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**Authors:** Alexander J. Hoffman, Jason S. Bates, John R. Di Iorio,  
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# Rigid Arrangements of Ionic Charge in Zeolite Frameworks Conferred by Specific Al Distributions Preferentially Stabilize Alkanol Dehydration Transition States

Alexander J. Hoffman<sup>[a]</sup>, Jason S. Bates<sup>[b]</sup>, John R. Di Iorio<sup>[b]</sup>, Steven V. Nystrom<sup>[a]</sup>, Claire T. Nimlos<sup>[b]</sup>, Rajamani Gounder<sup>[b]\*</sup>, David Hibbitts<sup>[a]\*</sup>

**Abstract:** Zeolite reactivity depends on the solvating environments of their micropores and the proximity of their Brønsted acid sites. Turnover rates (per H<sup>+</sup>) for methanol and ethanol dehydration increase with the fraction of H<sup>+</sup> sites sharing six-membered rings of chabazite (CHA) zeolites. DFT shows that activation barriers vary widely with the number and arrangement of Al (1–5 per 36 T-site unit cell), but cannot be described solely by Al–Al distance or density. Certain Al distributions yield rigid arrangements of anionic charge that stabilize cationic intermediates and transition states via H-bonding to decrease barriers. This is a key feature of acid catalysis in zeolite solvents, which lack the isotropy of liquid solvents. The sensitivity of polar transition states to specific arrangements of charge in their solvating environments and the ability to position such charges in zeolite lattices with increasing precision herald rich catalytic diversity among zeolites of varying Al arrangement.

## Introduction

Zeolites are crystalline aluminosilicates that are ubiquitous as acid catalysts in carbon conversion routes<sup>[1,2]</sup> because they provide diversity of function arising from differences in crystal topology and habit, which can be synthetically engineered.<sup>[3–7]</sup> Their microporous voids are of molecular dimension (<2 nm) and provide solvating environments that confine and stabilize, via non-specific van der Waals interactions, the covalent and ion-pair intermediates and transition states that mediate gas-phase reactions.<sup>[8–11]</sup> Zeolite micropores also contain Brønsted acidic OH groups that charge-compensate framework Al and Si–OH groups at lattice defects which act as hydrophilic binding sites that influence the extended H-bonded clusters of water,<sup>[12]</sup> alkanols and other polar protic molecules commonly used as solvents in liquid-phase reactions.<sup>[13–17]</sup> Catalytic regimes exist absent a condensed intrapore phase, however, wherein molecular clusters and networks of specific structure and size are stabilized within microporous voids, and their stability depends intimately on the geometry and charge distribution within the environment provided by the solvating zeolite lattice. Such features of the zeolite

“solvent” and their effects on acid catalysis, however, are not well-understood.

Additionally, the effects of zeolite acid site proximity on turnover rates have been documented,<sup>[18,19]</sup> but only anecdotally and with little understanding or consensus about the mechanistic origin of such rate differences. Acid strength—as determined by deprotonation energy (DPE), a theoretical metric defined as the energy to remove a proton from its conjugate base to a non-interacting distance—increases for Brønsted acid sites in proximal configurations (Al–O–(Si–O)<sub>x</sub>–Al,  $x=1,2$ ),<sup>[20]</sup> indicating that interactions between proximal sites may increase turnover rates solely due to electronic effects by facilitating charge separation during transition state formation.<sup>[21,22]</sup> Turnover rates of protolytic alkane activation<sup>[23]</sup> and alkene oligomerization<sup>[24–26]</sup> are reported to increase with acid site proximity in MFI zeolites, yet firm mechanistic interpretations are precluded by the numerous Al–Al pair configurations present within the low-symmetry (12 tetrahedral-site; T-site) MFI framework. Here, we examine the effects of acid site proximity in zeolite catalysis using the high-symmetry (1 T-site) CHA framework and alkanol dehydration as the probe reaction, which offers significant promise for developing mechanistic understanding of how acid site proximity influences turnover rates, given that the abundant surface intermediates and reaction mechanisms in this chemistry are well understood.<sup>[27]</sup> Moreover, gas-phase alkanol dehydration reactions can occur at high coverages of adsorbed alkanol clusters and networks reminiscent of those present during liquid-phase catalysis, promising to offer molecular insight into interactions among proximal acid sites mediated by co-adsorbates that have gone largely ignored in prior studies of H-form zeolites, but constitute a mechanism by which turnover rates vary with site arrangement.

## Results and Discussion

SSZ-13 zeolites (CHA, Fig. 1) have one crystallographically unique T-site; as such, each acid site has an identical local environment, thus avoiding complications of different acid site locations and reducing the combinatorial complexity of proximal Al–Al site ensembles. CHA is comprised of 6-membered ring (6-MR) units adjacent to 8-MR windows (3.7 Å in diam.) that separate CHA cages (7.4 × 9.8 Å).<sup>[28]</sup> CHA can be synthesized with a specific acid site density (Si:Al ratio) but with varying distributions of proximal Al–Al site ensembles, for example by varying the ratios of *N,N,N*-trimethyl-1-adamantylammonium (TMAda<sup>+</sup>) to Na<sup>+</sup> structure-directing agents (SDAs) present during crystallization at fixed total SDA cation concentration.<sup>[29,30]</sup> This synthesis procedure enables systematically varying the number

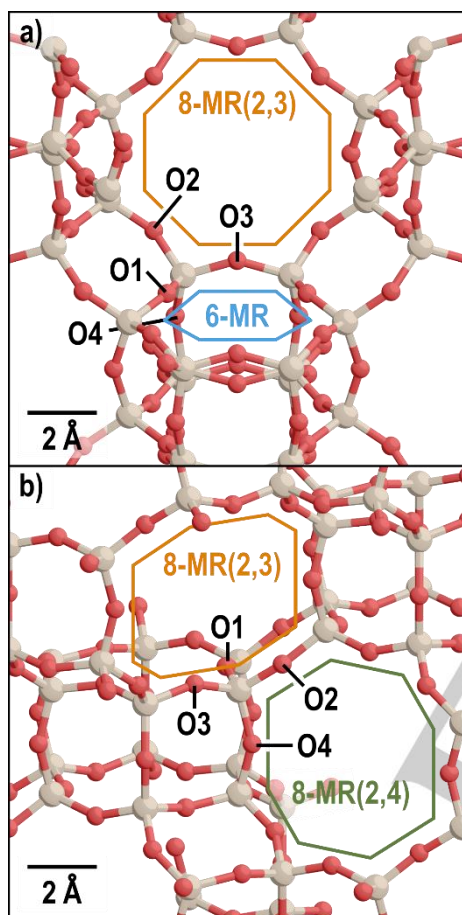
[a] A. J. Hoffman, Dr. S. V. Nystrom, Prof. Dr. D. D. Hibbitts  
Department of Chemical Engineering  
University of Florida  
1030 Center Dr, Gainesville, FL 32608 (USA)  
E-mail: [hibbitts@che.ufl.edu](mailto:hibbitts@che.ufl.edu)

[b] Dr. J. S. Bates, Dr. J. R. Di Iorio, C. T. Nimlos, Prof. Dr. R. Gounder  
Charles D. Davidson School of Chemical Engineering  
Purdue University  
480 Stadium Mall Drive, West Lafayette, IN 47907 (USA)  
E-mail: [rgounder@purdue.edu](mailto:rgounder@purdue.edu)

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of Al–Al pairs within a 6-MR (as measured by  $\text{Co}^{2+}$  titration<sup>[29,31]</sup>), thus avoiding effects caused by simultaneously varying bulk acid site density to allow better elucidation of acid site proximity effects on catalysis.<sup>[23,24,32,33]</sup> This and other methods to influence Al proximity in zeolites<sup>[19,25,34,35]</sup> have motivated renewed investigation of the kinetic and mechanistic consequences of acid site proximity in zeolite catalysis.<sup>[18]</sup>



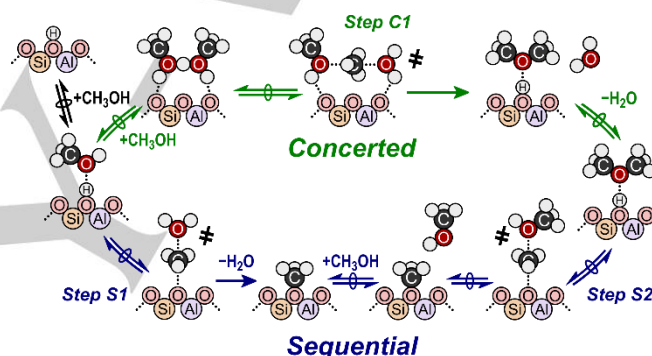
**Figure 1.** The CHA structure showing (a) the 6-MR and 8-MR(2,3) (containing O2 and O3) structures and (b) the 8-MR(2,3) and 8-MR(2,4) around one T-site. The four symmetric O atoms around one T-site are labelled.

Methanol dehydration to dimethyl ether (DME) is an informative probe reaction for solid Brønsted acids, used previously to decouple acid strength and confinement effects among zeolites of different acid strength and topology<sup>[21,22,36–39]</sup> and polyoxometalates.<sup>[22,40,41]</sup> Methanol dehydration can occur by two competing mechanisms: a sequential (dissociative) or a concerted (associative) mechanism (Scheme 1), where the former creates a surface methyl ( $\text{CH}_3\text{-Z}$ ) that reacts with methanol to form DME, while the latter forms DME in a single bimolecular reaction. At 415 K, a typical condition, density functional theory (DFT) calculations show that methanol dehydrates via the concerted mechanism at all relevant methanol pressures ( $>0.3$  kPa) in CHA.<sup>[27]</sup> Abundant surface species vary

from methanol monomers to dimers and larger methanol clusters over the pressure ranges of kinetic studies, and their change in molecularity causes a transition from a first- to a zero-order regime on all H-form zeolites.<sup>[21,27,36,42]</sup> With increasing methanol pressure, rates become inhibited by  $\text{CH}_3\text{OH}$  on small-pore, cage-window zeolites (CHA, AEI, LEV, LTA; structures shown in Figs. S1–S3, SI) because methanol clusters ( $\geq 3$   $\text{CH}_3\text{OH}$  per  $\text{H}^+$ ) larger than the molecularity of kinetically relevant transition states form, requiring desorption of some extraneous methanol prior to reaction.<sup>[27,31]</sup> Such methanol dehydration rates are described by:

$$r_{\text{DME}} = \frac{k_{\text{first}} P_{\text{M}}}{1 + \frac{k_{\text{first}}}{k_{\text{zero}}} P_{\text{M}} + \frac{k_{\text{first}}}{k_{\text{inhib}}} P_{\text{M}}^2} \quad (1)$$

where  $k_{\text{first}}$ ,  $k_{\text{zero}}$ , and  $k_{\text{inhib}}$  are the first-order, zero-order, and inhibitory rate coefficients and  $P_{\text{M}}$  is the methanol pressure (derivation in Section S9, SI).<sup>[27,31]</sup>

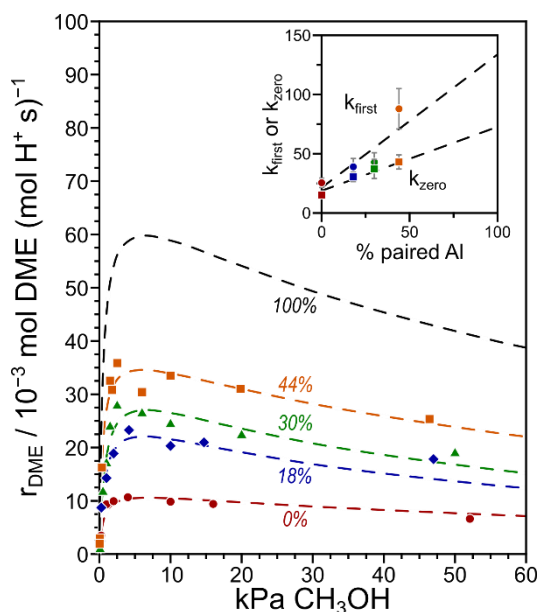


**Scheme 1.** Two parallel methanol dehydration mechanisms: concerted (top) and sequential (bottom). Both routes can occur with spectating methanol molecules forming extended complexes with the species depicted; adapted from prior work.<sup>[27]</sup>

Recently, we reported that methanol dehydration turnover rates (per  $\text{H}^+$ ) increase systematically with the percentage of 6-MR paired acid sites in H-CHA (0–44%; Fig. 2).<sup>[31]</sup> First- and zero-order rate coefficients (Eq. 1) fit to the kinetic data also increase systematically with site-pairing in the 6-MR, and rate constants extrapolated to a hypothetical CHA sample with 100% of its sites in paired configurations (Fig. 2) are 7.2 $\times$  and 4.4 $\times$  larger for first-order and zero-order rate coefficients, respectively, than on isolated sites.<sup>[31]</sup> The first-order rate coefficient reflects the effective free energy barrier ( $\Delta G^\ddagger$ ) to form a transition state with one additional methanol from an adsorbed methanol complex—e.g., forming a bimolecular transition state from a methanol monomer.<sup>[22,27]</sup> The zero-order rate coefficient instead reflects the intrinsic activation free energy ( $\Delta G_{\text{act}}$ ) to form a transition state with the same number of methanol molecules as its precursor—e.g., forming a bimolecular transition state from an adsorbed methanol dimer.<sup>[22,27]</sup> The higher rate coefficients on paired sites correspond to apparent and intrinsic free energy barriers ( $\Delta G^\ddagger$  and  $\Delta G_{\text{act}}$ ) values that are lower by 7 and 5  $\text{kJ mol}^{-1}$ , respectively, on paired than on isolated sites. Transition states can be unimolecular, bimolecular, or have additional H-bonded methanol

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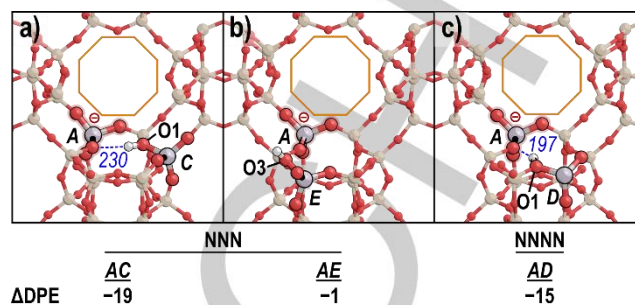
molecules that act to stabilize them; similarly, adsorbed methanol complexes can include 1–4 methanol molecules over this range of methanol pressures.<sup>[27]</sup>



**Figure 2.** DME formation rates (per  $\text{H}^+$ , 415 K) as a function of methanol pressure on H-CHA with 0% ( $\bullet$ , red), 18% ( $\blacklozenge$ , blue), 30% ( $\blacktriangle$ , green), 44% ( $\blacksquare$ , orange), and the extrapolation to 100% (black) 6-MR paired Al. Dashed lines represent regression to Eq. 1. Inset: first-order ( $k_{\text{first}}$ ,  $\bullet$ ,  $10^{-3}$  mol DME ( $\text{kPa mol H}^+ \text{s})^{-1}$ ) and zero-order ( $k_{\text{zero}}$ ,  $\blacksquare$ ,  $10^{-3}$  mol DME ( $\text{mol H}^+ \text{s})^{-1}$ ) rate coefficients as a function of the percentage of 6-MR paired Al. Adapted from prior work.<sup>[31]</sup>

There are three possible Brønsted acid site-pair locations within the 6-MR (AC, AD, AE, Fig. 3), omitting pairs in which Al occupy neighboring T-sites that would violate Löwenstein's Rule.<sup>[43]</sup> The Al in both the AC and AE site-pairs are separated by one Si atom (Fig. 3a,b), forming two next-nearest neighbor (NNN) site-pairs. These arrangements behave as a single indistinguishable site-type during catalysis (see Section S4, SI), and thus are treated collectively. The Al in the AD site-pair are separated by two Si atoms and are thus in next-next-nearest neighbor (NNNN) positions (Fig. 3c). Acid sites in these 6-MR paired configurations are stronger acids because deprotonation of one site in the pair allows the other proton to H-bond with and stabilize the conjugate base (Fig. 3),<sup>[20]</sup> evident in decreases in DPE of 19 and 15  $\text{kJ mol}^{-1}$  for the AC (NNN) and AD (NNNN) site-pairs, respectively, compared to an isolated site. In contrast, the AE site-pair does not allow the framework to contort upon deprotonation of the A site to facilitate the  $\text{O1-H-AlO}_4^-$  interaction (Fig. S5, SI); instead, the remaining proton on site E binds to O3, precluding H-bonding to the conjugate base and resulting in the same DPE as an isolated site ( $\Delta\text{DPE} = -1$   $\text{kJ mol}^{-1}$ , Fig. 3b). These calculations show that H-bonding can stabilize conjugate bases across 6-MR motifs in both NNN and NNNN Al site-pair arrangements; importantly, they reveal how cationic species—here, a proton—are stabilized by

specific arrangements of Al and associated anionic charge positioned in the solvating environment.



**Figure 3.** Change in acid strength relative to an isolated site, given by the DPE difference ( $\text{kJ mol}^{-1}$ ). The conjugate base upon A-site deprotonation is shown for the NNN site-pairs (a) AC and (b) AE, and the NNNN site-pair (c) AD. Blue dashed lines represent H-bonds (lengths in pm). Adapted from prior work.<sup>[20]</sup>

These interactions between a conjugate base and a bare proximal proton may be relevant to reactions that prevail at low acid site coverages, such as high-temperature ( $>700$  K) alkane cracking.<sup>[9,32,44–51]</sup> Methanol dehydration, however, occurs at acid sites that are fully covered by methanol species ( $>0.15$  kPa  $\text{CH}_3\text{OH}$ ; *in situ* IR<sup>[27,31]</sup> and DFT<sup>[27]</sup> evidence). Co-adsorbed methanol species alter both the H-bonding interactions among site pairs that determine DPE and influence the stability of methanol dehydration transition states. Thus, we performed a theoretical evaluation of methanol dehydration at isolated and paired Al in CHA to elucidate the mechanistic origins of the observed rate enhancement.

Methanol dehydration reactions were modeled at site A while the second site (C, D, E) binds spectating molecules. We focus on reactions with 1–2 adsorbed  $\text{CH}_3\text{OH}$  per site (results with bare proximal sites in Section S5, SI), and the simplest routes that form DME—sequential and concerted paths without spectators—because these capture barriers for  $\text{CH}_3\text{OH}$  adsorption to form a protonated dimer and then to form DME, which are the dominant factors affecting turnover rates. These methanol dehydration routes involve three transition states (Scheme 1) corresponding to zeolite methylation (Step S1), methanol methylation (Step S2), and concerted DME formation (Step C1).<sup>[27]</sup> The free energies and structures presented here for reactants and transition states are the most stable among  $\sim 150$  unique optimized structures for each reaction that were identified by altering the framework O atoms involved in methylation/demethylation reactions and by systematic reorientations of exemplars. These reorientations (described in Section S6, SI) are necessary because DFT methods only identify reactants and transition states near input structures, while numerous structures exist differing in energy by up to 200  $\text{kJ mol}^{-1}$ .<sup>[52]</sup> Reorientations indicate that H-bonding is critical for these transition states—configurations that maximize H-bonding with minimal framework restructuring and internal transition state distortions are consistently preferred.



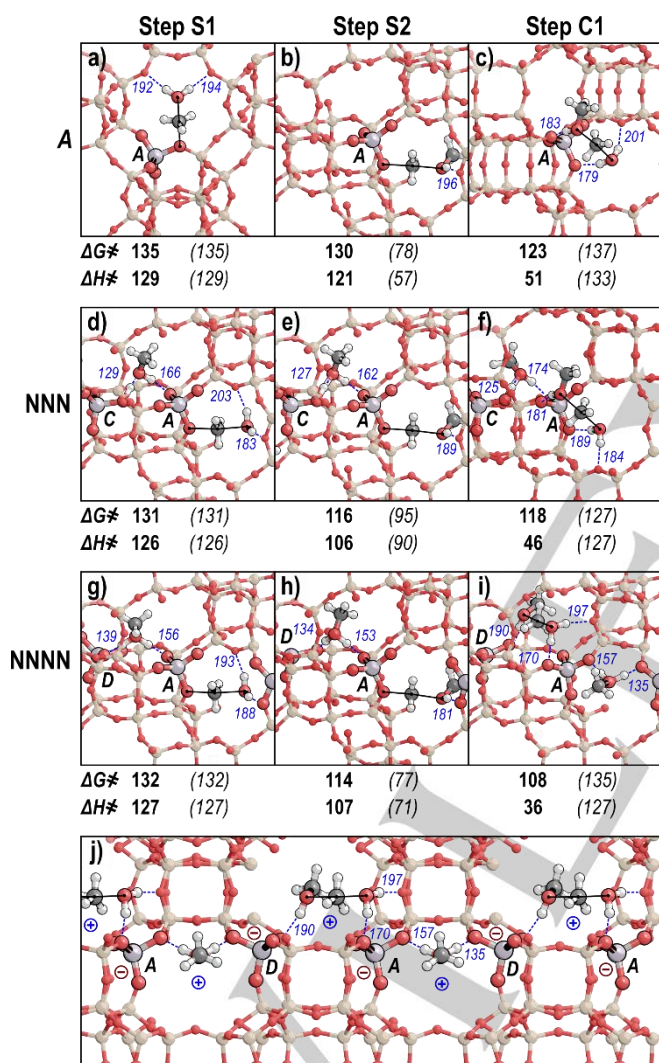
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We first examine the most stable transition states at isolated Al (A) sites (Fig. 4a-c). Zeolite methylation (Step S1) has a transition state structure of  $\text{H}_2\text{O}-\text{CH}_3^+-\text{O}_2^-$  and is most stable within 8-MR(2,3) with  $\text{H}_2\text{O}$  forming two H-bonds with framework O atoms (Fig. 4a); it has an intrinsic activation barrier ( $\Delta G_{\text{act}}$ , 415 K, 1 bar) of 135  $\text{kJ mol}^{-1}$  when methylating O3. The transition state for methanol methylation (Step S2) is nearly identical with  $\text{CH}_3\text{OH}$  instead of  $\text{H}_2\text{O}$  ( $\text{CH}_3\text{OH}-\text{CH}_3^+-\text{O}_2^-$ ) and one H-bond with the framework; it prefers demethylating O4 and has a  $\Delta G_{\text{act}}$  of 78  $\text{kJ mol}^{-1}$  (Fig. 4b). The transition state of the concerted reaction (Step C1,  $\text{CH}_3\text{OH}-\text{CH}_3^+-\text{H}_2\text{O}$ ) resides in the CHA cage and H-bonds to two O of the conjugate base in 8-MR(2,4) (Fig. 4c); it has a  $\Delta G_{\text{act}}$

of 137  $\text{kJ mol}^{-1}$  and an effective barrier ( $\Delta G^\ddagger$ ) of 123  $\text{kJ mol}^{-1}$ , referenced to an adsorbed methanol monomer that is the low-pressure most abundant surface intermediate (MASI). This  $\Delta G^\ddagger$  for concerted DME formation (123  $\text{kJ mol}^{-1}$ ) is 12  $\text{kJ mol}^{-1}$  lower than that of zeolite methylation (Step S1,  $\Delta G^\ddagger = 135 \text{ kJ mol}^{-1}$ ), which limits the rate of the sequential pathway. The lower barrier for the concerted mechanism indicates that it prevails over the sequential pathway at isolated acid sites and at conditions relevant to kinetic studies, as shown in our prior work.<sup>[27]</sup>

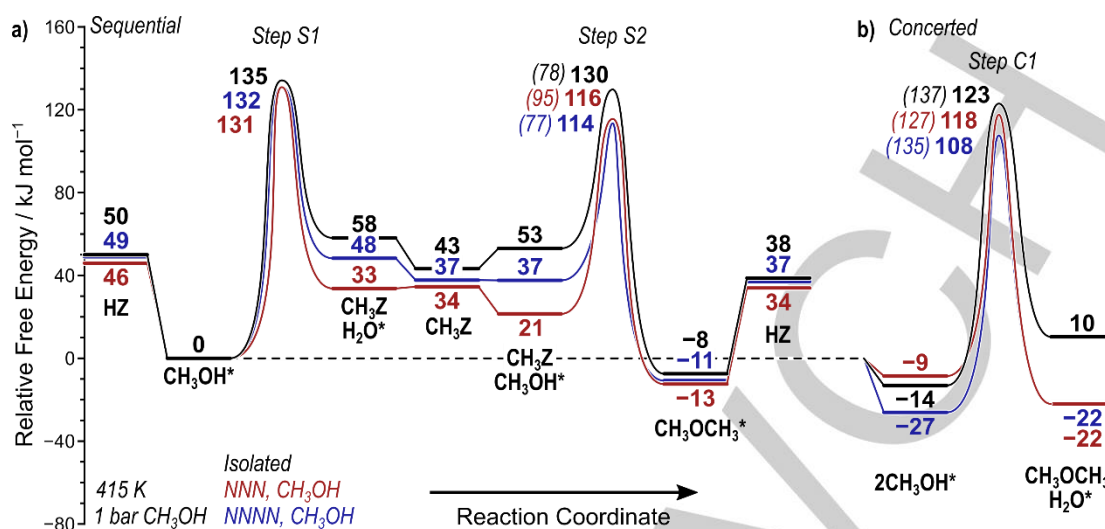
We next examine the most stable transition states at paired Al sites in NNN positions (AC), where the second site has one adsorbed methanol (Fig. 4d-f). Step S1 transition states are most stable in the 8-MR(2,4) to avoid interacting with the methanol bound to site C. This methanol facilitates an interaction between the proximal Al sites through two strong H-bonds (129 and 166 pm) that reduce the  $\Delta G_{\text{act}}$  for surface methylation to 131  $\text{kJ mol}^{-1}$  on the NNN site-pair (Fig. 4d) from 135  $\text{kJ mol}^{-1}$  on an isolated site (Fig. 4a). Adding the second site decreases the barrier by 17  $\text{kJ mol}^{-1}$  for methylating in 8-MR(2,4), but rearranging the transition state from 8-MR(2,3) to 8-MR(2,4) requires +13  $\text{kJ mol}^{-1}$ , resulting in an effective decrease of 4  $\text{kJ mol}^{-1}$  (Fig. S9, SI). An analogous interaction exists for Step S2 (Fig. 4b,e) and decreases  $\Delta G^\ddagger$  by 14  $\text{kJ mol}^{-1}$  on the NNN site-pair compared to the isolated Al site, although this step is not kinetically relevant during methanol dehydration.<sup>[27,31]</sup> Concerted methanol dehydration also occurs with lower  $\Delta G^\ddagger$  and  $\Delta G_{\text{act}}$  on the NNN site-pair (118 and 127  $\text{kJ mol}^{-1}$ ) than on isolated sites (123 and 137  $\text{kJ mol}^{-1}$ ; Fig. 5), and this pathway should thus prevail over the sequential pathway at these conditions.<sup>[27]</sup> These reductions of 5 and 11  $\text{kJ mol}^{-1}$  in  $\Delta G^\ddagger$  and  $\Delta G_{\text{act}}$ , respectively, for the concerted methanol pathway are similar to the decrease in free energy barriers (7 and 5  $\text{kJ mol}^{-1}$ ) measured in experimental kinetic data.<sup>[31]</sup> All three methanol dehydration transition states are stabilized by the presence of a methanol monomer associated with a second Al site at the NNN position in the 6-MR, indicating that alkanol co-adsorbates facilitate inter-site H-bonding interactions similar to how bare protons do so to increase acid strength (Fig. 3) rather than stabilizing transition states via van der Waals interactions. These differences in free energy barriers of 4–15  $\text{kJ mol}^{-1}$  are consistent with the observed rate increases but are relatively small compared to the overall accuracy of DFT methods, so a second site pair (NNNN) can corroborate these findings.

We also examine the most stable transition states at paired Al sites in NNNN positions (AD) to determine whether these decreases in free energy barriers are ubiquitous for site-pairs located in the 6-MR of CHA. Transition states for Steps S1 and S2 exhibit similar interactions across the 6-MR (Fig. 4g,h). Barriers for Steps S1 and S2 at the NNNN site-pair decrease by 3 and 16  $\text{kJ mol}^{-1}$ , similar to those on the NNN site-pair (Fig. 5). The concerted transition state for the NNNN site-pair is unique as it bridges the 6-MR (Fig. 4i) while the co-adsorbed methanol interacts simultaneously with the site-pair via the 8-MR(2,4). The simultaneous interaction of this site pair across the 6-MR and 8-



**Figure 4.** The most stable transition state structures for methanol dehydration in CHA for Step S1, Step S2, and Step C1 for isolated acid sites (A) and NNN and NNNN site-pairs. Effective free energy (415 K, 1 bar) and enthalpy barriers ( $\Delta G^\ddagger$  and  $\Delta H^\ddagger$ ), and intrinsic barriers ( $\Delta G_{\text{act}}$  and  $\Delta H_{\text{act}}$  in parenthesis and italicized) are in  $\text{kJ mol}^{-1}$ . Solid black lines indicate incipient and breaking bonds. Blue dashed lines indicate H-bonds (lengths in pm). (j) Interactions between alternating cations and anions from periodic boundary conditions in the Step C1 transition state on the NNNN site-pair. Alternate and larger views given for all DFT-obtained structures in Figures S14–S16 (SI).

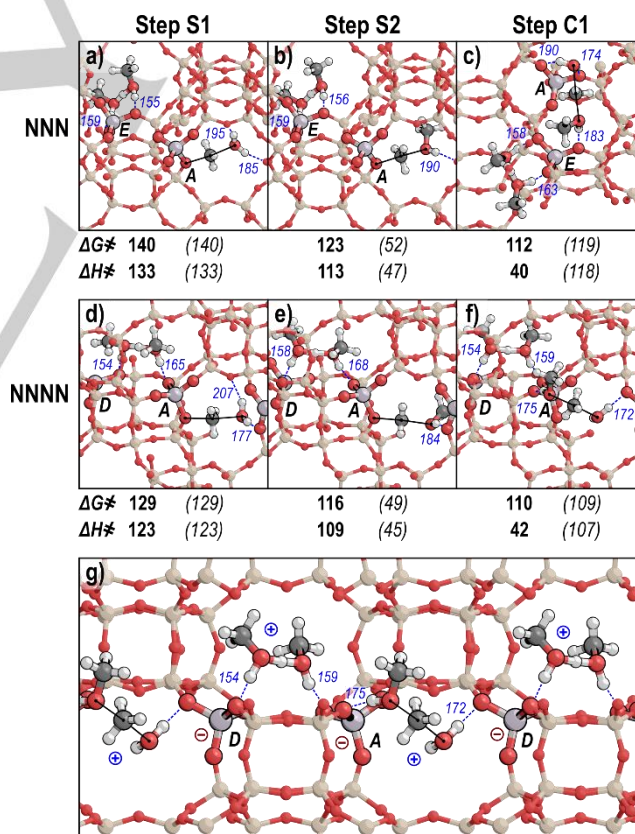
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**Figure 5.** Reaction coordinate diagram for (a) sequential and (b) concerted methanol dehydration on isolated sites (black), the NNN (red) and NNNN (blue) site-pairs with a  $\text{CH}_3\text{OH}$  monomer at the second acid site. Free energies (415 K, 1 bar) are shown relative to a methanol monomer on all sites. Intrinsic barriers are shown for Steps S2 and C1 in parentheses. Structures for kinetically relevant structures are shown in Fig. 4 and Figs. S12–S16 (SI).

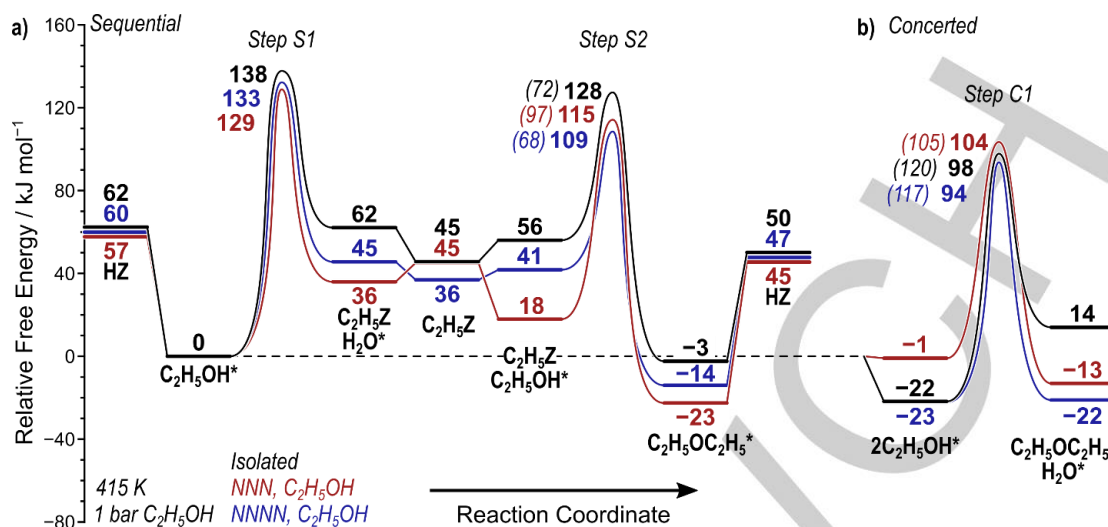
MR(2,4) occurs because of the periodic boundary conditions and the relatively small CHA unit cell. This forms an infinite chain of Al-centered anions and cationic species (Fig. 4j), resulting in lower  $\Delta G^\ddagger$  and  $\Delta G_{\text{act}}$  for the concerted transition state on NNNN site-pairs (108 and 135  $\text{kJ mol}^{-1}$ ) compared to NNN (118 and 127  $\text{kJ mol}^{-1}$ ), and again lower than for the isolated site (123 and 137  $\text{kJ mol}^{-1}$ , Fig. 5). This result indicates that ideal zeolites would have the ability to create long anion-cation chains that stabilize charge-separated transition state structures, like those found in polyphosphobetaines with zwitterionic structures of alternating positive and negative charges in their polymeric chains.<sup>[53,54]</sup> Tailored design of such zeolite catalysts, with site-specific high Al content, would confer additional stability to charged transition states; this design motif could apply to any combination of zeolite framework topology and reactions with cationic intermediates, assuming the intermediates are large enough to interact concurrently with nearby sites.

Methanol dehydration barriers are lower on paired sites with co-adsorbed  $\text{CH}_3\text{OH}$  than on isolated sites (Fig. 5), but  $\text{CH}_3\text{OH}$  dimers form at higher pressures; therefore, the increases in  $k_{\text{zero}}$  suggest that similar promotional effects of site pairing are expected at high  $\text{CH}_3\text{OH}$  coverages. Two co-adsorbed  $\text{CH}_3\text{OH}$  molecules deprotonate Brønsted acid sites to form a protonated dimer complex (Scheme 1).<sup>[21,22,27,36,38,39]</sup> When located near the conjugate base of a methanol dehydration transition state, these protonated  $\text{CH}_3\text{OH}$  dimers interact with both deprotonated Al-centers simultaneously across the 6-MR (Fig. 6). For the prevailing concerted DME formation mechanism, the effective and intrinsic ( $\Delta G^\ddagger$  and  $\Delta G_{\text{act}}$ ) barriers are lower (112 and 119  $\text{kJ mol}^{-1}$ , Fig. 6c) for the NNN site-pair than for isolated sites (123



**Figure 6.** The most stable transition state structures for methanol dehydration in CHA for Step S1, Step S2, and Step C1 for NNN and NNNN site-pairs with protonated methanol dimers on the second site. Effective ( $\Delta G^\ddagger$  and  $\Delta H^\ddagger$ ) and intrinsic barriers ( $\Delta G_{\text{act}}$  and  $\Delta H_{\text{act}}$  in parenthesis and italicized) are shown in  $\text{kJ mol}^{-1}$ . Solid black lines indicate incipient and breaking bonds. Blue dashed lines indicate H-bonds (lengths in pm). (g) Interactions between alternating cations and anions from periodic boundary conditions in the Step C1 transition state on the NNNN site-pair. Alternate views given in Figures S14–S16 (SI).

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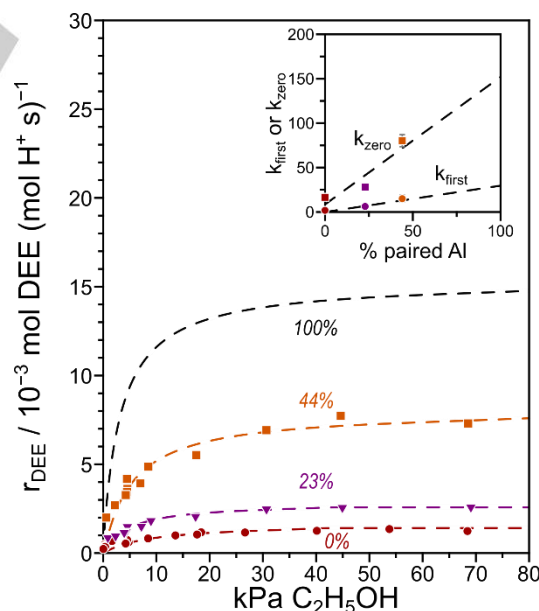


**Figure 7.** Reaction coordinate diagram for (a) sequential and (b) concerted ethanol dehydration to DEE on isolated sites (black), the NNN site-pair (red), and the NNNN site-pair (blue) with a  $C_2H_5OH$  monomer at the second acid site. Free energies (415 K, 1 bar) are shown relative to an ethanol monomer on all sites. Intrinsic barriers are shown for Steps S2 and Step C1 in parentheses. Structures of kinetically relevant states are shown in Figs. S21–S25 (SI).

and  $137 \text{ kJ mol}^{-1}$ , Fig. 4c). Similarly, the NNNN site-pair reduces  $\Delta G^\ddagger$  and  $\Delta G_{\text{act}}$  by 13 and  $28 \text{ kJ mol}^{-1}$ , respectively, than those of isolated sites. The “chain” of alternating charges observed in Fig. 4j persists here with a cationic  $CH_3OH$  dimer above the 6-MR and the cationic bimolecular transition state interacting across the 8-MR(2,4), resulting in lower barriers (Figs. 6f and 6g). Both NNN and NNNN configurations show significant ( $>10 \text{ kJ mol}^{-1}$ ) decreases in the intrinsic barriers ( $\Delta G_{\text{act}}$ ) for concerted DME formation even at these higher coverages (4  $CH_3OH$  per CHA cage), once again consistent, if overestimating, the estimated decrease in  $\Delta G_{\text{act}}$  ( $5 \text{ kJ mol}^{-1}$ ) obtained by changes in  $k_{\text{zero}}$  extrapolated from kinetic data.

We further illustrate how proximal sites enhance turnover rates by probing dehydration of a larger alcohol. Ethanol dehydration proceeds via pathways similar to methanol dehydration, but forms diethyl ether (DEE) and water. Ethanol can also dehydrate monomolecularly to form ethylene and water; however, DEE is the primary product formed at the experimental conditions studied here. Paired acid sites ethylate the zeolite with lower barriers ( $129$  and  $133 \text{ kJ mol}^{-1}$  for NNN and NNNN, Fig. 7) than isolated sites ( $138 \text{ kJ mol}^{-1}$ ) because ethanol at the proximal site facilitates an H-bonding interaction that stabilizes the conjugate base (Fig. S18, SI), analogous to the methanol case (Fig. 4). Effective and intrinsic barriers ( $\Delta G^\ddagger$  and  $\Delta G_{\text{act}}$ ) for concerted DEE formation also decrease from an isolated site ( $\Delta G^\ddagger = 98 \text{ kJ mol}^{-1}$ ;  $\Delta G_{\text{act}} = 120 \text{ kJ mol}^{-1}$ ) to an NNNN site-pair ( $\Delta G^\ddagger = 94 \text{ kJ mol}^{-1}$ ;  $\Delta G_{\text{act}} = 117 \text{ kJ mol}^{-1}$ , Fig. 7), for which the same interactions that stabilize DME formation transition states prevail. For ethanol dehydration near co-adsorbed ethanol dimers, barriers for the concerted reaction slightly increase for AI in NNN arrangements, but decrease for AI in NNNN arrangements (Section S10, SI). These DFT data suggest that ethanol dehydration should occur at faster

rates on paired than isolated sites in CHA. Indeed, experimental kinetic data show that CHA zeolites with higher percentages of 6-MR paired sites have higher ethanol dehydration turnover rates (per  $H^+$ , Fig. 8). Notably, DEE formation on CHA zeolites is not inhibited at high pressures like methanol,<sup>[27]</sup> and the rate of DEE formation is described by:



**Figure 8.** Turnover rate ( $r_{\text{DEE}}$ ) of diethyl ether (DEE) formation (415 K, per  $H^+$ ) as a function of ethanol pressure on H-CHA with 0% (●, red), 23% (▼, purple), and 44% (■, orange) of AI in paired configurations. Dashed lines represent regression to Eq. S23 (SI). Inset:  $k_{\text{zero}}$  ( $10^{-4} \text{ mol DEE (mol H}^+ \text{s kPa)}^{-1}$ ) and  $k_{\text{first}}$  ( $10^{-4} \text{ mol DEE (mol H}^+ \text{s)}^{-1}$ ) in Eq. 2 as functions of the fraction of paired AI.



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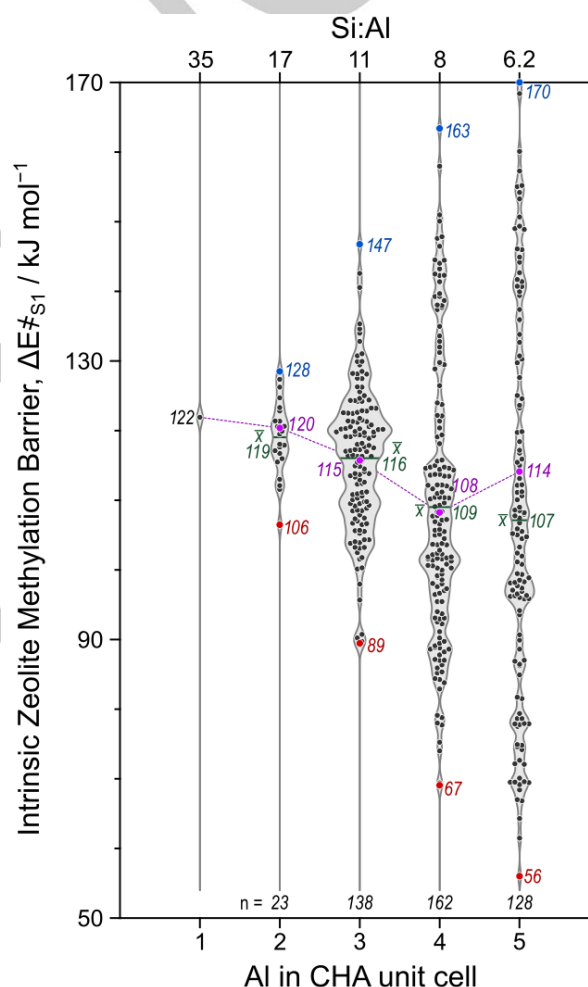
$$r_{DEE} = \frac{k_{first} P_E}{1 + \frac{k_{first}}{k_{zero}} P_E} \quad (2)$$

where  $P_E$  is the ethanol pressure (derivation in Section S11, SI). This indicates that, unlike methanol dehydration, ethanol dehydration transition states are not comprised of more molecules than the MASI at any pressures studied here. Measured first- and zero-order rate coefficients increase by 5x as 6-MR paired sites increase from 0% to 44% (Fig. 8). First- and zero-order rate coefficients (Eq. 2) fit to the kinetic data predict rate coefficients that are 20x larger on paired than on isolated sites based on extrapolation to the 100% paired Al limit (Fig. 8). These increases in  $k_{first}$  and  $k_{zero}$  reflect lower effective barriers by 11 and 10 kJ mol<sup>-1</sup> for first- and zero-order regimes on paired than isolated sites, comparable to differences observed for methanol dehydration between paired and isolated sites.

DFT-calculated free energy barriers indicate that H-bonding interactions between 6-MR paired Al sites stabilize cationic transition state structures more than their reactant precursors, even when those precursors are themselves cationic, resulting in an increase in both first- and zero-order rate coefficients. These interactions between proximal sites are seldom considered, yet often present in Brønsted acid-catalyzed reactions in zeolites, especially at low Si:Al ratios. Critically, we find that such inter-site communication via H-bonding is strengthened by co-adsorbed alkanols compared to a bare proton site (discussed in Section S7, SI), regardless of the size of the alkyl moiety. Such interactions resemble those observed during reactions in zeolites when capillary condensation occurs, wherein intrapore solvents can surround and interact with guest species to alter reaction mechanisms,<sup>[17,55]</sup> affect barriers,<sup>[56,57]</sup> or stabilize charged intermediates.<sup>[16,58]</sup> During alkanol dehydration on paired sites, however, the stabilizing interaction does not require a dense phase, because H-bonding interactions among polar adsorbates enable communication between two proximal Al sites. Zeolite frameworks have long been considered to behave as a pseudo-solvent, given their ability to stabilize reacting intermediates and transition states through non-specific (e.g., dispersive) and specific (e.g., H-bonding) interactions.<sup>[10,42,59–61]</sup>

Al centers in the zeolite framework not only generate proton active sites, but also influence the ionic properties of the structured solvent provided by the zeolite framework. Both effects can influence reactivity, just as altering the pH or the ionic strength of a bulk solvent can alter rates of homogeneous reactions.<sup>[62]</sup> Al centers in zeolites, however, are held within a rigid crystalline framework, and thus anisotropically position anionic charges in the solvating environment, unlike in liquid phases. Thus, the effect of a spectating Al center depends on its position relative to the active Al center in a zeolite framework. To investigate this in more detail, we consider the case of zeolite methylation at the O3 oxygen of an active site (i.e., Step S1). This step was examined in the absence of co-adsorbed methanol with increasing site density (1–5 Al per unit cell; Si:Al=35–6.2, 452 total Al

arrangements) such that no Al are at NN positions which violate Löwenstein's rule.<sup>[43]</sup> An isolated site catalyzes this step with an intrinsic energy barrier ( $\Delta E^\ddagger_{S1}$ ) of 122 kJ mol<sup>-1</sup> (Fig. 9). Among the 23 different two-Al configurations examined,  $\Delta E^\ddagger_{S1}$  varies from 106 to 128 kJ mol<sup>-1</sup>, representing a decrease of up to 16 kJ mol<sup>-1</sup> or an increase of up to 6 kJ mol<sup>-1</sup> compared to the isolated Al case (Fig. 9). Importantly, these barriers do not correlate with any simple geometric descriptor such as Al–Al distance or Al–C distance (Fig. S29, SI); instead, proximal Al centers placed in specific arrangements can either raise or lower barriers. For example, Al placed across 4-MR structures from the reacting site increase barriers (consistent with weaker acids predicted by DPE for such arrangements)<sup>[20]</sup>, while those placed across 6- or 8-MR structures generally lower barriers (Fig. S31, SI).



**Figure 9.** Intrinsic potential energy barriers ( $\Delta E^\ddagger_{S1}$ ) for zeolite methylation (Step S1) occurring on O3 of the A site with 1–5 Al in the CHA unit cell (Si:Al = 6.2–35). For each Al content, the minimum (red), maximum (blue), average ( $\bar{x}$ , green), and effective (purple, assuming Al occupy sites according to a Boltzmann distribution at 415 K) barriers are labeled, along with the total number of Al arrangements studied ( $n$ ; discussed in Section S12, SI).

With each addition of a proximal site, barriers shift by as much as  $\pm 37$  kJ mol<sup>-1</sup>, resulting in barriers as low as 56 kJ mol<sup>-1</sup> or as high



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as 170 kJ mol<sup>-1</sup> (Fig. 9; Si:Al=6.2). These shifts in barriers are nearly symmetric, such that the average barrier decreases slightly to 107 kJ mol<sup>-1</sup> at the highest Al content (Fig. 9). If Al are assumed to be distributed thermodynamically (i.e., based on a Boltzmann average), then the effective barrier can be computed for each Si:Al ratio. This Boltzmann-averaged barrier decreases nearly monotonically as Al content increases, to a minimum of 108 kJ mol<sup>-1</sup> at Si:Al of 8 and 114 kJ mol<sup>-1</sup> at Si:Al of 6.2 (Fig. 9). These spectating Al are not directly involved in the methylation transition state; rather, they modify the solvation environment provided by the zeolite by altering its polarity and providing H-bonding sites that interact with the transition state or conjugate base of the reacting site, in a manner sensitive to the specific positions of the Al centers, resulting in dramatic effects (both increases and decreases) in activation barriers.

## Conclusions

DFT calculations and experimental kinetic and spectroscopic data reveal that altering the proximity and arrangement of framework Al and their associated Brønsted acid sites modifies the solvating environments of zeolite pores, by altering their polarity and H-bonding capacity, so as to significantly increase or decrease activation barriers for zeolite-catalyzed reactions. These shifts in barriers cannot be predicted by Al–Al distances or Al densities alone, but depend strongly on the specific locations of spectating Al centers. For example, Al located across 4-MR in CHA generally result in weaker acids and higher activation barriers, whereas Al located across 6-MR or at certain 8-MR positions decrease activation barriers. Alkanol dehydration rates are enhanced on paired sites located in 6-MR of CHA, because H-bonding interactions are facilitated by co-adsorbed alkanols that stabilize the conjugate base of reacting sites to lower barriers. Such inter-site cooperation is distinct from that observed in dense intrapore phases, wherein solvent species act as proton shuttles or interact with co-adsorbates;<sup>[17]</sup> rather, these H-bonding interactions are mediated by co-adsorbed reactants without an intermediary solvent phase, thus requiring Al sites to be positioned in specific locations to accommodate the geometric limitations imposed by the size and structure of the reactant complex and the connecting H-bonds. Such interactions are likely sensitive to the topology of the zeolite framework; thus, these findings on CHA are expected to hold for small-pore zeolite frameworks with similar 6- and 8-MR motifs (e.g., LTA, AEI), but extending these conclusions to zeolites characterized by different topological features (e.g. MFI) requires further investigation. Additionally, our results indicate that further barrier reductions can occur with chains of alternating cationic and anionic charges that mimic long chains of alternating charges in some zwitterionic polymers.<sup>[53,54]</sup> These mechanistic insights into the specific interactions of proximal binding sites and co-adsorbates during Brønsted acid-catalyzed reactions in zeolites have gone previously unrecognized, and provide new targets for synthesizing zeolites with framework Al arrangements tailored to match the geometries of reactants and transition states to maximize H-bonding among these moieties.

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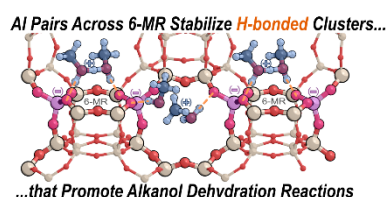
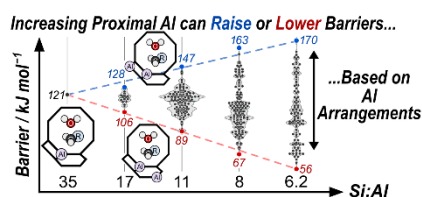
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Alexander J. Hoffman, Jason S. Bates,  
John R. Di Iorio, Steven V. Nystrom,  
Claire T. Nimlos, Rajamani Gounder\*,  
David Hibbitts\*

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**Proximal Al in Zeolites:** have a dramatic impact on activation barriers depending on their specific siting within 4-, 6-, and 8-MR motifs in the CHA zeolite. Alkanol dehydration turnover rates increase for paired Al in 6-MR because the Al stabilize one another via H-bonded cationic transition states and alkanol clusters.

**Rigid Arrangements of Ionic Charge in Zeolite Frameworks Conferred by Specific Al Distributions Preferentially Stabilize Alkanol Dehydration Transition States**