TiPSY/1 complex, (to avoid signal overlap, the sample was prepared with nearly exclusive population of the *E*-complex; see above for sample preparation). The proton H_1 of the catalyst (shaded in red) was selected as probe. The offset-Lorentzian decay indicates presence of chemical exchange ($k_{\text{ex}} \approx 9 \text{ ks}^{-1}$ at 180 K).

TRIM/2 and TRIM/3

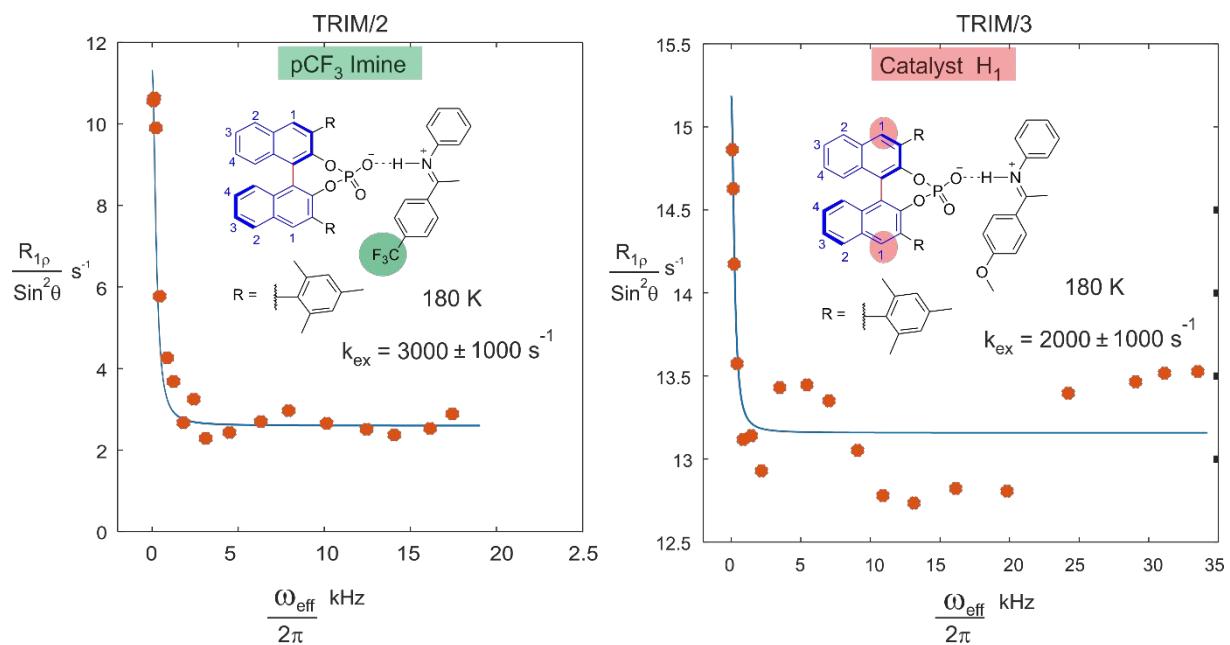


Figure 6: $R_{1\rho}$ measurements for TRIM/2 and TRIM/3 complexes. For TRIM/2, the CF_3 group of the imine was used as a probe (shaded in green) and for TRIM/3 the proton H_1 of the catalyst (shaded in red) was selected as probe. The offset-Lorentzian decay in both system indicates the presence of chemical exchange ($k_{\text{ex}} \approx 2000-3000 \text{ s}^{-1}$ at 180 K).

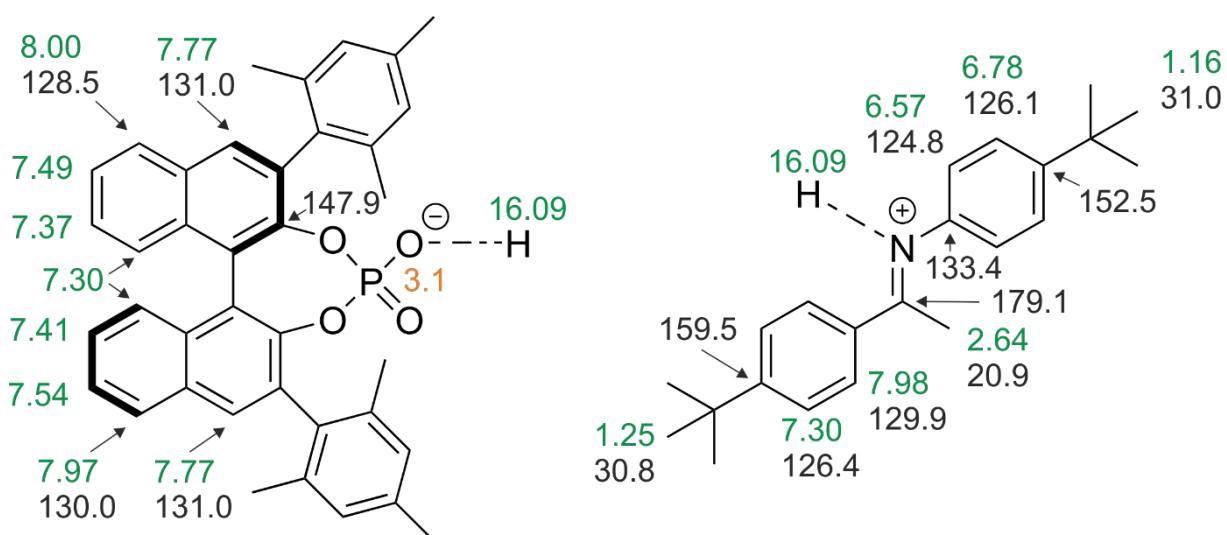
For TRIM/2 and TRIM/3, the offset-Lorentzian decay fit does not match the experimental decay as precisely as for the other systems. Hence, a larger experimental error is expected. However, the $R_{1\rho}$ measurements for TRIM/2 and TRIM/3 complexes indicate, that similar exchange rates ($k_{\text{ex}} \approx 2000-3000 \text{ s}^{-1}$ at 180 K) are obtained for TRIM/1-3 regardless of the hydrogen bond strength.

NMR-Structural analysis of CPA/imine complexes

The investigated complexes TRIP/**1-3**, TiPSY/**1** and TRIM/**1-3** were already described in the previous work.^{2,9} The NMR-structural analysis of TRIM/**4** is described below. In addition, the backbone splitting of TiPSY/**1** is discussed in detail below.

Chemical Shift assignment of TRIM/**4**

The ¹H (black) and ¹³C (green) chemical shifts (in ppm) of all investigated complexes were assigned with standard 2D NMR experiments (¹H,¹H COSY, ¹H,¹H TOCSY, ¹H,¹H NOESY, ¹H,¹³C HSQC, ¹H,¹³C HMBC) at 180 K. The ³¹P (orange) chemical shift (in ppm) was assigned by ¹H,³¹P HMBC. Due to signal overlap and lacking resolution and intensity of 2D correlations, not all signals could be assigned.



In accordance with our previous analysis of hydrogen bond strengths,¹ the ¹H chemical shift of the hydrogen bond strength reflects the H-bond strength (in our system, higher chemical shifts correlate with stronger H-bonds). Hence, the H-bond strength of TRIM/**4E** (16.09 ppm) is in between TRIM/**1E** (16.26 ppm) and TRIM/**3E** (15.80 ppm).⁹

Identification of Type I E and Type II E

Analogous to our previous work,^{2,9} for the *Type I E* structure, the *tert*-butyl group of the ketone part of the imine (marked in blue) is placed above the BINOL backbone. The blue NOE cross signals A and B to the Naphthyl backbone reveal the presence of the *Type I E* structure (Figure 7) for TRIM/**4E**. The NOE cross signal D (black) is a result of overlapping NOE signals between the *tert*-butyl group and the naphthyl backbone and the *tert*-butyl group and the attached

phenyl ring (backbone: 8.00 ppm, phenyl ring of imine: 7.98 ppm). For the *Type II E* structure, the tert-butyl group of the aniline part of the imine is placed above the BINOL backbone and thus the red NOE cross signals A-D reveal the presence of the *Type II E* structure.

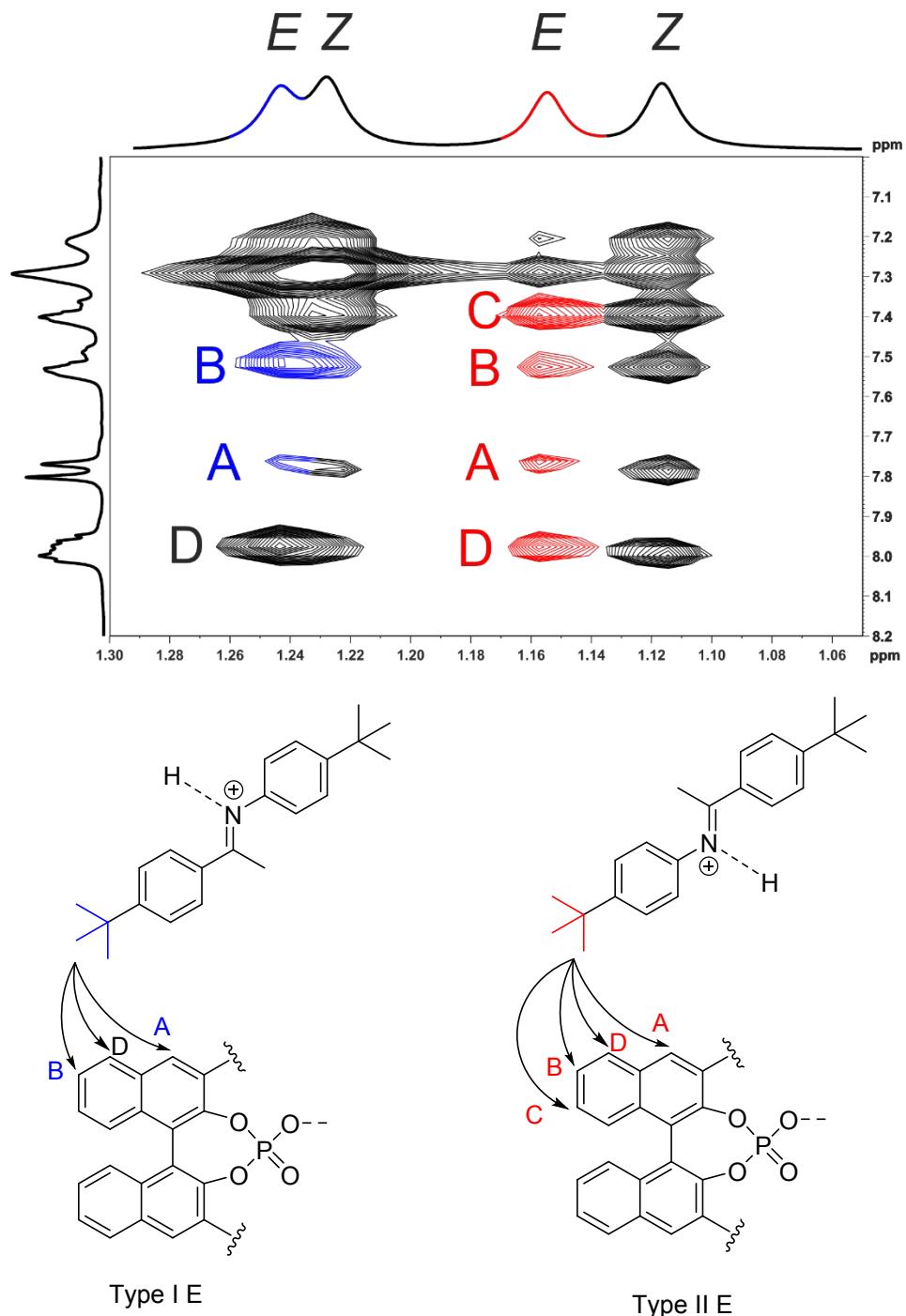


Figure 7: Excerpt of the $^1\text{H},^1\text{H}$ NOESY spectrum of TRIM/4 at 180 K and 600 MHz (100 mM sample) in CD_2Cl_2 . The blue NOE cross signals indicate the presence of the *Type I E* structure, while the red NOE signals indicate the presence of the *Type II E* structure.

DOSY measurements of TRIM/4

To prove, that the observed species in TRIM/4 are monomers of the binary complexes, DOSY measurements (Diffusion ordered spectroscopy) were performed to derive the molecular radii of the present species.

The DOSY measurements were performed with the convection suppressing DSTE (double stimulated echo) pulse sequence developed by Jerschow and Müller in a pseudo 2D mode.¹⁰ TMS was used to reference the viscosity of the solvent at 180 K. The diffusion time delay was set to 45 ms. The gradient pulse lengths (p16, SMSQ10.100 pulse shape) were optimized for each species to give a sigmoidal signal decay for varying gradient strengths. Optimal pulse lengths of 3.0 ms and 6.0 ms were found for TMS and TRIM/4, respectively. For each species, twenty spectra with linear varying gradient strength of 5% - 95% have been measured. The used probe signals for the analysis are listed in table 1. The signal intensities of the respective groups were analyzed as a function of the gradient strength by Bruker TopSpin 3.2 software T1/T2 relaxation package by employing the Stejskal-Tanner equation.¹¹ No line broadening occurred for increasing gradient strength. The sigmoidal fit provided the translational self-diffusion coefficients D_i listed in Table 1. The molecular radii were derived by the Stokes-Einstein equation¹² using Chens correction.¹³

$$D_i = \frac{k_B T}{6\pi\eta r_H} * (1 + 0.695 * (\frac{r_{solv}}{r_H})^{2.234}) \quad (Eq. 4)$$

D_i is the self-diffusion coefficient derived by the measurement, η is the viscosity of the solvent, r_H is the hydrodynamic radius of the observed molecule and r_{solv} the radius of the solvent . No form factor correction was applied. The viscosity was determined by measuring the diffusion coefficient of the reference tetramethylsilane (TMS) and solving the equation for η with the literature value¹⁴ of the radius of 2.96 Å. The solvent radius of CD₂Cl₂ (2.46 Å) was taken from the reference.¹⁵

Table 1: Probe signals, measured diffusion coefficients and derived molecular radii for TRIM/5.

Entry	Species	p16 [ms]	Observed signal [ppm]	D_i [m^2/s] · 10 ⁻¹²	Averaged	r_H [Å]
1	TMS	3.10	0.00	21.32		
2	TRIM/4E	6.00	1.16	4.76		
3	TRIM/4E	6.00	2.64	4.80	4.69	9.54
4	TRIM/4E	6.00	7.77	4.50		

The derived molecular radius (9.54 Å) is similar to the one reported previously for TRIM monomers.⁹ Thus, the DOSY measurements showed, that the species investigated by the R_{1ρ} measurement is the binary TRIM/4E complex.

Estimation of the rotational correlation time of binary CPA/imine complexes

A general rotational correlation time of 10-50 ns was estimated for the investigated binary complexes of chiral phosphoric acids and imines with their hydration shell. Stokes Law (Eq. 5) was used to calculate the rotational correlation time based on the results of the DOSY measurements of TRIM/4E.

$$\tau_c = \frac{4\pi\eta r^3}{3kT} \quad (\text{Eq. 5})$$

The radius of the binary complex of TRIM/4 with its hydration shell was determined by DOSY measurements (see respective chapter) and a radius of 9.54 Å was determined for the complex with its hydration shell. The viscosity (3.05×10^{-2} kg m⁻¹ s⁻¹) of the solution at 180 K was determined by Equation 4 based on the DOSY measurement on TMS. A rotational correlation time of 44.6 ns was calculated for TRIM/4E with its hydration shell. Given the bulkiness of the two tert-butyl substituents of imine 4, it is expected that the other investigated complexes have smaller radii, resulting in smaller rotational correlation times. Hence, a general rotational correlation time around 10 – 50 ns was estimated.

TiPSY/1

In the TiPSY/1 complex, the backbone splitting could not be resolved at 180 K and 600 MHz in CD_2Cl_2 (see figure). However, in our previous investigations we observed, that the backbone splitting in TiPSY complexes is very small and could often not be resolved sufficiently (see SI of literature).⁹ Goodman *et al.* demonstrated, that the binding pocket for TiPSY is significantly smaller than for TRIP (no rotation possible).¹⁶ The biggest binding pocket for the investigated CPAs was determined for TRIM (rotation possible). Hence, a rotation of the imine inside the binding pocket of TiPSY can be excluded, even if the backbone splitting can not be resolved sufficiently.

The sample was prepared with an exclusive population of the *E*-imine. To achieve that, the sample was prepared according to the general procedure described above. However, before adding the solvent, the NMR-tube with TiPSY and **1** was cooled to -90 °C in acetone/liquid N_2 and the precooled solvent was added to the tube. The low temperature suppressed $E \rightarrow Z$ isomerization of the imine.

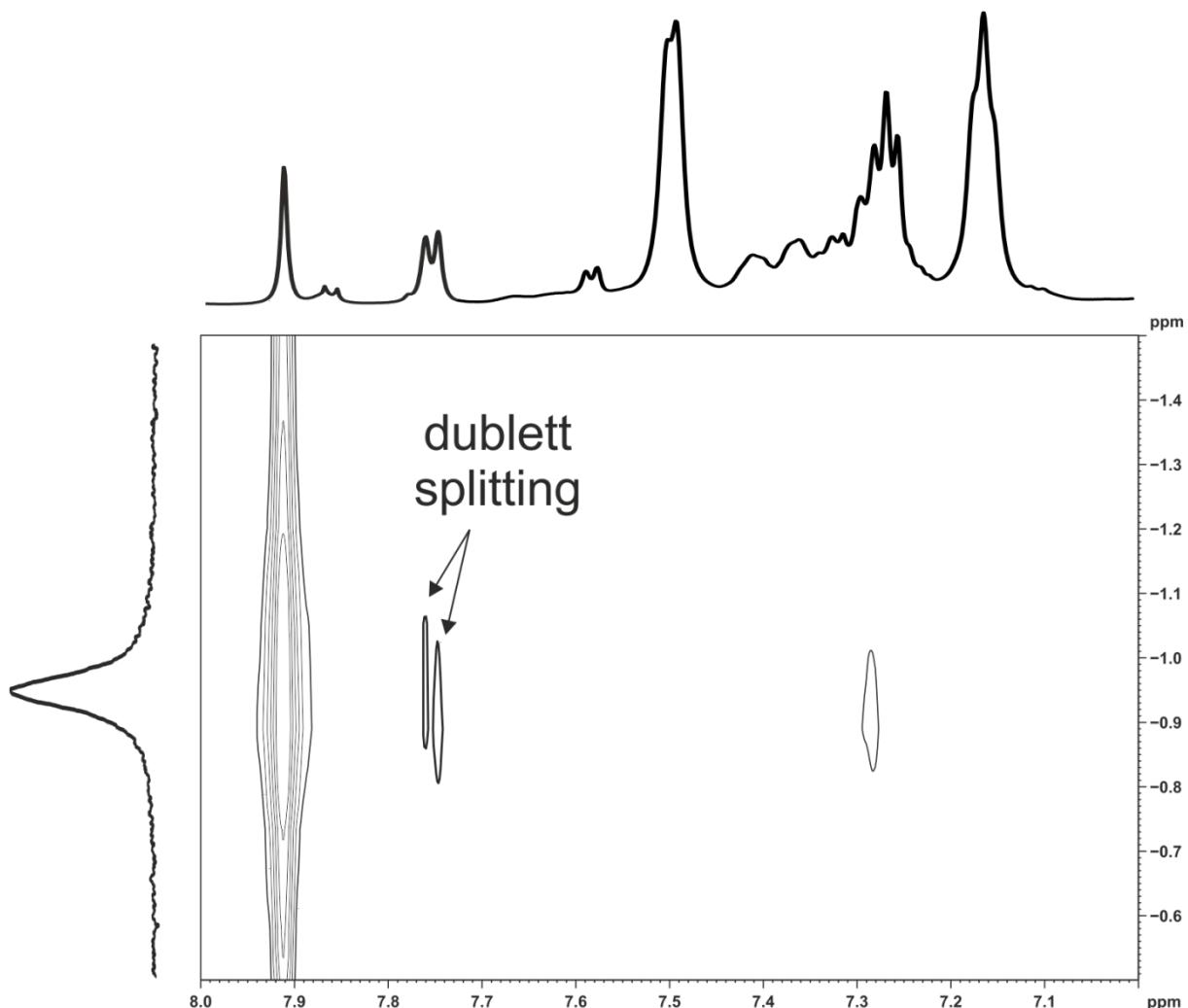


Figure 8: Excerpt of the $^1\text{H}, ^{31}\text{P}$ HMBC spectrum of TiPSY/1 (E-only, 25 mM, TiPSY:**1** = 1:1) at 180 K and 600 MHz in CD_2Cl_2 . The sample was prepared with an exclusive population of *E*-imine species.

Exchange with free imines

Typically, in 1:1 complexes of TRIP, TRIM and TiPSY, the complexation of the imine was incomplete ($\approx 25\%$ free imine present). Thus, for complexes with imines **1-3**, the assignment of the free imines was done for the CPA/imine samples. For imine **4**, a NMR sample with the imine only (50 mM, CD_2Cl_2 , 180 K and 600 MHz) was measured and compared to the chemical shifts in the CPA/**4** sample. For TRIM/**4**, no free imine was observed in the binary complex sample.

As the free imines **1-4** gave chemical shifts well separated from the respective CPA/*E*-imine complexes (see figure 9 for free imines and assignments of the binary complex in literature^{2,9}), the exchange between free and complexated imine is slow on the NMR time scale. Thus, the exchange pathway via dissociation and re-association causing averaging of the BINOL backbone can be excluded to significantly contribute to the measured exchange rates in the $R_{1\sigma}$ measurements.

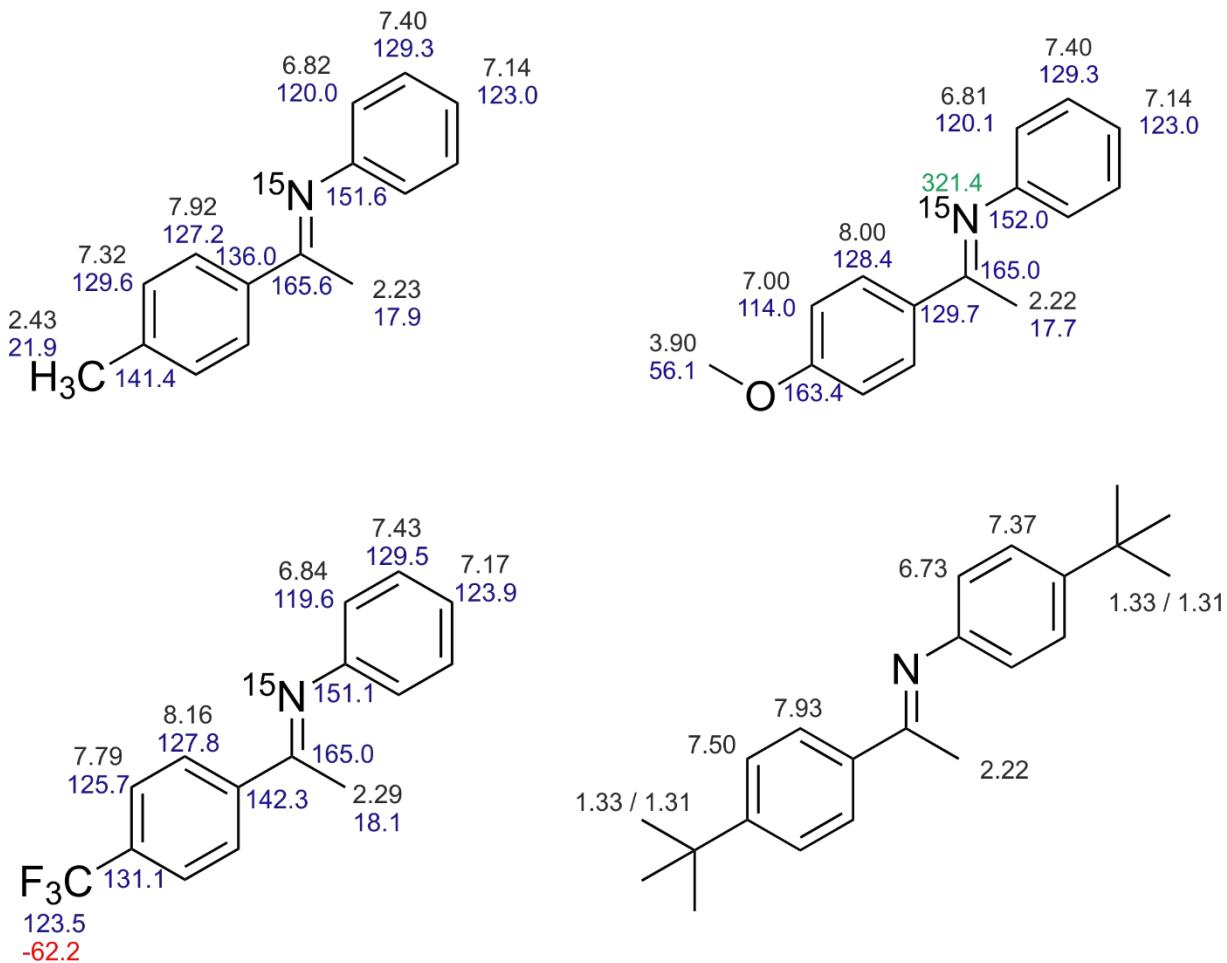


Figure 9: Chemical shift assignment of the free imines **1-4** at 180 K in CD_2Cl_2 .

Additional fitting of TRIM/1E

Due to the strong signal decay within the first few percent of the horizontal axis for TRIM/1E (Figure 6B in the manuscript), fewer data points are in the decay region relevant for the fit, which could affect the quality of the fit and thus the precision of the extracted rate. However, the fit curves for lower (500 s^{-1}) and higher ($10\,000 \text{ s}^{-1}$) exchange rates as the optimized one (2500 s^{-1}) significantly deviate from the data set (Figure 10).

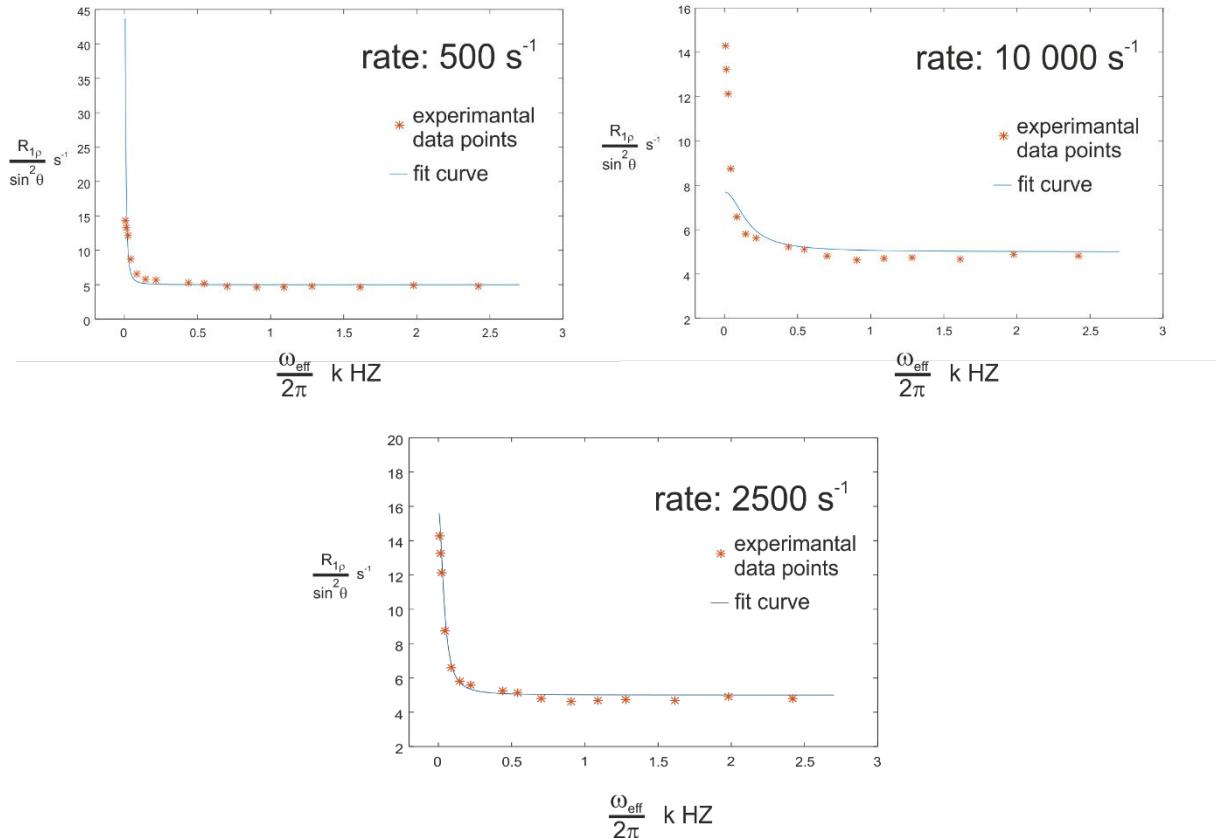


Figure 10: Additional curve fitting for the signal decay of TRIM/1E with different exchange rates at a concentration of 5 mM. The determined exchange rate of 2500 s^{-1} fits the data set best, while lower or higher exchange rates result in a significant offset. Other parameters, e.g. the population of E-I and E-II were not changed.

In addition, a TRIM/1E sample at different concentration (0.25 mM) was measured with more ω_{eff} increments. The extracted exchange rate of 2500 s^{-1} is identical to the previous one and the fit for lower or higher exchange rates does not match the data set (Figure 11).

Hence, the quality of the fit for TRIM/1E is adequately precise to reveal, that the exchange rate for TRIM/1E is significantly lower than for TRIP/1E ($\approx 10\,000 \text{ s}^{-1}$).

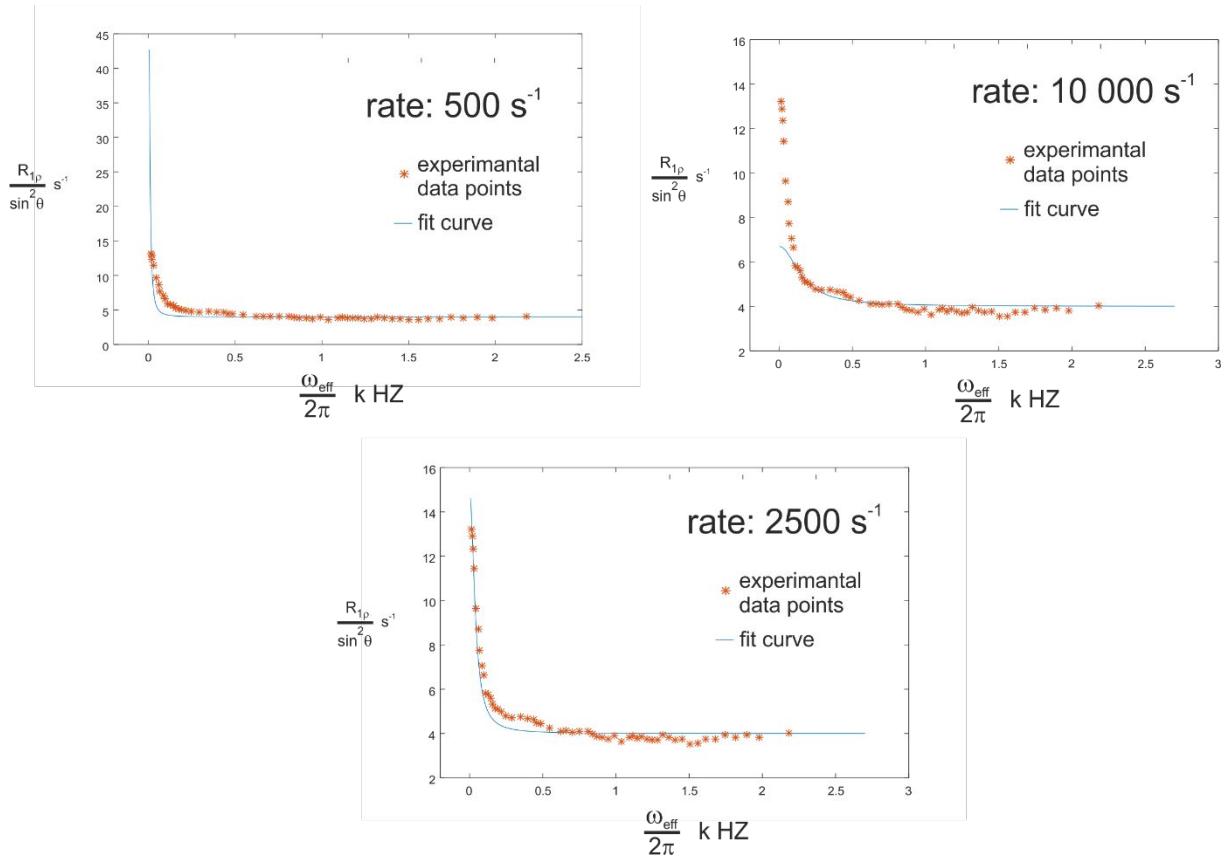


Figure 10: Additional curve fitting for the signal decay of TRIM/1E with different exchange rates at a concentration of 0.25 mM and additional ω_{eff} increments. The determined exchange rate of 2500 s^{-1} fits the data set best, while lower or higher exchange rates result in a significant offset. Other parameters, e.g. the population of E-I and E-II were not changed.

Computational Details

All structures were optimized at TPSS/def2-SVP^[17] level of theory using D3^[18] correction in continuum of CH₂Cl₂ (SMD^[19]). The dielectric constant was adapted to 16.20 to mimic low temperature condition in the experiment. Vibrational and thermochemical analyses were performed at the same level of theory as the geometry optimization. Subsequently, single point calculations at SCS-MP2/CBS^[20] level of theory were conducted. Def2-SVP and def2-TZVP basis sets were used for extrapolation to approach complete basis set. Solvent and thermochemical corrections were added to the single point energy. Software used for the geometry optimization, frequency analysis and solvent correction was Gaussian 09 version D.01.^[21] For single point calculation, ORCA 4.1.1 was used.^[22] Constant temperature (NVT) molecular dynamic was done using semi-empirical xtb software in DCM.^[23]

Extrapolation procedure

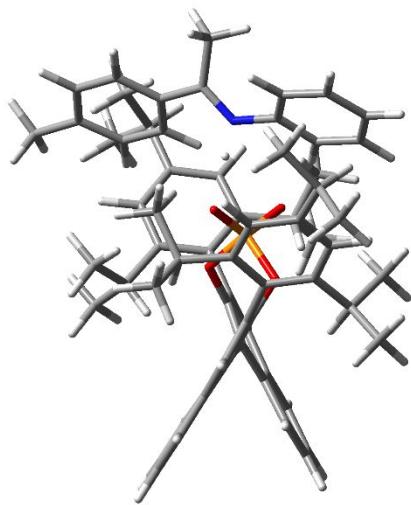
The CBS basis set extrapolation proceeds under the two point extrapolation procedure as in implemented in ORCA 4.1.1 with the basis set Def2-SVP and Def2-TZVP.

$$E_{\text{SCF}}^X = E_{\text{SCF}}^\infty + A e^{-\alpha \sqrt{X}}$$

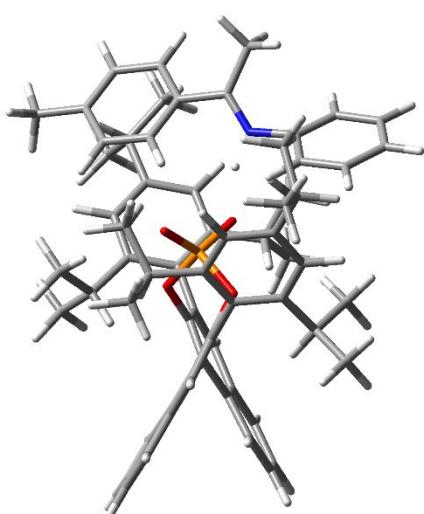
$$E_{\text{corr,MP2}}^\infty = \frac{X^\beta E_{\text{corr,MP2}}^X - (X-1)^\beta E_{\text{corr,MP2}}^{X-1}}{X^\beta - (X-1)^\beta}$$

$$E_{\text{MP2}}^\infty = E_{\text{corr,MP2}}^\infty + E_{\text{SCF}}^\infty$$

TRIP/1E Type III.1 and E-III.2 structures (most stable conformations)



Type E-III.1

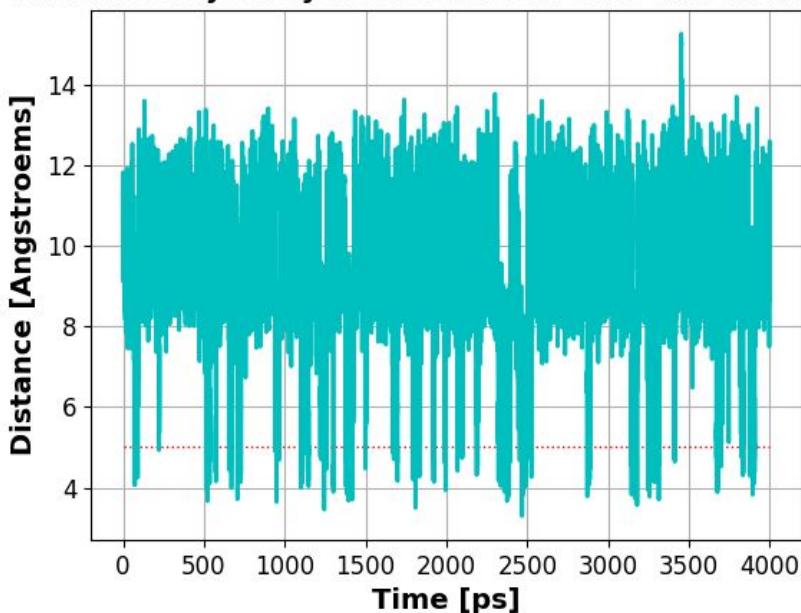


Type E-III.2

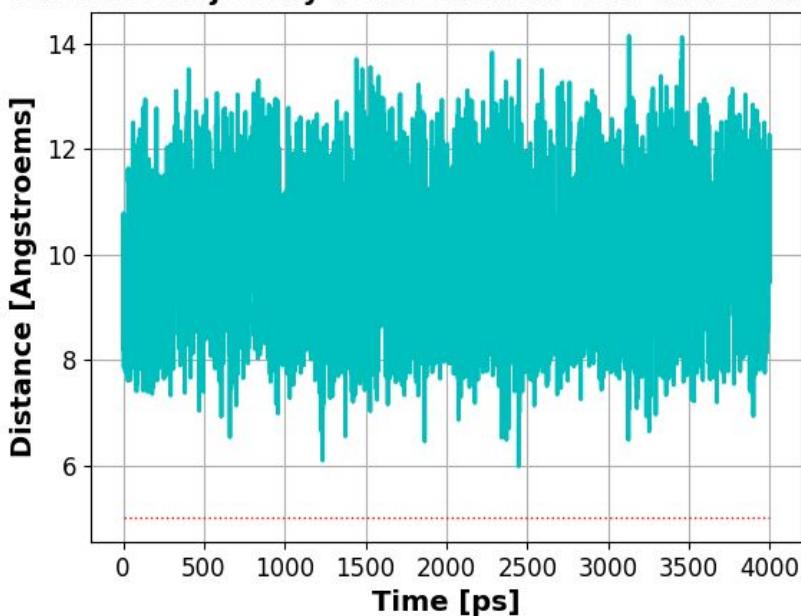
xtb-MD of TRIP/1E binary complex at 370 K

MD of TRIP/1E complex shows the exchange of E-II and E-I conformations via tilting and switching. The E-II conformation is marked by the close contact between C134 and H10, while the E-I conformation shows close contact between C140 and H25. No rotation is observed during a simulation of 4000 ps at 300 K to 370 K, i.e. C134 never meets H25 or C140 never meets H10. Beyond this temperature, dissociation of complex is observed.

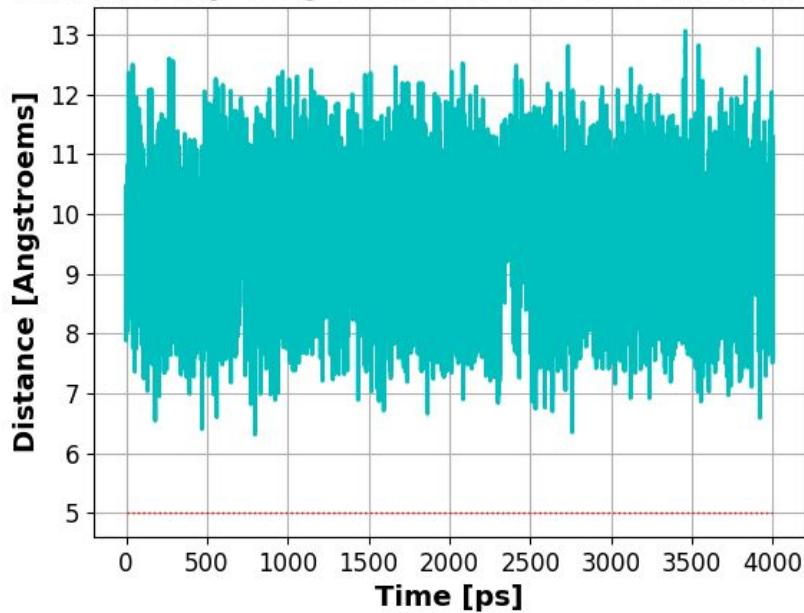
Distance trajectory between atom C140 and atom H25



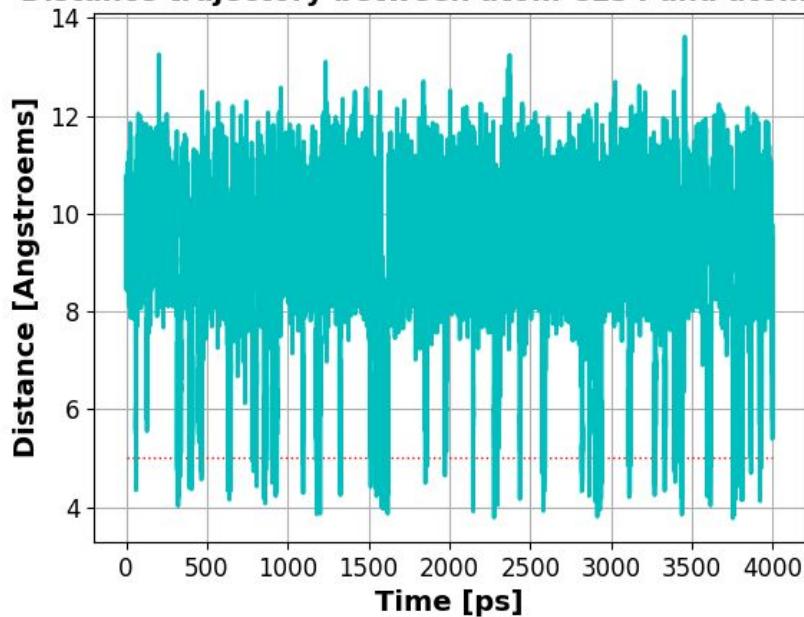
Distance trajectory between atom C140 and atom H10



Distance trajectory between atom C134 and atom H25



Distance trajectory between atom C134 and atom H10



Optimized Geometries

TRIM/1E Type I

```
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\0\#p tpsstpss/def2svp empiricaldispersion=gd3 int=ultrafine opt scrf
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1.1839383304O,-1.4175075558,0.5618941922,-1.4320413577O,0.6889832102
,0.3763631976,0.0168282895O,0.6300817555,-0.5399021282,-2.4784978792\
O,-0.5766080585,-1.7420153175,-0.4359277394\C,-1.446100498,2.147809437
,0.3711360845\O,-2.0698317941,1.122837238,-0.3502138217\O,-3.395018365
8,0.6663965667,-0.0681477603\O,-4.0703770977,1.2454430496,0.9985929916
\H,-5.095082782,0.9299435094,1.2232029452\O,-3.4593497944,2.2287931273
,1.8282185621\O,-2.1205478813,2.6764874421,1.5309470667\O,-1.502671696
6,3.5956395341,2.4329780968\H,-0.4706962922,3.9085577634,2.251357641\O
```

,-2.1840971756,4.0819139807,3.5389827427\H,-1.6850717191,4.7804794202,
 4.2194778391\C,-3.5190578898,3.6730083748,3.8040883989\H,-4.0466914589
 ,4.0672317544,4.679316788\C,-4.1381357835,2.7585716694,2.9673759161\H,
 -5.1579812303,2.4147168377,3.1741719223\C,-0.0995474827,2.6345132883,-
 0.0476174044\C,0.9455071297,1.7164195775,-0.1873343455\C,2.2926672419,
 2.1001225065,-0.4723187204\C,2.558703778,3.4483819763,-0.6657499207\H,
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 -0.6571408924\H,0.7839495245,6.7474855465,-0.8872017123\H,1.0082027204
 ,7.7998634367,-1.092917887\H,1.798284324,5.8032516275,-0.889462277\H,2
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 C,-5.1126323678,-0.9944206736,-3.0292151972\H,-5.3318232431,-2.3046804
 372,-2.5644701541\H,-5.0403127902,-3.6466576615,-0.8904703152\H,-5.451
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 788629\H,-1.7348499764,0.7905369911,5.6834699971\H,-2.803088998,0.5594
 643122,5.5276926166\H,-1.4973498997,0.687292941,6.7549605809\H,-1.5936
 724284,1.8509620385,5.4006266169\H,-4.2514305334,1.372187681,-2.780680
 954\H,-3.1760422211,1.5471962801,-2.9686362199\H,-3.7234763064,-2.1051
 234165,0.9133792143\H,-5.9947651326,-3.3402812992,-3.4460684157\H,-6.4
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 06\\Dipole=1.9357807,-1.8507178,5.1028764\\Quadrupole=-5.3970991,8.00075
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 \\@"

TRIM/1E Type II

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 .3378371932\\O,-1.3323487876,-0.1477418071,-1.7685757054\\O,0.4699021972
 ,0.6376544369,-0.117007209\\O,1.0960184283,-0.2617512622,-2.5438422821\\O,
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 5,-0.047677137\\C,-2.2773858888,0.0321480603,-0.7756252488\\C,-3.2471058
 074,-0.9997039478,-0.5809999831\\C,-4.2152945273,-0.8056545488,0.395002
 1261\\H,-4.9823699696,-1.5717223468,0.5524632722\\C,-4.2209376288,0.3501
 188667,1.2262881953\\C,-3.225216969,1.3750918933,1.0268646148\\C,-3.1934
 6662,2.4692479859,1.9443982728\\H,-2.4173230832,3.2323805375,1.83842853
 67\\C,-4.1146485351,2.5660998959,2.9772334299\\H,-4.0613104839,3.4096361
 667,3.6742146243\\C,-5.121032943,1.5768782932,3.1444291675\\H,-5.8471699

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TRIM/1E Switching TS

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lvent=ch2cl2,read,externaliteration,dovacuum)\|title\0,1P,0,0.347012
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 2,-2.580411,3.998989\|C,0,-4.606352,-3.03591,2.999637\|H,0,-4.465903,-3.
 996337,3.508803\|C,0,-0.237275,-3.803668,0.687657\|C,0,0.776864,-3.90790
 5,1.676037\|C,0,-0.10901,-4.500115,-0.539959\|C,0,1.906551,-4.701811,1.4
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TRIP/1E Type II

Conf0

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Conf2

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solvent=ch2cl2,read,externaliteration,dovacuum)\|title\0,1\P,0,-0.155
228,-0.579395,-0.967524\|O,0,-1.438658,0.42075,-1.366627\|O,0,0.754101,0
.513805,-0.029774\|O,0,0.513917,-0.942732,-2.258201\|O,0,-0.587418,-1.60
3041,0.077683\|C,0,-1.442528,2.273002,0.141688\|C,0,-2.06485,1.127965,-0
.360351\|C,0,-3.355556,0.682816,0.064064\|C,0,-3.997133,1.417735,1.05174
6\|H,0,-4.999718,1.124407,1.378874\|C,0,-3.376638,2.537694,1.676662\|C,0,
-2.073467,2.970274,1.234824\|C,0,-1.441817,4.038812,1.94066\|H,0,-0.4350
51,4.34979,1.648056\|C,0,-2.074505,4.668043,3.003484\|H,0,-1.563772,5.47
7591,3.536325\|C,0,-3.373777,4.26429,3.412975\|H,0,-3.864271,4.77124,4.2
51036\|C,0,-4.007391,3.217442,2.761794\|H,0,-4.999691,2.880162,3.082335\|C
,0,-0.134269,2.709553,-0.424584\|C,0,0.941952,1.814709,-0.45113\|C,0,2.2
70396,2.212857,-0.799977\|C,0,2.471272,3.521275,-1.218696\|H,0,3.482306,
3.854152,-1.47594\|C,0,1.393297,4.445004,-1.337501\|C,0,0.066365,4.03988
8,-0.942596\|C,0,-1.001988,4.972788,-1.109781\|H,0,-2.020526,4.668775,-0
.851344\|C,0,-0.768606,6.24621,-1.607759\|H,0,-1.605769,6.941729,-1.7335
18\|C,0,0.544994,6.654106,-1.964138\|H,0,0.716416,7.664687,-2.350753\|C,0
,1.601508,5.766428,-1.835488\|H,0,2.616792,6.062514,-2.12326\|C,0,3.3800
33,-4.498677,-0.906033\|C,0,2.9726,-3.150063,-0.884041\|H,0,3.138571,-2.
516376,-1.759657\|C,0,2.373378,-2.594182,0.248702\|H,0,2.128611,-1.52925
7,0.256458\|C,0,2.163234,-3.379228,1.4049\|C,0,2.604879,-4.725187,1.4047
02\|H,0,2.458762,-5.351126,2.290671\|C,0,3.18613,-5.273761,0.26038\|H,0,3
.500474,-6.323492,0.268332\|C,0,1.521772,-2.812329,2.602018\|C,0,2.03259
,-3.190567,3.958784\|H,0,1.852763,-2.390769,4.69428\|H,0,3.109442,-3.409
324,3.902064\|H,0,1.519917,-4.102952,4.316322\|N,0,0.534941,-1.96088,2.4
26616\|H,0,0.175942,-1.785633,1.416753\|C,0,-0.21323,-1.238951,3.405937\|
C,0,-0.522414,0.101418,3.1021\|H,0,-0.166834,0.535963,2.16361\|C,0,-1.27
8039,0.855422,4.00759\|H,0,-1.504575,1.900551,3.778827\|C,0,-1.750818,0.
273151,5.194446\|C,0,-1.46658,-1.072443,5.474595\|H,0,-1.853256,-1.53970
5,6.386283\|C,0,-0.699195,-1.836435,4.584427\|H,0,-0.51618,-2.893487,4.7
88585\|C,0,3.417597,1.259815,-0.647689\|C,0,0.4039479,0.687945,-1.788694\|
C,0,3.906187,0.965319,0.655219\|C,0,5.165107,-0.13767,-1.605589\|C,0,5.0
34339,0.135074,0.784315\|C,0,5.687539,-0.41705,-0.33139\|H,0,5.648612,-0
.57182,-2.488369\|H,0,5.428801,-0.084659,1.783075\|C,0,-3.992199,-0.5131
01,-0.576324\|C,0,-4.176093,-1.71483,0.160919\|C,0,-4.428565,-0.437254,-
1.928834\|C,0,-4.838244,-2.793398,-0.45754\|C,0,-5.069459,-1.548562,-2.5
03418\|C,0,-5.296624,-2.734295,-1.78317\|H,0,-4.992347,-3.719527,0.10895
1\|H,0,-5.404763,-1.482562,-3.545152\|H,0,-2.350516,0.865497,5.893445\|C,

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 095451,-4.368036,-2.94997\H,0,5.072992,-5.421324,-1.883397\C,0,-4.2410
 03,0.821196,-2.778528\H,0,-3.707247,1.569361,-2.169105\C,0,-3.658689,-
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 0324,-1.638714\C,0,-4.802213,-1.777047,2.62324\H,0,-4.405745,-1.853795
 ,3.652485\H,0,-5.342695,-0.818646,2.537555\H,0,-5.536534,-2.591929,2.4
 8001\C,0,-5.123525,-4.542873,-3.547949\H,0,-4.975629,-3.822056,-4.3732
 7\H,0,-4.12733,-4.832203,-3.168357\H,0,-5.606618,-5.444278,-3.968369\C
 ,0,-7.397789,-3.576437,-2.944915\H,0,-7.902256,-4.474182,-3.346977\H,0
 ,-8.028368,-3.161826,-2.138014\H,0,-7.350087,-2.827253,-3.756669\C,0,-
 3.371413,0.540385,-4.021301\H,0,-3.866838,-0.16954,-4.709253\H,0,-3.18
 778,1.475924,-4.581414\H,0,-2.396134,0.118589,-3.725892\C,0,-5.598815,
 1.442687,-3.165234\H,0,-5.449154,2.382607,-3.727831\H,0,-6.189471,0.75
 8859,-3.802499\H,0,-6.200082,1.673453,-2.267273\C,0,6.927798,-1.28999,
 -0.156235\H,0,7.16665,-1.298142,0.924082\C,0,3.533673,0.964387,-3.2059
 14\H,0,2.567179,1.489157,-3.115409\C,0,3.272983,1.598974,1.896123\H,0,
 2.223227,1.833055,1.651385\C,0,6.663771,-2.747211,-0.586875\H,0,7.5638
 43,-3.369667,-0.429419\H,0,5.834836,-3.188119,-0.007521\H,0,6.396891,-
 2.803394,-1.657988\C,0,8.144554,-0.70187,-0.899101\H,0,7.981635,-0.696
 785,-1.992557\H,0,8.342965,0.337318,-0.580402\H,0,9.050899,-1.302042,-
 0.698073\C,0,3.251307,0.665426,3.119475\H,0,2.802298,-0.310245,2.86854
 7\H,0,4.264805,0.481944,3.519907\H,0,2.653982,1.117515,3.931128\C,0,3.
 972332,2.934604,2.230116\H,0,3.929303,3.631498,1.374335\H,0,3.488969,3
 .424127,3.09581\H,0,5.03651,2.76724,2.479369\C,0,4.503952,1.896014,-3.
 961884\H,0,4.661004,2.842682,-3.414765\H,0,5.491228,1.416118,-4.096913
 \H,0,4.10649,2.141742,-4.963936\C,0,3.268215,-0.331829,-3.996825\H,0,2
 .892861,-0.08725,-5.007868\H,0,4.186636,-0.934521,-4.121598\H,0,2.5013
 15,-0.93652,-3.485767\C,0,-2.879221,-3.198276,1.777228\H,0,-3.535891,-
 4.08198,1.677976\H,0,-2.066978,-3.273439,1.036588\H,0,-2.433127,-3.232
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 \State=1-A\HF=-3216.0959114\RMSD=2.239e-09\Di pole=1.8737331,-1.9436175
 ,4.8971892\Quadrupole=-10.7882883,10.2639406,0.5243477,-17.931839,-2.4
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Conf3

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solvent=ch2cl2,read,externaliteration,dovacuum)\title\0,1\P,0,-0.216
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0.534188,-0.142033\O,0,0.375417,-0.823764,-2.417141\O,0,-0.549913,-1.6
1786,-0.045375\C,0,-1.460074,2.267169,0.202987\C,0,-2.103595,1.124232,
-0.277059\C,0,-3.354082,0.65363,0.231815\C,0,-3.929704,1.358872,1.2796
81\H,0,-4.9008,1.043347,1.674288\C,0,-3.278431,2.473956,1.881475\C,0,-
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3.4.321958,1.680605\C,0,-1.90373,4.587637,3.160051\H,0,-1.363613,5.390
35.3.673981\C,0,-3.16536,4.159423,3.65365\H,0,-3.598375,4.64079,4.5372
7\C,0,-3.834796,3.120931,3.025518\H,0,-4.797485,2.76471,3.409404\C,0,-
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39,2.268219,-0.924196\C,0,2.365036,3.584396,-1.332181\H,0,3.360407,3.9
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,-0.936268\C,0,-1.116623,4.993769,-1.031009\H,0,-2.116253,4.675525,-0.
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,-0.116669\C,0,2.379366,-3.349478,0.987916\C,0,2.87546,-4.674507,0.920
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,0,3.721797,-6.217726,-0.324495\C,0,1.826044,-2.833339,2.250151\C,0,2.
473182,-3.222961,3.543793\H,0,2.3237,-2.455353,4.319067\H,0,3.549598,-
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18,2.18778\H,0,0.341234,-1.830817,1.220107\C,0,0.104729,-1.358414,3.25
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-0.436166,-1.686292\C,0,-4.813502,-2.837467,-0.250334\C,0,-5.256883,-1
.543266,-2.229887\C,0,-5.393267,-2.752741,-1.52611\H,0,-4.898937,-3.78
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,0,-4.4845,0.846376,-2.514792\H,0,-3.9087,1.585061,-1.932559\C,0,-3.48

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 ,-4.218806\|H,0,-4.414142,-4.688336,-3.269654\|H,0,-5.993083,-5.325376,-
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 8\|H,0,-3.590011,1.561089,-4.380223\|H,0,-2.719897,0.188066,-3.633076\|C,
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 2317,0.784704,-3.371598\|H,0,-6.402146,1.664206,-1.819788\|C,0,6.691328,
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 79,-3.36809\|H,0,2.514622,1.644512,-3.26601\|C,0,3.261704,1.658368,1.742
 106\|H,0,2.19927,1.876977,1.539959\|C,0,7.952002,-0.747467,0.161711\|H,0,
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 96,-0.344431,1.177055\|C,0,6.320232,-2.664511,0.517729\|H,0,6.110784,-2.
 349191,1.55661\|H,0,5.423923,-3.180719,0.134475\|H,0,7.150208,-3.394065,0
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 96002\|H,0,4.676477,2.894661,-3.522756\|H,0,5.434569,1.446298,-4.241906\|
 H,0,4.092494,2.266147,-5.091823\|C,0,3.120337,-0.185411,-4.195198\|H,0,2
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 4.6541888Quadrupole=-9.4883586,10.703515,-1.2151564,-19.9919649,-1.92
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TRIP/1E Type I

Conf0

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14,-0.396129\|C,0,-1.312229,2.256433,0.304329\|C,0,-1.988344,1.187587,-0  

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\|\H,0,-4.878244,1.113224,1.525644\|C,0,-3.198893,2.438397,1.913361\|C,0,-  

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77,-0.625037\|C,0,-0.833204,5.063391,-0.669616\|H,0,-1.853229,4.756108,-  

0.420771\|C,0,-0.579725,6.376061,-1.040006\|H,0,-1.402474,7.098707,-1.07  

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84247,-0.672644,4.67252\|C,0,-0.968356,-1.891613,5.160822\|H,0,-1.320245  

,-2.279616,6.123\|C,0,-0.012532,-2.614593,4.439549\|H,0,0.362478,-3.5587  

86,4.845938\|C,0,0.428027,-2.154778,3.17601\|C,0,-0.078816,-0.926491,2.6  

8633\|H,0,0.274329,-0.519187,1.736303\|C,0,-1.006796,-0.200796,3.429945\|  

H,0,-1.371773,0.752816,3.037219\|C,0,1.403641,-2.937189,2.402355\|C,0,2.  

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2,1.090101\|H,0,0.606212,-2.27416,0.599507\|C,0,2.20945,-3.520338,0.1374  

94\|C,0,2.717731,-2.750206,-0.925311\|H,0,2.5404,-1.672167,-0.960912\|C,0  

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Conf1

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Conf2

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 .944829,2.377967,2.577633\H,0,3.921031,3.151992,1.790715\H,0,3.402426,
 2.768073,3.458622\H,0,4.999877,2.216261,2.866273\C,0,4.79577,1.912476,
 -3.656813\H,0,4.901606,2.813288,-3.027145\C,0,5.800162,1.465412,-3.778
 703\H,0,4.443054,2.232752,-4.654588\C,0,3.616055,-0.321392,-3.9526\H,0
 ,3.238195,0.008207,-4.937984\H,0,4.563033,-0.864439,-4.12729\H,0,2.873
 659,-1.011813,-3.520984\C,0,-3.098062,-3.029848,1.322043\H,0,-3.776786
 ,-3.879828,1.123778\H,0,-2.245072,-3.086189,0.626422\H,0,-2.715084,-3.
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Conf3

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 solvent=ch2cl2,read,externaliteration,dovacuum)\|title\0,1\P,0,-0.090
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 0.400252,-0.103878\O,0,0.584213,-0.744615,2.498014\O,0,-0.548967,-1.6
 78426,-0.266704\C,0,-1.263845,2.268832,0.181636\C,0,-1.942186,1.19706,
 -0.404913\C,0,-3.267009,0.808618,-0.031728\C,0,-3.891736,1.528175,0.97
 8073\H,0,-4.917402,1.274986,1.265318\C,0,-3.223964,2.573342,1.678608\C
 ,0,-1.880435,2.939244,1.300527\C,0,-1.201052,3.916347,2.089683\H,0,-0.
 165787,4.172117,1.847347\C,0,-1.825543,4.52711,3.167591\H,0,-1.278623,
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 776\H,0,0.078615,2.667111,-0.330283\C,0,1.105922,1.719156,-0.405246\C,
 0,2.462225,2.067911,-0.689656\C,0,2.743837,3.391516,-1.001459\H,0,0.377
 8809,3.688438,-1.200747\C,0,1.718256,4.376559,-1.085267\C,0,0.362284.
 017047,-0.750645\C,0,-0.652987,5.012094,-0.886043\H,0,-1.692441,4.7449
 43,-0.674527\C,0,-0.342615,6.299489,-1.299041\H,0,-1.140629,7.042724,-
 1.404642\C,0,0.999586,6.66072,-1.593823\H,0,1.23242,7.682644,-1.912531
 \C,0,2.006985,5.714345,-1.491409\H,0,3.043903,5.974773,-1.732839\C,0,-
 1.762359,-0.333249,4.717655\C,0,-1.343062,-1.555447,5.284581\H,0,-1.76
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 -3.306132,5.109864\C,0,0.135683,-1.983554,3.390664\C,0,-0.272199,-0.75
 2068,2.82254\H,0,0.151361,-0.410648,1.874398\C,0,-1.19352,0.056438,3.4
 85071\H,0,-1.48287,1.010085,3.034723\C,0,1.101822,-2.852306,2.702727\C
 ,0,2.031586,-3.714957,3.501162\H,0,2.937013,-3.967451,2.928421\H,0,1.5
 26347,-4.659247,3.775681\H,0,2.310341,-3.200263,4.433238\N,0,1.14428,-
 2.821242,1.389902\H,0,0.426483,-2.234905,0.831551\C,0,2.00202,-3.55293
 6,0.508707\C,0,2.642569,-2.830492,-0.515509\H,0,2.548722,-1.741904,-0.
 563241\C,0,3.417927,-3.518772,-1.45581\H,0,3.919458,-2.955712,-2.24782
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 ,3.004738,-6.715642,-0.297006\C,0,2.123765,-4.951909,0.589082\H,0,1.58
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 .542321,-1.743659\C,0,4.018945,0.660739,0.710352\C,0,5.316326,-0.29324
 4,-1.597693\C,0,5.144645,-0.177808,0.802031\C,0,5.826595,-0.643574,-0.
 33592\H,0,5.811885,-0.671926,-2.499058\H,0,5.519143,-0.467672,1.790306
 \C,0,-3.954991,-0.322138,-0.734553\C,0,-4.21732,-1.538027,-0.046037\C,
 0,-4.36713,-0.169834,-2.087684\C,0,-4.924413,-2.55639,-0.714403\C,0,-5
 .060202,-1.221271,-2.712287\C,0,-5.359786,-2.420721,-2.042182\H,0,-5.1
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 1,-5.445613,-2.11647\C,0,-2.797231,0.536664,5.385513\H,0,-3.722285,0.5
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 5.470161\C,0,-4.104441,1.112656,-2.879673\H,0,-3.533579,1.801353,-2.23
 4774\C,0,-3.737665,-1.77886,1.386639\C,0,-6.116747,-3.548967,-2.737656
 \H,0,-6.264066,-4.349022,-1.987874\C,0,-4.901059,-1.697252,2.396646\H,
 0,-4.531413,-1.847277,3.427728\H,0,-5.407726,-0.717364,2.35355\H,0,-5.
 657863,-2.477592,2.192205\C,0,-5.29617,-4.144183,-3.900844\H,0,-5.1266

4,-3.392854,-4.694167\H,0,-4.309163,-4.497647,-3.552858\H,0,-5.827996,
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 .941874,-3.660553\H,0,-8.109496,-2.690614,-2.382487\H,0,-7.430836,-2.3
 06896,-3.990304\C,0,-3.244823,0.841921,-4.131346\H,0,-3.772722,0.18988
 8,-4.851645\H,0,-3.00992,1.790372,-4.64894\H,0,-2.293696,0.35761,-3.85
 3145\C,0,-5.424222,1.8252,-3.241049\H,0,-5.220923,2.778282,-3.763266\H
 ,0,-6.049952,1.20279,-3.907052\H,0,-6.014662,2.050806,-2.334535\C,0,7.
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 .893402,-3.144462\H,0,2.695478,1.364794,-3.026066\C,0,3.33999,1.183056
 ,1.979426\H,0,2.27892,1.359501,1.733968\C,0,6.874788,-2.936232,-0.6945
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 ,0,6.636819,-2.950855,-1.773916\C,0,8.295183,-0.836922,-0.888572\H,0,8
 .144643,-0.782521,-1.982411\H,0,8.461818,0.19017,-0.51715\H,0,9.215266,
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 ,0,3.942962,2.53816,2.408664\H,0,3.866591,3.284781,1.598929\H,0,3.4125
 66,2.936519,3.293234\H,0,5.011556,2.42547,2.669511\C,0,4.620256,1.9105
 94,-3.83253\H,0,4.728251,2.830525,-3.231306\H,0,5.629587,1.482889,-3.9
 80399\H,0,4.223281,2.195326,-4.824435\C,0,3.483026,-0.356833,-4.0224
 ,0,0.3.068507,-0.063449,-5.004492\H,0,4.434445,-0.887215,-4.211203\H,0,2
 .767559,-1.046755,-3.547146\C,0,-2.978977,-3.113494,1.530161\H,0,-3.64
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 8,-5.0747244,-15.696795|PG=C01 [X(C65H72N1O4P1)]\\@

TRIP/1E Type III.I

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 solvent=ch2cl2,read,externaliteration,dovacuum)\\title\\0,1\H,0,-5.712
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 ,0.457582\C,0,-3.725315,2.370052,-1.825077\C,0,-5.030883,-0.626222,1.0
 298571\C,0,-6.953155,-1.306385,-0.545022\C,0,-2.42593,2.147774,0.85695
 \C,0,-3.930427,0.184386,1.371492\C,0,-4.860968,3.387137,-2.062421\C,0,-
 3.243194,1.746094,-3.149402\H,0,-2.875635,2.930265,-1.400273\H,0,-5.17
 4929,3.86192,-1.115281\H,0,-5.748612,2.903864,-2.510859\H,0,-4.528172,
 4.185379,-2.751306\H,0,-2.89849,2.536293,-3.841821\H,0,-4.055396,1.190
 618,-3.653172\H,0,-2.402716,1.055504,-2.970615\H,0,-5.366268,-1.37636,
 1.752671\C,0,-8.201823,-0.395391,-0.504575\C,0,-7.168786,-2.549115,0.33
 21491\H,0,-6.827527,-1.64689,-1.590917\H,0,-8.090667,0.471759,-1.179232
 \H,0,-8.368168,-0.01002,0.518567\H,0,-9.105163,-0.954128,-0.811629\H,0
 ,-7.416856,-2.272411,1.373458\H,0,-6.273674,-3.194138,0.356186\H,0,-8.
 010507,-3.147846,-0.059391\C,0,-1.137088,2.130885,0.237809\C,0,-2.65269
 4,3.093675,1.849275\C,0,-3.242342,-0.042412,2.720695\O,0,-0.927382,1.2
 4762,-0.802506\C,0,-0.10798,3.008405,0.59368\H,0,-3.640855,3.153937,2.
 316966\C,0,-1.626806,3.968707,2.306727\C,0,-2.83326,-1.515782,2.922745
 \C,0,-4.124488,0.444851,3.889406\H,0,-2.313553,0.552578,2.725203\H,0,-
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 786,-2.183394,2.979935\H,0,-3.600558,0.312914,4.854166\H,0,-4.383492,1
 .513091,3.785832\H,0,-5.069366,-0.128315,3.936357\P,0,0.024963,-0.1127
 46,-0.558885\C,0,1.18566,2.95481,-0.143659\C,0,-0.321382,3.912207,1.69
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 0,0.442926,-0.593453,-1.923892\O,0,1.291191,0.597333,0.290182\C,0,1.86
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 .737011,2.23804\H,0,-2.860187,4.920213,3.824412\C,0,-0.841782,5.690978
 ,3.859415\C,0,3.183948,1.626376,-0.78989\C,0,1.076792,5.365092,-0.8697
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 01\H,0,1.268285,6.22248,3.691318\H,0,4.795324,2.724908,-1.694251\C,0,4
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 83,7.408177,-1.544071\C,0,2.99088,6.377503,-1.992591\H,0,4.688617,5.08
 904,-2.348807\C,0,5.19731,-1.335516,0.499726\C,0,4.307139,0.670098,1.7
 72286\C,0,4.852842,-1.629645,-1.865876\C,0,3.507492,0.000032,-3.276822
 \H,0,3.453025,7.255511,-2.456998\H,0,5.62096,-1.679258,1.450826\C,0,5.
 41594,-2.093282,-0.665503\C,0,5.6759,1.131322,2.31387\C,0,3.508777,-0.
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 11557\C,0,6.224673,1.721477,1.5579\H,0,6.309492,0.270971,2.598242\H,0,
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 1,-0.388831,2.458192\C,0,2.825261,-1.15497,-4.035107\H,0,2.72548,0.746581,-3
 .056585\H,0,4.137943,1.013369,-5.115793\H,0,5.037698,1.541057,-3.66456
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 524,1.007736\|H,0,8.18077,-2.365061,-0.601885\|H,0,8.241614,-4.090645,-0
 .133315\|H,0,5.35467,-4.2542,1.172832\|H,0,6.073209,-5.452389,0.060903\|H
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 .991379\|H,0,-2.324491,-6.132302,0.562205\|H,0,-2.058468,-5.886202,-1.18
 386\|H,0,-0.773756,-6.629507,-0.163137\|C,0,0.299726,-5.29013,2.099502\|C
 ,0,1.206082,-3.475074,0.750955\|H,0,-0.477895,-6.033995,2.298488\|C,0,1.
 387444,-5.181516,2.967819\|H,0,1.171075,-2.834609,-0.133895\|C,0,2.30350
 2,-3.397637,1.610735\|H,0,1.450188,-5.847668,3.835617\|C,0,2.407197,-4.2
 29581,2.745069\|H,0,0.3098599,-2.680581,1.389268\|C,0,3.565382,-4.10036,3
 .703011\|H,0,0.391352,-5.088827,4.050079\|H,0,3.255882,-3.530575,4.600165
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 Quadrupole=-5.0187217,26.9891532,-21.9704314,14.9074858,11.1982848,-7.
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TRIP/1E Type III.2

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solvent=ch2cl2,read,externaliteration,dovacuum)\|title\|0,1\P,0,0.1771
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672,0.424385\|O,0,0.540274,-0.785479,-1.710831\|O,0,-0.597575,-0.909954,  

0.68035\|C,0,0.412979,3.042288,0.519768\|C,0,-0.709295,2.264679,0.215221  

\|C,0,-1.992989,2.476696,0.808751\|C,0,-2.120315,3.519287,1.717083\|H,0,-  

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4.051997,1.542681\|C,0,1.407269,4.797432,2.035136\|H,0,2.406299,4.584585  

,1.644291\|C,0,1.24312,5.774,3.006635\|H,0,2.115023,6.328554,3.370867\|C,  

0,-0.043506,6.056097,3.53898\|H,0,-0.160304,6.834836,4.300557\|C,0,-1.14  

1657,5.329168,3.107756\|H,0,-2.135958,5.518492,3.528317\|C,0,1.699663,2.  

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,-0.696914\|C,0,4.257224,2.181195,-1.279331\|H,0,5.263705,1.973373,-1.65  

6693\|C,0,3.732498,3.498837,-1.41974\|C,0,2.429167,3.807377,-0.883509\|C,  

0,1.904896,5.118114,-1.097955\|H,0,0.901657,5.357225,-0.733689\|C,0,2.64  

0261,6.08286,-1.770959\|H,0,2.212132,7.078962,-1.928479\|C,0,3.939626,5.  

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TRIP/1E Switching TS

Conf0

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Conf1

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Conf2

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Conf3

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TRIP/2E Type I

Conf0

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Conf1

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Conf2

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Conf3

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TRIP/2E Type II

Conf0

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Conf1

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Conf2

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Conf3

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TRIP/2E Switching TS

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Conf2

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Conf3

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 9\|C,0,4.714477,-2.641136,3.102026\|F,0,5.297125,-3.658948,3.779686\|F,0,
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 64L-G09RevD.01State=1\|HF=-3513.6052377\|RMSD=4.196e-09\|Dipole=-0.677
 3552,-5.204177,-1.1071978\|Quadrupole=-20.2746366,41.2347341,-20.960097
 5,0.1185029,1.4294712,8.9734658\|PG=C01 [X(C65H69F3N1O4P1)]\\@"

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