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Possible role of molecular clustering in single-file diffusion of mixed and pure gases in dipeptide nanochannels

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ABSTRACT

Single-file diffusion (SFD) of CO, CO₂, and CH₄ gas molecules confined in L-Ala-L-Val nanochannels as both pure and binary mixtures is investigated using pulsed field gradient (PFG) NMR at the high field of 17.6 T. A unique SFD mobility is observed for each of the sorbates as pure gases. As binary mixtures, both sorbates exhibit identical SFD mobilities. The mixture mobilities were observed to be larger than those of the slowest-diffusing component in its pure form under the conditions when the total concentration of sorbate molecules in the channels was the same or comparable. It is proposed that molecular clustering in the studied channels is responsible for the relationship experimentally observed between the one-component and mixture mobilities. It is demonstrated that molecular clustering can also explain the lack of the experimental observation of a transition from the SFD mechanism to the center-of-mass diffusion mechanism, viz. diffusion mechanism characterized by correlated movements of all molecules in any particular channel.

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1. Introduction

Single-file (SF) channels forbid molecules inside the channels from passing each other during a diffusion process by geometrically constraining them and, hence, conserving their relative order in the channels. A unique property of diffusion in sufficiently long SF channels is the growth of the mean square displacement (MSD) with the square root of time

$$\langle z^2(t) \rangle = 2Ft^{0.5}, \quad (1)$$

where F is the single-file mobility factor [1]. Diffusion in SF channels obeying Eq. (1) is known as single-file diffusion (SFD). In finite SF channels, however, the fastest and rate-determining diffusion mechanism in the limit of long observation times is related to correlated displacements of all molecules in each channel [1]. The direction of such correlated displacement at any particular time is expected to be independent of any previous displacement resulting

in the applicability of the Einstein relation, which requires a proportionality between the MSD and diffusion time. The diffusion associated with this type of displacements of chains of molecules is known as center-of-mass diffusion (COM). Theoretical and computational studies predict a transition from the SFD to COM diffusion regime, which is marked by a shift in the MSD dependence on diffusion time, at sufficiently large displacements [2–4]. However, until now, such a transition has not been observed experimentally.

We have recently reported the first experimental study of SFD of molecular mixtures [5,6]. The combination of high magnetic field (17.6 T) and large magnetic field gradients (up to 23 T/m) in this pulsed field gradient (PFG) NMR study allowed for measurement of MSDs of CO, CO₂, and CH₄ molecules on a microscopic length scale as a function of diffusion time. Self-assembled L-alanyl-L-valine (AV) nanochannels were selected as model SF channels for the SFD observation of these gas molecules since the radius, r , of AV channels ($r \sim 0.27$ nm) is smaller than the kinetic diameters, d , of these molecules ($d_{\text{CO}} \sim 0.34$ nm, $d_{\text{CO}_2} \sim 0.33$ nm, $d_{\text{CH}_4} \sim 0.41$ nm) [7,8]. In the current paper, we provide a review of our SFD data for this sorbate/sorbent system together with the analysis of the relationship between the observed single-file mobilities of the pure gases and

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their corresponding binary mixtures. We also discuss and explain the absence of a transition from the SFD to COM regime, which is predicted by the previously reported model [2,3] to occur under our experimental conditions.

2. Experimental

L-alanyl-L-valine crystallizes to form hydrophobic channels that have a helical structure. For this study, AV was purchased commercially from Bachem and used without further purification. The average length (l) of the AV nanochannels was determined to be around 50 μm from scanning electron microscopy images [9]. A fixed amount of AV crystals (~55 mg) was loaded into five-millimeter medium wall NMR tubes (Wilmad-LabGlass) that can sustain pressures up to 20 bar. The tubes were connected to a custom-made vacuum system and the samples were activated overnight at an elevated temperature of 373 K under high vacuum, i.e. at pressures below 0.001 mbar. Following activation, a measured mass of either pure gas or gas mixture was cryogenically condensed through the vacuum system and into the NMR tubes.

PFG NMR diffusion measurements were performed using a 17.6 T Bruker BioSpin NMR spectrometer operating at the ^{13}C resonance frequency of 188.8 MHz. The diff60 diffusion probe was used to generate gradients in the range of 0.3–23 T/m. The 13-interval bipolar PFG NMR sequence with eddy current delays between 3.5 and 5 ms and sine-shaped gradients with the effective duration of 1.5–2.2 ms were used for all diffusion measurements [6,10,11]. This sequence was also used to estimate T_2 and T_1 ^{13}C NMR relaxation times for each sorbate by systematically changing the time interval between the first and second $\pi/2$ rf pulse and the second and third $\pi/2$ rf pulse, respectively, while keeping all other intervals constant. The ^{13}C T_2 relaxation times of adsorbed CO, CO₂, CH₄ molecules were estimated to be around 16, 18, and 22 ms, respectively. The corresponding T_1 times were determined to be around 0.6, 2.7, and 1.5 s. All measurements were performed at 298 K.

The total pressure of the gas phase loaded into the five NMR tubes was around 10 bar at 298 K. In mixture samples, the partial pressure of each sorbate was 5 bar. Concentrations were obtained from the analysis of the recorded NMR spectra, as discussed in our previous work [5,6]. The measured concentration of each sorbate in each sample is presented in Table 1. This table shows that the total gas concentration in each of the samples is comparable, within the experimental uncertainty. This is significant as it allows us to directly compare the rates of diffusion of these molecules in single-sorbate and mixture samples.

Diffusion data is obtained from the change in PFG NMR signal intensity, which corresponds to the area under NMR lines, measured as a function of the applied magnetic field gradient while keeping all other parameters constant. The attenuation of the PFG NMR signal ($\Psi \equiv \frac{S(g)}{S(g=0)}$) is plotted against $q^2 t^{0.5}$, where q is equal to $2\gamma g \delta$, γ is the gyromagnetic ratio, g is the gradient strength, δ is the effective gradient duration, and t is the diffusion time. Given the helical nature of the AV nanochannels and their random orientation

in the PFG NMR sample, the expectation of the signal attenuation for sorbates exhibiting SFD is given by:

$$\begin{aligned} \psi(q, t) &= \sqrt{\frac{p^2 \pi}{2q^2 \langle z^2(t) \rangle_{\text{helix}}}} \text{erf} \left(\sqrt{\frac{q^2 \langle z^2(t) \rangle_{\text{helix}}}{2p^2}} \right) \\ &= \sqrt{\frac{p^2 \pi}{4q^2 F_{\text{helix}} t^{0.5}}} \text{erf} \left(\sqrt{\frac{q^2 F_{\text{helix}} t^{0.5}}{p^2}} \right) \end{aligned} \quad (2)$$

where p is a geometrical factor that relates the actual MSD, $\langle z^2(t) \rangle_{\text{helix}}$, of molecules along the helical diffusion path with the observed MSD, $\langle z^2(t) \rangle$, and is equal to 1.3 [1,6,12]. F_{helix} denotes the actual single-file mobility factor along the helical path of the AV channels. The dependence of the signal attenuation on q^2 in Eq. (2) is different from the monoexponential dependence expected for normal diffusion without any diffusion anisotropy because this equation was developed for one-dimensional diffusion assuming a random distribution over the channel directions in a sample.

3. Results and discussion

The measured PFG NMR attenuation curves for diffusion of CO, CO₂, and CH₄ in single-sorbate and mixture samples, as shown in Fig. 1, are scaled to test the agreement of the data measured at different diffusion times with Eq. (2) developed for SFD. The agreement with this equation, and thus the time scaling of MSD (Eq. (1)), is demonstrated by the observation that, for each sorbate and sample, the attenuation curves corresponding to different diffusion times coincide within the experimental uncertainty for a broad range of times used in the measurements.

For each gas in the single-component samples, a least-squares fit of Eq. (2) to all the attenuation curves in Fig. 1a measured for this gas at different diffusion times yielded a unique value of the single-file mobility, F_{helix} , which is presented in Table 2. Fig. 1b and c show coincidence of the attenuation curves of both sorbates in each of the two mixture samples, within the experimental uncertainty. Hence, a single least-squares fit by Eq. (2) is used to describe the attenuation curves for both molecules in each of the mixtures. The corresponding values of the single-file mobility of the mixture are shown in Table 2. The identical rates of transport observed for both sorbates in a mixture provides further confirmation for the SFD mechanism because this mechanism forbids the mutual passages of molecules in the channels. The small deviations observed between the experimental data and the least-squares fits (Fig. 1) can be assigned to differences between the actual channel orientation distribution and the completely random distribution assumption used to derive Eq. (2).

Fig. 2 a,b compares the time dependencies of MSDs obtained using Eq. (2) from the best-fits of the individual attenuation curves at every diffusion time for each sorbate in the single-sorbate and its corresponding mixture samples. The solid lines in the figure were obtained by fitting the time dependencies of MSD to Eq. (1), where $\langle z^2(t) \rangle_{\text{helix}}$ and F_{helix} are substituted for $\langle z^2(t) \rangle$ and F . The values of F_{helix} resulting from these MSD fits corroborate those reported in Table 2, within limits of the experimental error.

Table 1
Gas concentration (mol/kg of AV) determined for each sample.

	CO	CO ₂	CH ₄	Total concentration in mixture samples
Single- Sorbate	0.56 ± 0.06	0.52 ± 0.05	0.48 ± 0.05	—
CO + CH ₄	0.34 ± 0.03	—	0.20 ± 0.02	0.54 ± 0.02
CO + CO ₂	0.24 ± 0.02	0.34 ± 0.03	—	0.58 ± 0.03

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