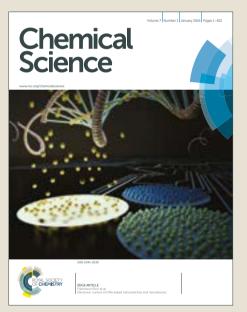
View Article Online View Journal

Chemical Science

Accepted Manuscript

This article can be cited before page numbers have been issued, to do this please use: M. Melikian, J. Gramueller, J. Hioe, J. Greindl and R. M. Gschwind, *Chem. Sci.*, 2019, DOI: 10.1039/C9SC01044K.



This is an Accepted Manuscript, which has been through the Royal Society of Chemistry peer review process and has been accepted for publication.

Accepted Manuscripts are published online shortly after acceptance, before technical editing, formatting and proof reading. Using this free service, authors can make their results available to the community, in citable form, before we publish the edited article. We will replace this Accepted Manuscript with the edited and formatted Advance Article as soon as it is available.

You can find more information about Accepted Manuscripts in the **author guidelines**.

Please note that technical editing may introduce minor changes to the text and/or graphics, which may alter content. The journal's standard <u>Terms & Conditions</u> and the ethical guidelines, outlined in our <u>author and reviewer resource centre</u>, still apply. In no event shall the Royal Society of Chemistry be held responsible for any errors or omissions in this Accepted Manuscript or any consequences arising from the use of any information it contains.



rsc.li/chemical-science

Chemical Science



ARTICLE

Received 00th January 20xx, Accepted 00th January 20xx

DOI: 10.1039/x0xx00000x

www.rsc.org/

Brønsted Acid Catalysis – The Effect of 3,3'-Substituents on the Structural Space and the Stabilization of Imine/Phosphoric Acid Complexes

Maxime Melikian, + Johannes Gramüller, + Johnny Hioe, Julian Greindl and Ruth M. Gschwinda*

BINOL derived chiral phosphoric acids (CPAs) are widely known for their high selectivity. Numerous 3,3' substituents are used for a variety of stereoselective reactions and theoretical models of their effects are provided. However, experimental data about the structural space of CPA complexes in solution is extremely rare and so far restricted to NMR investigations of binary TRIP/imine complexes featuring two *E*- and two *Z*-imine conformations. Therefore, in this paper the structural space of 16 CPA/imine binary complexes is screened and 8 of them are investigated in detail by NMR. For the first time dimers of CPA/imine complexes in solution were experimentally identified, which show an imine position similar to the transition state in transfer hydrogenations. Furthermore, our experimental and computational data revealed an astonishing invariance of the four core structures regardless of the different steric and electronic properties of the 3,3'-substituent. However, a significant variation of *E*/*Z*-ratios is observed, demonstrating a strong influence of the 3,3'-substituents on the stabilization of the imine in the complexes. These experimental *E*/*Z*-ratios cannot be reproduced by calculations commonly applied for mechanistic studies, despite extensive conformational scans and treatment of the electronic structure at a high level of theory with various implicit solvent corrections. Thus, these first detailed experimental data about the structural space and influence of the 3,3' substituent on the energetics of CPA/imine complexes can serve as basis to validate and improve theoretical predictive models.

Introduction

In the field of Brønsted acid catalysis^{1,2} a variety of different catalyst classes have emerged in recent years, one of the most famous being BINOL (1,1'-bi-2-naphthol) derived chiral phosphoric acids (CPAs).^{3,4} CPAs have been successfully applied in various enantioselective syntheses such as transfer hydrogenations (see Figure 1a),^{5,6} reductive aminations,^{7,8} Mannich type reactions,^{9–11} Strecker reactions,¹² and enantioselective additions.^{13,14} The main difference between these BINOL derived CPAs are the 3,3' substituents, which significantly influence not only yields, but especially the stereoselectivity of these reactions. Typical 3,3' substituents successfully employed are e.g. 2,4,6-triisopropylphenyl groups in the Brønsted acid catalyst TRIP, 3,5-bis(trifluoromethyl) groups in TRIFP, or triphenylsilyl groups in TiPSY (see Figure 1b).^{5,6,15} When the bulkiness of the 3,3' substituents is increased, the cited transfer hydrogenations and the reductive amination generally show an increase in enantiomeric excess, but also a sudden drop in yield and enantioselectivity when using extended aromatic systems such as naphthyl, or

^{a.} Institut für Organische Chemie, Universität Regensburg, D-93053 Regensburg, Germany 4-biphenyl groups.^{5–7} The reason for these differences is for most of these reactions largely unexplored and vary from reaction to reaction.¹⁶ As a result many synthetic methods using BINOL derived CPAs had to be developed via trial and error or screening methods.^{17,18}

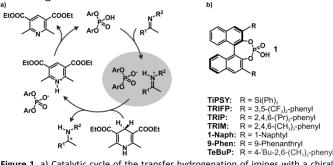


Figure 1. a) Catalytic cycle of the transfer hydrogenation of imines with a chiral phosphoric acid catalyst and a Hantzsch ester as reducing agent. The binary complexed highlighted in grey is the focus of this work; b) Brønsted acid catalysts with different 3,3' groups.

To shed light on the involved structures and to reveal the key interactions of reactivity and stereoselectivity so far mainly theoretical calculations and multivariant linear regression models have been applied.^{19,20} Thus, Goodman *et al.* linked the enantioselectivity to structural parameters of CPAs, such as the rotational barrier of the 3,3'-substituent describing the steric bulk close to the hydrogen bond and the cone angle AREA(Θ) of the substrate binding pocket, reflecting remote steric bulk.²¹ Furthermore, combining that with a steric classification of

⁺ Authors contributed equally to this work

Electronic Supplementary Information (ESI) available: [details of any supplementary information available should be included here]. See

DOI: 10.1039/x0xx00000x

Accepted Manus

Journal Name

ARTICLE

electrophiles and nucleophiles Goodman et al. developed a webtool for predicting suitable CPA catalysts based on reactant structures.^{22–24} Parallel to that, Sigman et al. demonstrated how a data-driven approach can be capable of revealing nonintuitive insights about interactions involved in stereoselectivity determination by analysing the dependence of ee on steric and electronic molecular descriptors.20,25-27

In contrast to the theoretical studies the information about the structures of CPAs/imine complexes based on experimental data is rather limited so far. To our knowledge, only two crystal structures were provided from the groups of MacMillan⁷ and Schneider²⁸ with strongly deviating structural features (see Figure 2 for structures and Figure 1 for substituents). While a TiPSY/imine complex adopts core structure Type II E proposed by theoretical calculations (see Figure 2a and c)²⁹ the second crystal structure of a TeBuP/imine complex exhibits an intermediate position of structures Type I E and Type II E (see Figure 2b and c).

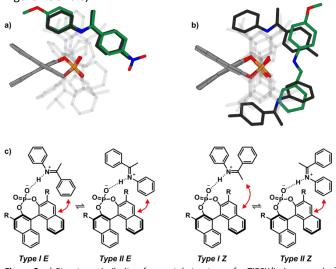


Figure 2. a) Structure similarity of a crystal structure of a TiPSY/imine complex⁷ (green) and structure *Type II E* of a TRIP/imine complex identified in our previous work²⁹ (black). b) structure deviation of the crystal structure of a TeBuP/imine complex²⁸ (green) vs both *Type I E* (black above) and *Type II E* (black below) of a TRIP/imine complex. The imine shows an intermediate position of *Type I E* and *Type II E*. c) The four core structures of the binary complex identified in our previous work.²⁹ The red arrows mark some of the identified NOE interactions.

In solution the experimental data base is similarly sparse. While two remarkable studies provided some NMR data about a ternary CPA complex³⁰ and a CPA reaction intermediate³¹ the first comprehensive insight about the structural space of CPA imine complexes and their hydrogen bond properties in solution was provided by our group on the example of TRIP/imine complexes.^{29,32} In depth NMR investigations revealed ionic complexes with extremely strong and highly covalent hydrogen bonds $^{\rm 32}$ and a structural space covering the four core structures (Type I/II E/Z, shown in Figure 2c),²⁹ which already predicted were previously for ternary CPA/imine/nucleophile complexes by theoretical calculations.³³ However, the effect of varying 3,3'-substituents on the structure of CPA/imine complexes as well as on their population has so far remained elusive.

Therefore, in this report we present the first experimental data about the influence of different 3,3'-substituents on the structures and E/Z populations of complexes between chiral phosphoric acids and imines. A screening covering, 16 binary complexes with 5 different chiral phosphonic adids and Finances and in depths structural investigations on selected examples reveal an astonishing invariance of the four main structures Type I E, Type II E, Type I Z and Type II Z. Moreover, for the first time dimeric CPA/imine complexes in solution were characterised, hence extending the structural space of those complexes in solution and forming a bridge to the dimeric crystal structure. In addition, comprehensive data about the E/Z population of these complexes are provided, which deviate significantly from the calculated values (e.g. for TRIP/5: experimental $\Delta G_{rel E/Z}$ = 2.0 kJ/mol; theoretical $\Delta G_{rel E/Z}$ = 9.3 kJ/mol).²⁹ This indicates a strong influence of refined dispersion interactions and/or solvent interactions not reflected by the commonly applied solvent models and offers the possibility to validate energetic results for theoretical prediction models in ion pairing catalysis.

RESULTS AND DISCUSSION

Investigated complexes and their NMR properties.

In order to investigate the influence of different 3,3'substituents of CPAs as well as different substituents of the imines on the structures of CPA/imine complexes and their E/Zpopulations, we selected several CPA catalysts and imines (see Figure 3) used in synthesis³⁴ and screened their NMR properties. Dichloromethane was used as solvent, since it gave small signal linewidths and the highest achievable signal dispersion at low temperatures.²⁹ Furthermore, temperatures between 170 K and 200 K were needed to sufficiently slow down exchange processes in solution and most of the structural investigations were performed at 180 K unless otherwise noted.

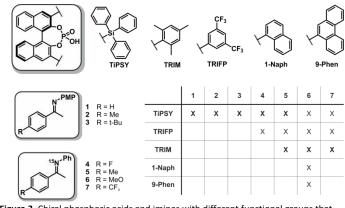


Figure 3. Chiral phosphoric acids and imines with different functional groups that were used for the NMR-spectroscopic investigations of CPA/imine complexes (PMP = p-methoxyphenyl). All screened CPA/imine combinations are shown in the table. A bold X marks complexes, which could be investigated in detail. For all table. A bold X marks complexes, which could systems, a 1:1 ratio of CPA and imine was used.

In principle, CPA/imine complexes with TiPSY and TRIFP produced basic signal pattern very similar to TRIP (see Figure 4). The spectra showed highly overcrowded aromatic regions and two well separated hydrogen bond signals for the E and Z complexes. Thus, quantification of E and Z complexes is

Journal Name

straightforward by integration of the H-bond protons. The access to a detailed structural analysis however depended individually on the overlap of key signals and their linewidths. For TiPSY complexes a structural analysis could be completed despite the significantly higher signal overlap in the aromatic region compared to TRIP. For TRIFP complexes line width factors and chemical shift overlap of key signals prevented any further structural analysis.

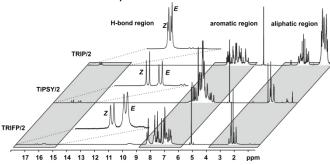


Figure 4. Spectral resolution of the ¹H spectra of complexes between catalysts TRIP, TiPSY and TRIFP and imine **2** at 180 K in CD_2Cl_2 ; in all cases well separated signals of the hydrogen bonds indicate the E/Z populations, while deviations in linewidths and chemical shift overlap of key signals allow the structural investigations via NOE analysis only for complexes with TIPSY.

For complexes with TRIM, 9-Phen, and 1-Naph as catalyst additional signals in the hydrogen bond region appeared at slightly lower chemical shifts (12.0 - 14.0 ppm) indicating an extended structural space for these complexes (see Figure 5). The detailed structural investigations of TRIM complexes (see below,) and dilution experiments for TRIM, 9-phen and 1-naph (see Supplementary S1-3) identified these species as dimers, most probably enabled by attractive interactions between the 3,3'- substituents of TRIM, 9-Phen, and 1-Naph and the imines. For 9-Phen and even more pronounced for 1-Naph, a plethora of additional signals appeared in the hydrogen bond region of the ¹H spectrum at 180 K. Given the asymmetry of the 3,3'substituents of these catalysts and a rotational barrier of \approx 14 kcal/mol,²¹ at least two slow exchanging rotational isomers of these catalyst are expected at 180 K, causing a signal splitting of the E and Z complexes, hence revealing the whole conformational space of these binary complexes.

Based on this NMR screening and initial 2D assignments, binary complexes with TiPSY and imines **1**, **2**, **3**, **4** and **5** as well as TRIM and imines **5**, **6**, **7** were selected for the in detail structural investigations described in the following.

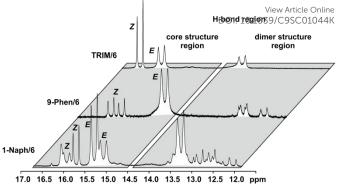


Figure 5 ¹H spectra of the H-bond region of catalysts TRIM, 9-Phen and 1-Naph at 180 K in CD₂Cl₂ reveal the extended structural space of these catalysts including dimeric complexes and additional conformations due to asymmetric 3,3 substituents in 9-Phen and 1-Naph.

Structural space of TiPSY/imine complexes

For complexes providing such degree of signal overlap as shown in the aromatic region in Figure 4, there are quite a few NMR spectroscopic techniques that are typically used to improve signal dispersion. These are mainly 3D experiments such as 3D-¹⁵N NOESY HSQC³⁵ or 3D ¹³C NOESY HSQC³⁶ or pure shift techniques.^{37–39} Especially the multidimensional approach^{40–42} is broadly applied in biochemistry and requires the use of specific labelling strategies.^{43–45} Due to the limited possibilities to include isotope labelling via different synthetic strategies into the catalyst or the imine, the chemical exchange even at low temperatures²⁹ and the lowered solubility at the required temperatures, all of these more advanced techniques proved to be too insensitive to provide structural information in our case. Instead we focused mainly on the most sensitive one- and twodimensional NMR methods (1H,1H / 1H,13C 2D spectra for assignments and selective 1D NOESY⁴⁶⁻⁴⁸/ 2D NOESY/ ¹H,¹⁹F HOESY^{49,50} spectra for structural investigations).

Table 1. Experimental E/Z ratios of the investigated CPA/imine complexes in CD₂Cl₂ at 180 K. Due to linewidth and signal overlap, a general error of 5% is expected.

	1	2	3	4	5	6	7
TRIP	n.d.	62:38	n.d.	n.d.	67:33	77:23	86:14
TiPSY	45:55	53:47	51:49	68:32	56:44	70:30	71:29
TRIFP	n.d.	n.d.	n.d.	83:17	70:30	81:19	69:30
TRIM	n.d.	n.d.	n.d.	n.d.	47:53	49:51	55:45

In TiPSY/imine mixtures (1:1) at 180 K varying *E/Z* complex ratios were found with a trend to higher Z amounts compared to TRIP/imine and TRIFP/imine complexes (see Table 1). A similar trend can be observed for TRIM/imine complexes. In TiPSY/4, the *E/Z* ratio reached 68:32, in TiPSY/2 53:47 and in TiPSY/3 51:49. For TiPSY/1 even a 45:55 *E/Z* ratio was found, a rare example with higher *Z*-imine concentration (see Supplementary S4-5 for spectra). A complete assignment of the TiPSY/imine complexes was done for imines 1, 2, 3 and 4 at 180 K (for spectra see Supplementary S6-S28). Despite a severe signal overlap in the crowded aromatic region, most of the signals were unambiguously assigned.

ARTICLE

The assignment of the *E Type* core structures of the TiPSY/imine complexes (*Type I E*, *Type II E*) is shown exemplarily on imine **4**. With the help of ¹⁹F spectroscopy the spectral resolution can be vastly improved by reducing the complexity of the obtained spectra.^{51–53} Orientation *Type I E* was identified in a 2D ¹H ¹⁹F HOESY experiment. A total of four HOE cross-peaks were found between the fluorine atom of imine **4** and the BINOL backbone of the catalyst (Figure 6).

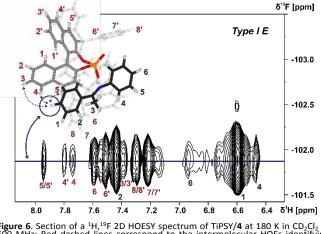
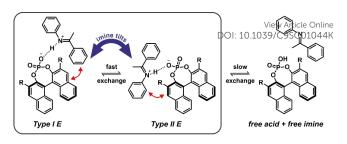


Figure 6. Section of a ¹H,¹⁹F 2D HOESY spectrum of TiPSY/4 at 180 K in CD₂Cl₂ at 600 MHz; Red dashed lines correspond to the intermolecular HOEs identifying complex structure *Type I E* (for detailed NMR parameters see Supplementary S19).

The presence of these HOE cross-peaks between imine 4 and the protons 3, 4 and 5 of the TIPSY backbone in combination with the hydrogen bond confirm the existence of the Type I E structure. Due to a dissociation, rotation and re-association of the TiPSY/imine complexes (Figure 7), HOEs are also observed between the fluorine atom of **4** and the opposite naphthyl moiety of the catalyst (Figure 6, HOE to proton 4'). This exchange is slow on the NMR timescale compared to the tilting mechanism that leads to an exchange between structures Type I E and Type II E (Figure 7). A rotation of the E-imine inside the complex can be excluded, since the rotational barrier for this process would be by far too high due to the steric hindrance of imine inside the complex. Orientation Type II E in the TiPSY/4 complex was similarly identified by using selective 1D NOESY experiments with saturation on proton 6 (see Supplementary S20) Similar NOE patterns for conformations Type I E and Type II E were observed for TiPSY complexes with the imines 1, 2 and 3 (Supplementary S14-S18).

In contrast to the TiPSY/*E*-imine structures, only one set of signals is observed for the BINOL backbone of the catalyst in the TiPSY *Z*-imine complexes. This can be explained by the reduced steric hindrance of the *Z*-imine enabling an exchange between orientation *Type I Z* and *Type II Z* via rotation of the imine around the hydrogen bond (Figure 7). This exchange pathway in addition to the exchange via tilting of the imine (see TiPSY/*E*-imine), results in only one average set of ¹H signals for the BINOL backbone of the catalyst.²⁹



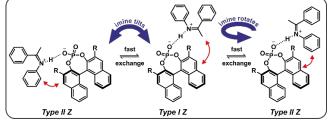
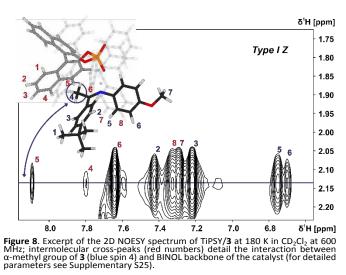


Figure 7. Complex exchange processes in TiPSY/imine complexes; an exchange via tilting of the imine is observed between structures *Type I E* and *Type II E* as well as a disassociation and association process of the TiPSY/E-imine complexes; the exchange via tilting is fast on the NMR time-scale and leads to a different interaction pattern for each half of the catalyst; the dissociation and association process is slow on the NMR time-scale and leads to each path time-scale and leads to each process the reduced steric hindrance of the Z-imine enables an additional fast exchange between structures *Type I Z* and *Type II Z* via rotation, leading to a different signal pattern.

Both orientations *Type I Z* and *Type II Z* were detected for the TiPSY/*Z*-imine complexes. Complex TiPSY/**3** is used exemplarily for the assignment of *Type I Z* (Figure 8). In the ¹H NOESY spectrum, two specific intermolecular NOEs between the α -methyl group of the imine and the BINOL backbone of the catalyst were found. This interaction, in combination with the strong hydrogen bond between catalyst and imine shows the existence of orientation *Type I Z* in solution. Similarly, orientation *Type II Z* could be confirmed in selective 1D NOESY spectra (Supplementary S26) and the same NOE pattern were observed for all investigated complexes TiPSY/**1,2,3,4** (Supplementary S21-S28).

Thus, despite significantly deviating 3,3' substituent/imine interactions causing different E/Z ratios in TiPSY/imine complexes (see Table 1), for all TiPSY/imine complexes studied in detail the structures *Type I E* and *Type II E* as well as *Type I Z* and *Type II Z* could be elucidated by NMR.



Journal Name

This article is licensed under a Creative Commons Attribution-NonCommercial 3.0 Unported Licence

Open Access Article. Published on 08 April 2019. Downloaded on 4/25/2019 3:32:10 PM.

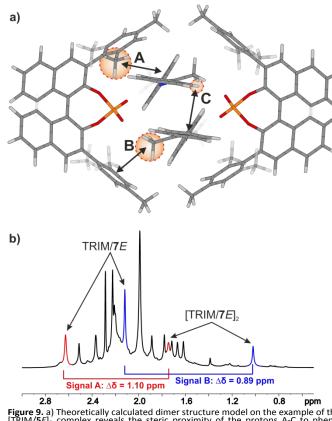
ARTICLE

Journal Name

Structural space of TRIM/imine complexes and dimerization trends of CPAs

For TRIM complexes with imines 5, 6, and 7 the same four core structures Type I/II E/Z were experimentally identified by NOE analysis (for assignment and spectra see Supplementary S33-42). In contrast to TRIP and TiPSY only one set of catalyst signals for both Type I/II E structures even at 180 K is observed, which can be explained by an additional exchange pathway. This pathway is potentially enabled via facilitated rotation like it is the case for Type I/II Z and/or an enhanced dissociationassociation process of the complex. (for detailed explanation see Supplementary S43).²¹

Furthermore, the additional hydrogen bonded signals as discussed above (see Figure 5) were identified as [TRIM/Eimine]₂ dimers and were in detail investigated on complex TRIM/7.^{± 1} Chemical exchange between TRIM/7E and [TRIM/7E]₂ was identified by EXSY signals and dilution of the sample lead to a complete shift of the monomer-dimer equilibrium towards the monomers (see Supplementary S1-3). In addition, DOSY measurements corroborated the assignment as a dimer (see Supplementary S44). Theoretical calculations supported by distinct changes of the measured chemical shifts (see Figure 9) confirm the observed dimer structures to be similar to the reported crystal structure of Schneider²⁸ (see Figure 11 below).



1.8 1.8 1.9 1.8 1.2 0.8 ppm Figure 9. a) Theoretically calculated dimer structure model on the example of the [TRIM/5*E*]₂ complex reveals the steric proximity of the protons A-C to phenyl entities which causes paratropic shielding effects and induces severe highfield shifts. b) Excerpt of the ¹H spectrum of TRIM/7 at 180 K in CD₂Cl₂ at 600 MHz highlighting the highfield shift of proton A (red) and B (blue) of the dimer compared to the monomer. The significant highfield shifts of A-C compared to the monomeric structure (At 1.10 ppm, B: 0.89 ppm, C: 1.56 ppm) experimentally corroborate the computed structure and indicate CH- π and π - π interactions.

The 3,3'-substituent dependent formation of CPA/imine dimers gave rise to the question, which 3,3'-substituent/properties enable or restrict dimer formation. Remarkably, all catalysts forming dimers possess medium rotational barriers of the 3,3'substituent (13.63-21.58 kcal/mol) as described by Goodman.²¹ This hints at the influence of putative van der Waals interactions between the catalyst 3,3'-substituent and the imine on the energetic interaction profile supporting dimer formation. The presence of such interactions is indicated for $[TRIM/7E]_2$ by a strong highfield shift of proton A (Figure 9), caused by steric proximity to a phenyl entity of the imine. For TRIFP and TiPSY a low rotational barrier (2.02 and 1.35 kcal/mol respectively)²¹ points out the lack of moieties capable of such interactions, while the high rotational barrier of TRIP (28.40 kcal/mol)²¹ potentially indicates steric repulsion overwriting attractive interactions. Furthermore, putative CH- π interactions between the α -Methyl group of the imine and the phenyl entities of the 3,3'-substituent (Interaction B in Figure 9 a) as well as π - π interactions between the two imines (Interaction C in Figure 9 a) might compensate the loss of interactions between the imine and BINOL-backbone present in the monomeric structures.²⁹ Moreover, other 3,3'-substituent properties such as electrostatic repulsion, e.g. due to the CF3-groups in a hypothetic TRIFP dimer, or severe entropic penalties upon dimerization for catalysts with a high degree of conformational freedom of the 3,3'-substituents such as TiPSY and TRIP^{‡2} might also affect dimerization trends.

It was reported that interlocked CPA catalysts can form P=O···(HO)P hydrogen bonded pseudo-dimers which provide significantly higher stereoselectivity as their non-interlocked analogues.⁵⁴ However, for synthetic applications CPA/imine dimers are not expected to have any influence as their population diminishes when approaching catalyst concentrations used in synthesis.6,7,55

Hence, the existence of a dimer in the crystal structure of TeBuP and in the NMR spectra of TRIM, 9-Phen, and 1-Naph corroborates our previous assumption²⁹ that attractive interactions between 3,3'-substituent and imine play a key role in the energetics of CPA complexes. Thus, the dimerization trend might be tentatively proposed as a qualitative measure for steric properties and polarizability of the 3,3'-substituent.

Effect of substituents on enantioselectivity in the transfer hydrogenation of ketimines

After ensuring the analogy of all catalytic relevant and NMR accessible structures, the influence of the 3,3'-substituents on the enantioselectivity was investigated. Therefore, the ee values for the transfer hydrogenation of imines 5, 6 and 7 with ethyl Hantzsch ester catalyzed by TRIP, TiPSY, TRIFP, TRIM, 1naph and 9-phen were determined (see Supplementary S45). As anticipated based on earlier work of Rueping,⁵ List⁶ and MacMillan⁷ and on the webtool BINOPtimal developed by Goodman,²⁴ TiPSY (90-95 % ee) gave the highest enantiomeric excess, followed by TRIP (83-86 % ee) and TRIFP (74-91 % ee), while TRIM (56-70%), 1-naph (46-56%) and 9-phen (47-59%)

Journal Name

ARTICLE

gave lower enantioselectivities. While imine 5 and 6 gave almost identical ee values for all investigated catalysts, the CF3 substituted imine 7 deviated up to ±15 % for TRIFP, TRIM, 1naph, and 9-phen. Since imines 5, 6 and 7 have a similar steric bulk this deviation suggests that steric interactions are not the only key factor, but that also electrostatic interactions can significantly modulate the enantioselective outcome. Next, linear correlations between the observed enantioselectivities and steric descriptors (AREA(θ) or rotational barrier) of the 3,3' substituents as previously postulated by Goodman²¹ were tested. However, neither with area θ nor with the rotational barrier a linear correlation was found for our data. (see Supplementary S46). Moreover, the enantioselectivities also showed no correlation with the E/Z ratios of the binary complexes (see Supplementary S46). This shows that a single parameter derived from catalyst alone or binary complex is not sufficient to adequately predict the stereoselective outcome at least of the investigated reaction. Rather multiple factors seem to affect the energetics of the reaction pathway, which will be subject of further investigations.

Theoretical model structures and structure comparison

The above mentioned experimentally identified dimeric structures seem to be stabilized by attractive interactions, often associated with London dispersion forces. The presence of such interaction modes was already investigated and proven for TRIP/imine complexes in the previous theoretical structural studies utilizing NCI analysis and NOE analysis.^{29,32} The binary complex is mainly anchored by a strong hydrogen bond,³² however equally important, it is also bolstered by numerous weak attractive interactions between the imine and the catalyst's backbone as well as the bulky 3,3'-substituents.²⁹ These interactions are also pervasive in other catalysts with different 3,3' motives.

Our theoretical calculations of TRIP, TiPSY, TRIFP, TRIM/**5-7** confirm the general existence of the four core structures (*Type I/II E/Z*) (for computational details and data on additional CPAs, see Supplementary S49) in accordance with previous results for similar complexes.²⁹ Each of the *E*- and *Z*-imine complexes features two different orientations of the imine (Figure 10). In the *Type I* orientation, the ketone moiety is located in close proximity to the BINOL backbone of the catalyst. In the *Type II* orientation, the imine is rotated around the hydrogen bond by ~180° and the aniline moiety is located close to the BINOL backbone.

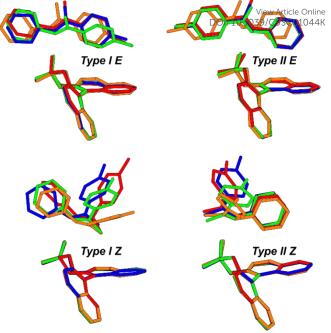


Figure 10. Theoretical calculations of the TiPSY, TRIP, TRIP, TRIPS complex show the invariance of four core structures. Despite that, the *E/Z* ratios and the tendency towards dimerization vary strongly within the CPA/imine complexes. Moreover, quantum chemical calculations of *E/Z*-ratio showed a significant offset to the experimental values. Colour code: TiPSY red, TRIP blue, TRIPP orange, TRIM green.

Geometrical comparison of the four core structures from the calculation revealed a high degree of invariance in the CPA/imine complexes, which is, to our surprise, retained despite significant variation of 3,3'-substituent of the catalyst (Figure 10). As in TRIP/imine complexes²⁹ the two orientations of the E- and Z-imine are obtained in TiPSY, TRIFP, TRIM/5-7 binary complexes. Furthermore, similar van der Waals interaction types (π - π , CH- π) are clearly recognizable from the structural analysis. In accordance with the previous studies,³² the acidic proton is located closer to the imine in TiPSY and TRIFP binary complexes, which indicates a strong predominant zwitter ionic, but also partly covalent character (POH angle \approx 110° - 120°). Due to the high degree of invariance of the core structures, the E/Z-ratio in the binary complexes is altered due to the difference in stabilization of E- and Z-imine by the 3,3'substituent. It is noteworthy to mention that despite a good agreement between experimentally and theoretically determined structures, the theoretical E/Z-ratio deviates significantly from the measured E/Z-ratio (Table 2).

Mostly, only the qualitative trend could be predicted correctly with exception of TRIM-complexes, i.e. experimentally the *E*complex is less stable than the *Z*-complex. Despite extensive conformational search (for computational details see below), application of various dispersion corrected density functionals (GGA, hybrid GGA, meta-hybrid GGA), post-HF methods (MP2, SCS-MP2, DLPNO-CCSD(T)) and implicit solvation models (SMD, COSMO-RS), the thermodynamic stability of the considered CPA/*Z*-isomers is gravely underestimated. One of the most common encountered error sources in the theoretical calculation during the simulation of large flexible molecules is the initial sampling using molecular mechanics not being able to find the initial structure near to global minimum. This problem

Journal Name

is augmented, especially in our case, by the poor parameterization of strongly hydrogen-bridged and zwitterionic complexes. Nevertheless, given the number of sampling conformations, experimental NOE contacts and the remarkable agreement of calculated coupling constants with the experimental values,³² the possibility of not finding the global minimum of the *Z*-complex is reduced significantly.

Since the weak interactions, e.g. dispersion effect in the complex, were already accounted in the post-HF methods, the underlying cause for the deviation might be either the missing explicit solvent-solute interaction favouring the Z-isomers due to its compactness, and hence a larger available solvent-solute interaction space, or the inaccurate implicit solvation of such zwitter-ionic species. For the former, tremendous increase of computational cost is expected making full ab initio calculations very restricted. An initial calculation of solvated complexes in a solvent box consisting of 300 solvent molecules has been performed and accurate calculations are planned for the future. For the latter, recent thorough investigation of solvent effect on the attenuation of inter- and intramolecular dispersive interaction showed that the current frequently employed implicit solvent models fail to describe dispersive solvent-solute interaction.⁵⁶ In summary, several underlying reasons for the failure to predict the E/Z-ratio are multiple. First, considering the huge conformational space of the complex, accurate initial sampling is necessary to catch the nearest minimum for DFT refined optimization. In this case, accurate semi-empirical methods, such as tight-binding method⁵⁷ and sampling using meta-dynamics may resolve the sampling issue in large electronic structures. Second, accurate description for solvation might be still underdeveloped for strongly bound zwitter-ionic species with large aromatic surfaces causing the solvent attenuation of dispersive interaction. Therefore, further improvement for implicit solvent correction is necessary.

Table 2. <i>E/Z</i> ratio of different catalyst with imine 5 .							
	Experimental E:Z	Theoretical E:Z					
TRIP/ 5	79:21	99.80 : 0.20					
TRIFP/5	70 : 30	96.80 : 3.20					
TiPSY/ 5	56 : 44	99.97 : 0.03					
TRIM/5	47 : 53	80.41 : 19.59					

In addition, a structure model for dimeric CPA/imine complexes was computed on the example of TRIM/**5** (see figure 9 and 11). This dimeric structure features two imines, nested between two catalysts. An extended dispersion force between the 3,3'-substituents of the two catalysts and the two extended aromatic imines is observed. The additional imine in the dimeric complex seems to be positioned at the nucleophilic attack site, when comparing the structure to the hydride transfer transition state with Hantzsch ester as the reducing agent. An overlap of the imine substrates and the Hantzsch ester is observed which suggests similar interaction modes (Figure 11c).

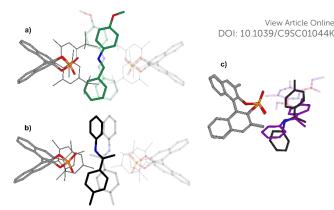


Figure 11. Reported crystal structure of a TeBuP/imine dimer (a) and calculated structure of a TRIM/imine dimer (b). The structures are similar despite comparing an aldimine (a) to a ketamine structure (b). c) Comparison of the dimeric structure of a TRIM complex (black imine) with the calculated transition state consisting of a CPA (the 3.3' substituents of the catalysts have been omitted to improve the visibility), an imine (purple) and a Hantzsch ester (purple, transparent). The steric influence of the second imine in the dimeric structure (b) is similar to the additional bulk of the Hantzsch ester in the transition state.

Computational Details

All binary complexes were optimized at DFT level of theory using TPSS functional with D3 correction in continuum of DCM. To mimic experimental low temperature condition the dielectric constant of DCM was increased to 16.20. Preconformational sampling was performed using force-field with MMFF parameter. Overall 1137 conformations of TRIP/imine were generated in the force-field procedure and subjected to ab initio geometry optimization. Low lying conformations were selected according to experimental constrains (NOE contacts). For other catalyst/imine conformations, core structures based on TRIP/imine complexes were taken and adapted. Post-HF single point calculations at SCS-MP2/CBS level of theory were performed on the optimized geometry (see Supplementary 46 for extrapolation procedure).

CONCLUSION

A combination of detailed NMR spectroscopic studies and theoretical calculations was used to investigate the structural space of CPA/imine complexes with varying 3,3'-substituents. Our experimental data revealed the coexistence of four core structures (Type I/II E and Type I/II Z) in solution, similar to those described for binary complexes of TRIP. These core structures show a high invariance, independent of the used CPA, which is corroborated by theoretical calculations of TiPSY/imine, TRIFP/imine and TRIM/imine complexes. For the first time dimeric CPA/imine complexes were identified in solution for TRIM, 9-Phen and 1-Naph with an imine occupying the space analogous to the transition state in the transfer hydrogenation. Surprisingly, this dimer consisting of two binary complexes is stabilized exclusively by attractive non covalent interactions. Hence, not only the steric repulsion term but also the attractive term of the large 3,3'-substituents has to be considered for the energetics of CPA complexes. Despite the good agreement of experiment and theory for the identified structures, a significant disparity was found for the E/Z ratios of the complexes. All binary complexes exhibit an increased

Ð

ence

Journal Name

View Article Online

ARTICLE

population of the Z-structures, which cannot be reproduced in the calculations. A possible reason is an insufficient description of solvent-complex interactions. This study may allow to validate and improve theoretical predictive models by providing for the first time detailed experimental data about the structural space and influence of the 3,3'-substituent on the energetics of CPA/imine complexes.

Conflicts of interest

There are no conflicts to declare.

Acknowledgements

Financial Support was provided by the European Research Council (ERC-CoG 614182 - IonPairsAtCatalysis) and intellectual support was provided by SPP 1807 dispersion. We would like to thank Dr. Nils Sorgenfrei and Thomas Hausler for preliminary investigations of catalyst imine complexes and Dr. Fabio Morana for establishing in the beginning of our ion pair investigations the syntheses of catalysts and substrates.

Notes and references

‡1 Due to chemical exchange between TRIM/imine, [TRIM/imine]₂ and free imine and the resulting line broadening as well as signal overlap, only the complex TRIM/7 could be analysed in detail, as it gave adequate line widths.

^{‡2} Significant line broadening for distinct proton signals of TRIPs isopropyl and TiPSYs phenyl-moieties at 180 K caused by medium chemical exchange between different substituent conformations demonstrated a high degree of conformational freedom for these catalysts.

- D. Parmar, E. Sugiono, S. Raja and M. Rueping, Chem. Rev., 1 2014, 114, 9047-9153.
- 2 D. Parmar, E. Sugiono, S. Raja and M. Rueping, Chem. Rev., 2017, 117, 10608-10620.
- 3 T. Akiyama, J. Itoh, K. Yokota and K. Fuchibe, Angew. Chem. Int. Ed., 2004, 43, 1566-1568.
- 4 D. Uraguchi and M. Terada, J. Am. Chem. Soc., 2004, 126, 5356-5357.
- 5 M. Rueping, E. Sugiono, C. Azap, T. Theissmann and M. Bolte, Org. Lett., 2005, 7, 3781-3783.
- 6 S. Hoffmann, A. M. Seayad and B. List, Angew. Chem. Int. Ed., 2005, 44, 7424-7427.
- 7 R. I. Storer, D. E. Carrera, Y. Ni and D. W. C. MacMillan, J. Am. Chem. Soc., 2006, 128, 84-86.
- 8 K. Saito and T. Akiyama, Chem. Commun., 2012, 48, 4573-4575.
- 9 M. Yamanaka, J. Itoh, K. Fuchibe and T. Akiyama, J. Am. Chem. Soc., 2007, 129, 6756-64.
- 10 X. Xu, Y. Qian, L. Yang and W. Hu, Chem. Commun., 2010, 47, 38 797-799.
- E. P. Ávila, R. M. S. Justo, V. P. Gonçalves, A. A. Pereira, R. 11 Diniz and G. W. Amarante, J. Org. Chem., 2015, 80, 590-594.
- K. Shen, X. Liu, Y. Cai, L. Lin and X. Feng, Chem. A Eur. J., 12

2009, 15, 6008-6014.

- 13 C. Gharui, S. Singh and S. C. Pan, Org Biomore Chem. 2017, 15, 7272-7276.
- 14 S. Saha and C. Schneider, Chem. Eur. J., 2015, 21, 2348-2352.
- 15 S. G. Ouellet, A. M. Walji and D. W. C. Macmillan, Acc. Chem. Res., 2007, 40, 1327-1339.
- 16 C. Min and D. Seidel, Chem. Soc. Rev., 2017, 46, 5889–5902.
 - Q. Kang, Z.-A. Zhao and S.-L. You, Org. Lett., 2008, 10, 2031-2034.
- K. Saito, K. Horiguchi, Y. Shibata, M. Yamanaka and T. 18 Akiyama, Chem. Eur. J., 2014, 20, 7616-7620.
- 19 R. Maji, S. C. Mallojjala and S. E. Wheeler, Chem. Soc. Rev. 2018, 47, 1142-1158.
- 20 A. Milo, A. J. Neel, F. D. Toste and M. S. Sigman, Science, 2015, 347, 737-743.
- 21 J. P. Reid and J. M. Goodman, J. Am. Chem. Soc., 2016, 138, 7910-7917.
- 22 J. P. Reid, L. Simón and J. M. Goodman, Acc. Chem. Res., 2016, **49**, 1029–1041.
- 23 L. Simón and J. M. Goodman, J. Org. Chem., 2011, 76, 1775-1788.
- J. P. Reid and J. M. Goodman, Chem. Commun., 2019, 55, 24 1778-1781.
- 25 C. B. Santiago, J. Y. Guo and M. S. Sigman, Chem. Sci., 2018, 9, 2398-2412.
- 26 A. J. Neel, A. Milo, M. S. Sigman and F. D. Toste, J. Am. Chem. Soc., 2016, 138, 3863-3875.
- M. Orlandi, J. A. S. Coelho, M. J. Hilton, F. D. Toste and M. S. 27 Sigman, J. Am. Chem. Soc., 2017, 139, 6803-6806.
- 28 M. Sickert, F. Abels, M. Lang, J. Sieler, C. Birkemeyer and C. Schneider, Chem. Eur. J., 2010, 16, 2806-2818.
- 29 J. Greindl, J. Hioe, N. Sorgenfrei, F. D. Morana and R. M. Gschwind, J. Am. Chem. Soc., 2016, 49, 15965-15971.
- 30 W. Tang, S. Johnston, J. A. Iggo, N. G. Berry, M. Phelan, L. Lian, J. Bacsa and J. Xiao, Angew. Chem. Int. Ed., 2013, 52, 1668-1672.
- L. Liu, M. Leutzsch, Y. Zheng, M. W. Alachraf, W. Thiel and B. 31 List, J. Am. Chem. Soc., 2015, 137, 13268-13271.
- N. Sorgenfrei, J. Hioe, J. Greindl, K. Rothermel, F. Morana, N. 32 Lokesh and R. M. Gschwind, J. Am. Chem. Soc., 2016, 138, 16345-16354.
 - L. Simón and J. M. Goodman, J. Am. Chem. Soc., 2008, 130, 8741-8747.
 - T. Akiyama, Chem. Rev., 2007, 107, 5744-5758.
 - E. R. P. Zuiderweg and S. W. Fesik, Biochemistry, 1989, 28, 2387-2391.
- 36 J. Schleucher, M. Schwendinger, M. Sattler, P. Schmidt, O. Schedletzky, S. J. Glaser, O. W. Sørensen and C. Griesinger, J. Biomol. NMR, 1994, 4, 301-306.
 - M. Foroozandeh, R. W. Adams, M. Nilsson and G. A. Morris, J. Am. Chem. Soc., 2014, 136, 11867-11869.
 - J. Mauhart, S. Glanzer, P. Sakhaii, W. Bermel and K. Zangger, J. Magn. Reson., 2015, 259, 207-215.
 - K. Zangger, Prog. Nucl. Magn. Reson. Spectrosc., 2015, 86-87, 1-20.
 - D. Uhrín, J. Bramham, S. J. Winder and P. N. Barlow, J.

33

34

35

37

39

40

17

This article is licensed under a Creative Commons Attribution-NonCommercial 3.0 Unported Licence.

Open Access Article. Published on 08 April 2019. Downloaded on 4/25/2019 3:32:10 PM.

Journal Name

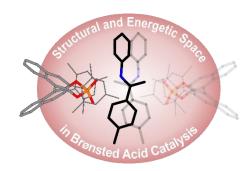
Biomol. NMR, 2000, **18**, 253–259.

- 41 A. Verma and B. Baishya, J. Magn. Reson., 2016, **266**, 51–58.
- 42 S. Akoka and P. Giraudeau, *Magn. Reson. Chem.*, 2015, **53**, 986–994.
- 43 R. J. Lichtenecker, N. Coudevylle, R. Konrat and W. Schmid, *ChemBioChem*, 2013, **14**, 818–821.
- 44 C. Prasanna, A. Dubey and H. S. Atreya, in *Methods in Enzymology*, Academic Press, 2015, vol. 565, pp. 167–189.
- 45 D. P. Frueh, A. C. Goodrich, S. H. Mishra and S. R. Nichols, *Curr. Opin. Struct. Biol.*, 2013, **23**, 734–739.
- H. Kessler, H. Oschkinat, C. Griesinger and W. Bermel, J. Magn. Reson., 1986, 70, 106–133.
- 47 K. Stott, J. Stonehouse, J. Keeler, T.-L. Hwang and A. J. Shaka, J. Am. Chem. Soc., 1995, **117**, 4199–4200.
- 48 D. Jeannerat and J. Furrer, *Comb. Chem. High Throughput Screen.*, 2012, **15**, 15–35.
- 49 C. Yu and G. C. Levy, J. Am. Chem. Soc., 1984, **106**, 6533– 6537.
- P. S. Pregosin, P. G. Anil Kumar and I. Fernández, *Chem. Rev.*, 2005, **105**, 2977–2998.
- 51 K. Shikii, S. Sakurai, H. Utsumi, H. Seki and M. Tashiro, *Anal. Sci.*, 2004, **20**, 1475–1477.
- 52 D. P. Cistola and K. B. Hall, J. Biomol. NMR, 1995, 5, 415–419.
- L. Fusaro, E. Locci, A. Lai and M. Luhmer, *J. Phys. Chem. B*, 2010, **114**, 3398–3403.
- 54 R. Mitra, H. Zhu, S. Grimme and J. Niemeyer, *Angew. Chem. Int. Ed.*, 2017, **56**, 11456–11459.
- 55 M. Sickert and C. Schneider, *Angew. Chem. Int. Ed.*, 2008, 47, 3631–3634.
- 56 R. Pollice, M. Bot, I. J. Kobylianskii, I. Shenderovich and P. Chen, *J. Am. Chem. Soc.*, 2017, **139**, 13126–13140.
- 57 S. Grimme, C. Bannwarth and P. Shushkov, *J. Chem. Theory Comput.*, 2017, **13**, 1989–2009.

View Article Online DOI: 10.1039/C9SC01044K

Table of Content Entry

View Article Online DOI: 10.1039/C9SC01044K



Advanced NMR and theoretical studies reveal detailed insights into monomeric and dimeric structures plus deviating experimental and theoretical *E/Z*-populations.

Chemical Science Accepted Manuscript