

Supporting Information For

Molecular insights into the Microscopic Behavior of CO₂ Hydrates in Oceanic Sediments: Implications for Carbon Sequestration

Fengyi Mi^{†,‡}, Wei Li[†], Jiangtao Pang[†], Othonas A. Moutos[‡], Fulong Ning^{*,†}, Thijs J.H. Vlugt^{*,‡}

[†]National Center for International Research on Deep Earth Drilling and Resource Development, Faculty of Engineering, China University of Geosciences, Wuhan, Hubei 430074, China

[‡]Engineering Thermodynamics, Process & Energy Department, Faculty of Mechanical Engineering, Delft University of Technology, Leeghwaterstraat 39, Delft, 2628CB, the Netherlands

*Authors to whom correspondence should be addressed. nflzx@cug.edu.cn, t.j.h.vlugt@tudelft.nl

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Video S1. Growth process of CO₂ hydrate in oceanic sediments for the H_{3%}Urea system.

Video S2. Dissociation process of CO₂ hydrate in oceanic sediments for the H_{3%}Urea system.

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Supporting Reference

Additional Details for Simulation Models and Methods

Simulation Models

The illite unit cell was obtained by modifying the pyrophyllite unit cell taken from the United States Mineral Crystal Structure Database¹. To construct the illite crystal structure with the chemical formula $K_1(\text{Si}_7\text{Al})\text{Al}_4\text{O}_{20}(\text{OH})_4$, a Si atom in the pyrophyllite unit cell was substituted with an Al atom. An illite layer is created by replicating the illite unit cell ($24 \times 8 \times 1$). An illite nanopore consists of two identical illite layers, and the nanopore size was set to 46 Å. CO_2 hydrate belongs to SI-type hydrate, mainly composed of 5^{12} and $5^{12}6^2$ cages. A standard CO_2 hydrate crystal ($2 \times 6 \times 4$) was inserted into the middle region of the illite nanopore. The CO_2 hydrate crystal contains 288 CO_2 and 1656 H_2O molecules, and the size was $24.3 \times 72.3 \times 36.3$ Å in the xyz directions. An aqueous homogeneous solution was also inserted into the illite nanopore. The homogeneous solution contains 1277 CO_2 , 7340 H_2O , 82 Na^+ , 82 Cl^- , and 192 K^+ . 82 Na^+ and Cl^- resulting in a salinity of 3.5 wt%, which is close to real seawater. Two layers of illite clay generate 384 negative charges through isomorphic substitution. Therefore, we added 384 K^+ to compensate for the negative charge generated by the illite clay surfaces. It is worth noting that the 384 K^+ are evenly divided into two parts: (1) the homogeneous solution and (2) outside the illite nanopore. This avoids extreme salt concentrations in the illite nanopore. A 5 Å buffer layer was placed between the CO_2 hydrate crystal and the illite clay layer, which eliminates the interaction between the illite surface and the CO_2 hydrate crystal in the initial configuration. Different numbers of urea molecules were inserted into the illite nanopores to represent different urea concentrations (1.6wt%, 3 wt%, 4.5 wt%, and 6 wt%). This allows for a precise analysis of the interaction between urea and geofluid. Too high a urea concentration will result in an aggregate phase between urea and geofluid, while too low a concentration will not reveal the role of urea in the growth and dissociation of CO_2 hydrates, and mechanical instability of CO_2 hydrate-illite interface in the brine-urea-illite system. A system without urea molecules was simulated as a control system. Each simulation system contains about 60,000 atoms. The dissociation of CO_2 hydrate was simulated by increasing the temperature to 307 K. High temperatures can promote hydrate dissociation in a short time and save calculation costs². All force field parameters for H_2O , CO_2 , ions, illite, and urea molecules in the systems are shown in Table S2.

Table S1. Number of molecules of each species for the five different simulations.

System	simulation systems in illite silt-nanopore					
	N_{CO_2}	$N_{\text{H}_2\text{O}}$	N_{K^+}	N_{Na^+}	N_{Cl^-}	N_{urea}
H _{0%} Urea	1565	8996	384	82	82	0
H _{1.6%} Urea	1565	8996	384	82	82	64
H _{3%} Urea	1565	8996	384	82	82	127
H _{4.5%} Urea	1565	8996	384	82	82	191
H _{6%} Urea	1565	8996	384	82	82	254

Table S2. Parameters for the TIP4P/ice water model³, TraPPE CO₂⁴, OPLS-AA urea⁵, and the CLAYFF force field⁶. σ and ε are the Lennard-Jones parameters, in units of nm and kJ/mol, respectively; q is the partial charge of an atom in units of elementary charge (e); m is the atomic mass in units of g/mol.

atom	ε / [kJ/mol]	σ / [nm]	q / [e]	m / [g/mol]
H ₂ O				
O (MW)	0	0	-1.1794	0
O	0.8822	0.31668	0	16
H	0	0	0.5897	1.008
CO ₂				
C	0.224478	0.28	0.70	12.011
O	0.656806	0.305	-0.35	15.9994
O (OM)	0	0	0	0
Ion				
Cl	0.418998	0.439997	-1.0	35.453
Na	0.544572	0.235002	1.0	22.989
K	0.41858	0.333401	1.0	39.0983
Illite				
Si (st)	7.70065×10^{-6}	0.3302	2.1	28.09
Al (ao)	5.56388×10^{-6}	0.4271	1.575	26.98
Al (at)	7.70065×10^{-6}	0.3302	1.575	26.98
O (ob)	0.65017	0.316556	-1.05	16
O (obts)	0.65017	0.316556	-1.1688	16
O (oh)	0.65017	0.316556	-0.95	16
H (ho)	0	0	0.425	1.008
Urea				
C	0.43932	0.375	0.142	12.011
O	0.87864	0.296	-0.39	15.999
N	0.71128	0.325	-0.542	14.007
H	0	0	0.333	1.008

Wall Methods

In the simulation box, the upper and lower clay layers will interact (*i.e.*, stacking) with each other due to the periodicity in the z direction, which will inevitably affect the structure and stability of the clay layer and even the surface properties of the illite. Therefore, we do not set the periodicity in the z direction and use two layers of virtual walls to limit interactions between the upper and lower clay layers. The parameters of the atoms in the virtual walls are as follows: $\rho_w = 60$ atoms/nm³, $\epsilon_w = 1.13$ kJ/mol, $\sigma_w = 0.37$ nm, and a 9-3 Lennard-Jones potential. The atom type on the virtual wall is the C atom with a charge of 0. In the initial configuration, the length of the simulation box in the z direction is large enough so that there is sufficient distance between the walls and the clay layers. On the isothermal-isobaric NPT ensemble, we use a semi-isotropic pressure scaling method to achieve independent scaling in the xy and z directions⁷. The pressure scaling factor in the xy direction is set to 0, and the pressure scaling factor in the z direction is set to 4.5×10^{-5} . The simulation box only compresses the z direction to achieve the given pressure value. In all simulations, the distance between the wall and the solution was larger than 10 Å, which did not affect the simulation results.

Pull Methods

We apply a constant force to the atoms on the upper illite layer in the z - z direction for the tensile and compressive simulations. The surface area of the clay layer is about 89.09 nm². Therefore, pressure will be generated in the illite nanopores. The constant pull rate is 10 Å/ns. The simulation time is 1 ns, with the trajectory is saved every 1 ps. The illite clay layer is incompressible in the xy direction, and the force acts in the z direction. Therefore, the NPT ensemble cannot precisely control the pressure value by compressing the box size. In this study, we use the NVT system to control the temperature and use the external force to change the pressure. As a result, an NVT ensemble was adopted to perform deformation simulations, with the temperature controlled by the Nosé-Hoover⁸ thermostat at 273.15 K. In the deformation period, the interactions of urea molecules, water molecules, CO₂ molecules, and ions in the nanopores will deform CO₂ hydrate crystals.

Simulation Methods

The geometry of the H₂O molecules is maintained by applying the SETTLE algorithm⁷. The Lorentz-Berthelot mixing rules are used for describing the unlike interactions. The equations of motion were integrated according to the leapfrog algorithm with a 2.0 fs timestep. The well depth $\epsilon_{O(CO_2)-O(H_2O)}$ between the oxygen in CO₂ and the oxygen in H₂O was scaled by a factor of 1.08. The initial configurations were energy minimized by using the steepest descent algorithm. A pre-equilibrium simulation was performed for 2 ns, and the temperature was set to 250 K by controlling the v-rescale⁹ algorithm, and the pressure was 500 bar by the Berendsen¹⁰ algorithm. Growth simulations were performed in the isothermal-isobaric NPT ensemble for 2.0 μs at 250 K and 500 bar, and the temperature and pressure were changed to the Nosé-Hoover⁸ thermostat and Parrinello-Rahman¹¹ barostat,

respectively. At the end of the growth simulation, the dissociation simulation was performed in the isothermal-isobaric NPT ensemble for 10 ns at 307 K and 100 bar. The temperature and pressure were also changed to the Nosé-Hoover⁸ thermostat and Parrinello-Rahman¹¹ barostat, respectively. At the end of the growth simulation, the tension and compression simulations were performed in the NVT ensemble for 1 ns. To monitor the growth and dissociation of CO₂ hydrates, and mechanical instability of CO₂ hydrate-illite interface in the brine-urea-illite system, the cage analysis algorithm proposed by Jacobson *et al.*¹² is used to identify the seven cage types (5^{12} , $5^{12}6^2$, $5^{12}6^3$, $5^{12}6^4$, $4^15^{10}6^2$, $4^15^{10}6^3$, and $4^15^{10}6^4$). CO₂ hydrate belongs to SI-type hydrate, mainly composed of 5^{12} and $5^{12}6^2$ cages, and its ratio is 1:3.

Calculation principle of the residence time correlation function

The principle of correlation functions is well established and documented in GROMACS 2022⁷. The residence time correlation function is one such function. The definition of the residence correlation function $C_f(t)$ for residence $f(t)$ is:

$$C_f(t) = \langle f(\xi)f(\xi+t) \rangle_{\xi} \quad (1)$$

$\langle \rangle$ indicates averaging over ξ , which is over time origins.

The residence time τ is calculated by the numeric integration of the residence correlation function:

$$\tau = \int_0^{\infty} C_f(t) dt \quad (2)$$

In MD simulations, correlation functions are calculated based on data points with discrete time intervals Δt , so the residence correlation function $C_f(t)$ is:

$$C_f(j\Delta t) = \frac{1}{N-j} \sum_{i=0}^{N-1-j} f(i\Delta t)f((i+j)\Delta t) \quad (3)$$

Where N is the number of available time frames for the calculation.

In this study, each τ_{Res} is calculated within 1 ns for CO₂, H₂O, and ions near urea molecules. The GROMACS command `gmx_mpi hbond -ac -contact` can perform the calculation of the residence correlation function.

Calculation principle of the CO₂ hydrate growth rate (R_{HG})

To monitor CO₂ hydrate growth kinetics, the CO₂ hydrate growth rate (R_{HG}) is defined. The F_4 order parameter serves as an effective discriminator for distinguishing the water phase, with average values of -0.04, -0.4, and 0.7 for liquid water, ice, and hydrate, respectively¹³. The F_4 order parameter is defined by

$$F_4 = \langle \cos(3\emptyset) \rangle \quad (4)$$

The value of F_4 is computed by the water-water pair as a function of the torsional angle \emptyset between oxygen atoms within 3.5 Å and the outermost hydrogen atoms in the water-water pairs. The R_{HG} value can be used for

monitoring CO₂ hydrate growth kinetics. The R_{HG} is calculated as follows:

$$R_{HG} = \frac{F_4(t+\Delta t) - F_4(t)}{\Delta t} \quad (5)$$

The noise reduction (*i.e.*, smoothing) was used for F_4 function. The slope of the F_4 curve is CO₂ hydrate growth rate R_{HG} .

Calculation principle of the Increments of the number of hydrogen bonds per square nanometer ($\Delta Hbonds$)

To assess the changes in the number of hydrogen bonds near the upper illite surface, we define the Increments of the number of hydrogen bonds per square nanometer ($\Delta Hbonds$). This increment of the number of hydrogen bonds per square nanometer as a function of mechanical tensile/compressive strain (ϵ) is:

$$\Delta Hbonds = \frac{N_{H-bonds} - N_0}{S_{illite\ surface}} \quad (6)$$

Where $N_{H-bonds}$ is the number of hydrogen bonds between urea/H₂O molecules and the upper illite surface under tensile/compressive strains (ϵ). N_0 is the number of hydrogen bonds between urea/H₂O molecules and the upper illite surface at $\epsilon = 0$. In this study, $S_{illite\ surface}$ is the surface area of the upper surface of illite, which is 89.09 nm².

Calculation principle of the number of hydrogen bonds at different distances from the upper surface (HCD)

We calculated the number of hydrogen bonds at different distances from the upper illite surface. These hydrogen bonds are mainly formed between water molecules. The number of hydrogen bonds was calculated for every 2 Å from the upper illite surface. The VMD software¹⁴ was used to calculate hydrogen bonds. The cut-off length and angle are set to 3.5 Å and 60° respectively in the VMD software.

Supporting Figures

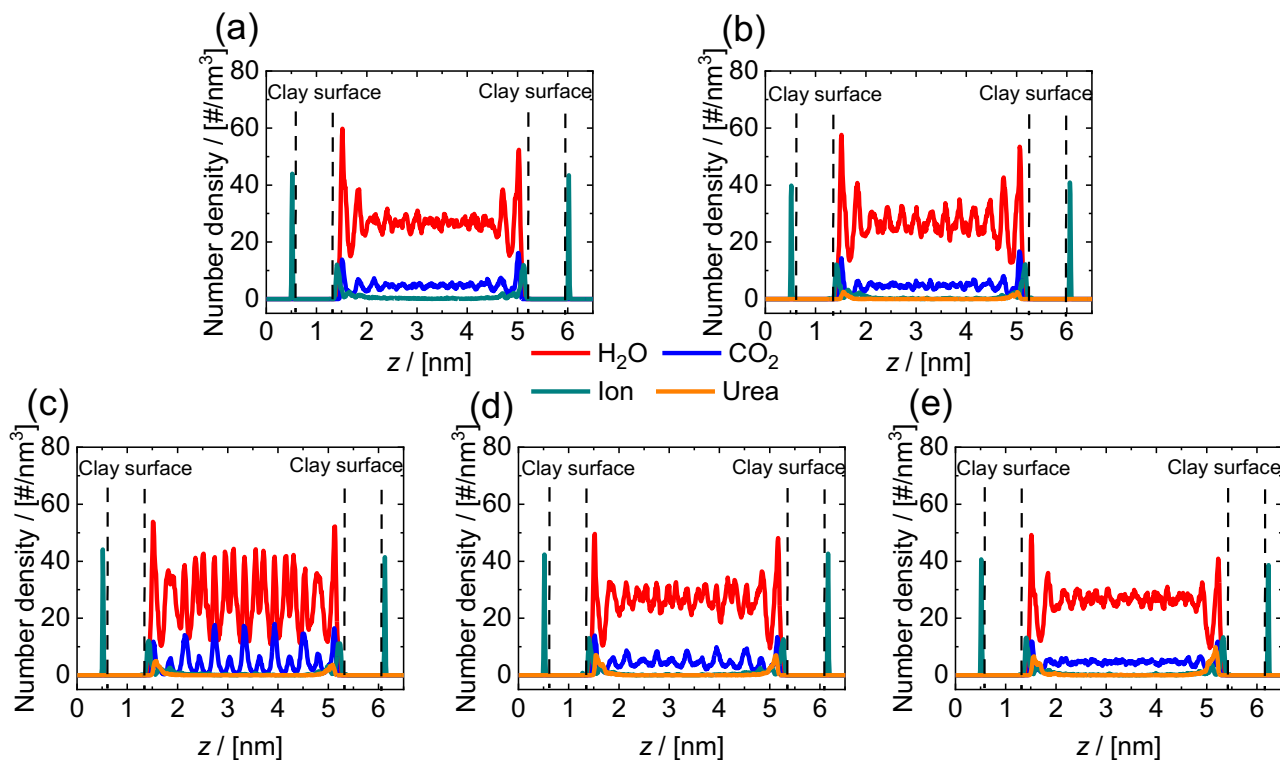


Figure. S1. Number density distribution of H₂O, CO₂, ions, and urea molecules along the z-axis direction for 1.95 - 2.0 μ s for the (a) H₀%Urea, (b) H_{1.6}%Urea, (c) H₃%Urea, (d) H_{4.5}%Urea, and (e) H₆%Urea systems.

We found that the fluctuations of H₂O and CO₂ molecules of the H₃%Urea system are larger than those of other systems, which is attributed to the location difference of CO₂ hydrate. At the end of the simulation, standard CO₂ hydrate crystals are parallel to the xy plane and perpendicular to the z direction in the H₃%Urea system (Figure S16(c)). Sharp, the CO₂ hydrate crystals in other systems are not perpendicular to the z direction (Figure S16(a-e)). Therefore, the number density of H₂O and CO₂ molecules in the z-direction in the H₃%Urea system will fluctuate more than for other systems.

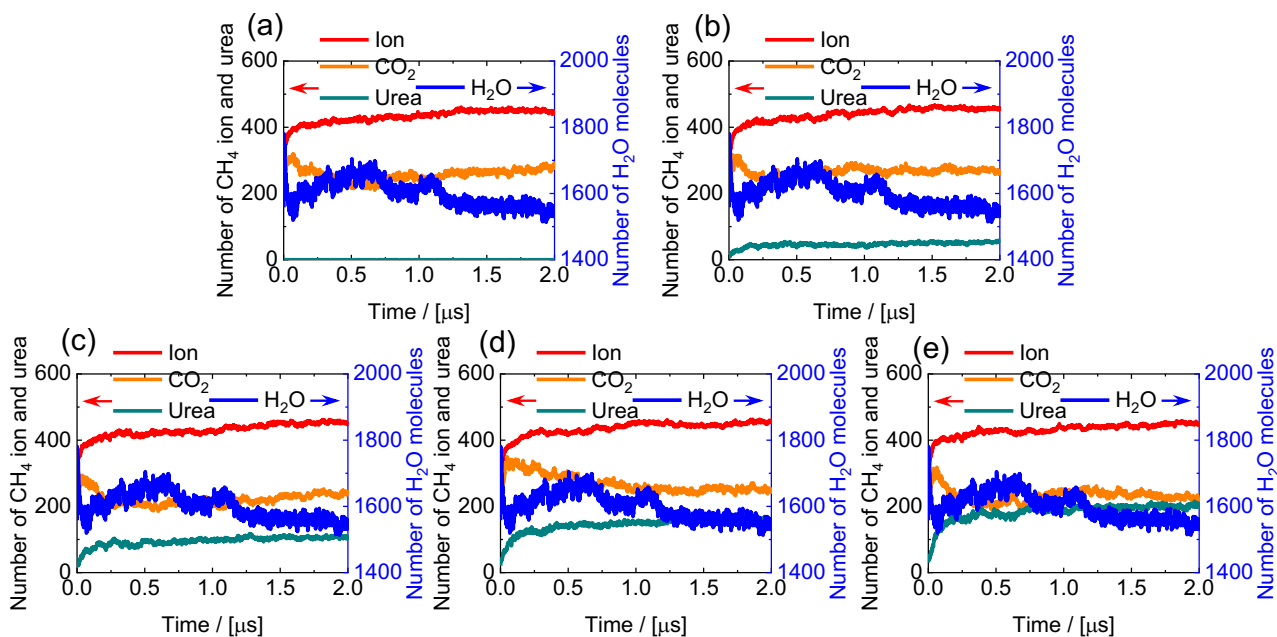


Figure S2. Evolution of the number of ions, CO₂, H₂O, and urea molecules near the illite surface within 5 Å in the (a) H_{0%}Urea, (b) H_{1.6%}Urea, (c) H_{3%}Urea, (d) H_{4.5%}Urea, and (e) H_{6%}Urea systems.

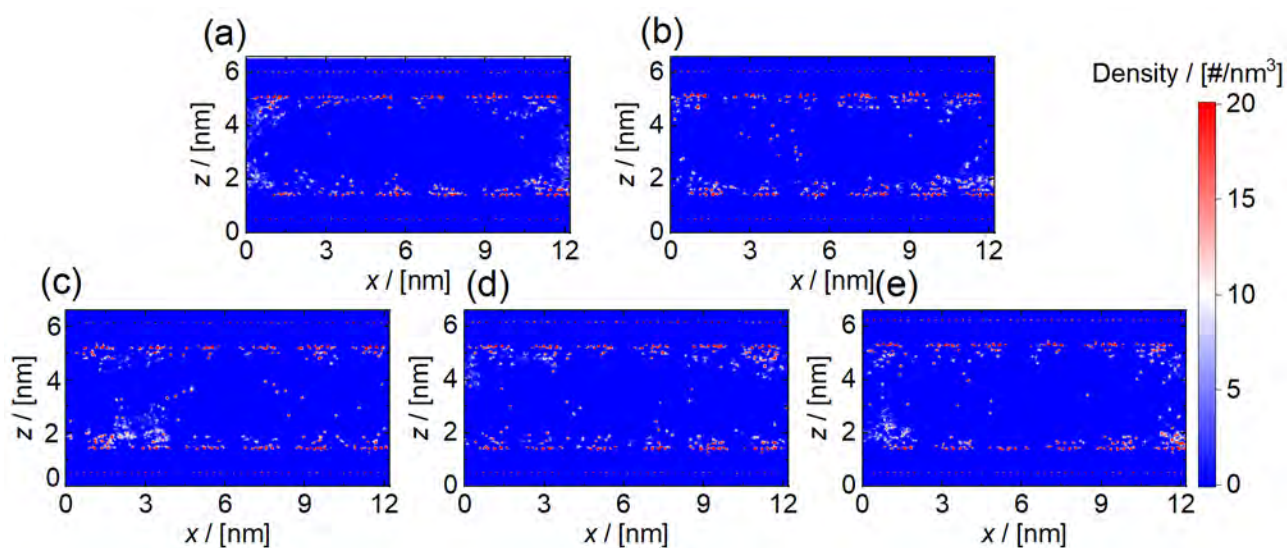


Figure S3. Number density distribution of ions for the last 0.01 μs in the (a) H_{0%}Urea, (b) H_{1.6%}Urea, (c) H_{3%}Urea, (d) H_{4.5%}Urea, and (e) H_{6%}Urea systems.

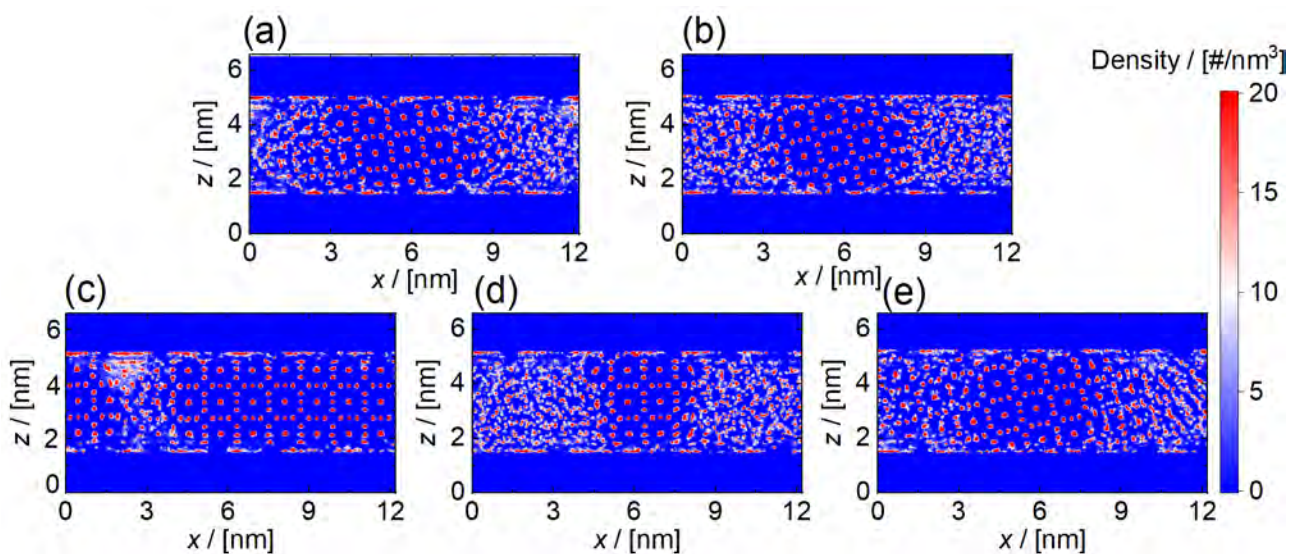


Figure S4. Number density distribution of CO₂ molecules for the last 0.01 μ s in the (a) H_{0%}Urea, (b) H_{1.6%}Urea, (c) H_{3%}Urea, (d) H_{4.5%}Urea, and (e) H_{6%}Urea systems.

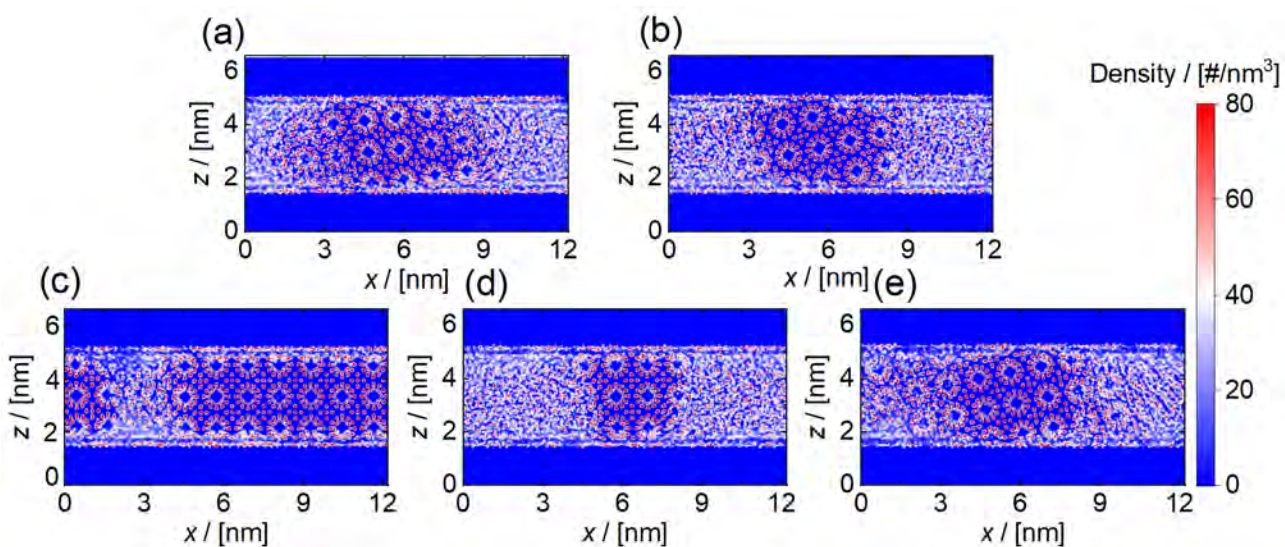


Figure S5. Number density distribution of H₂O molecules for the last 0.01 μ s in the (a) H_{0%}Urea, (b) H_{1.6%}Urea, (c) H_{3%}Urea, (d) H_{4.5%}Urea, and (e) H_{6%}Urea systems.

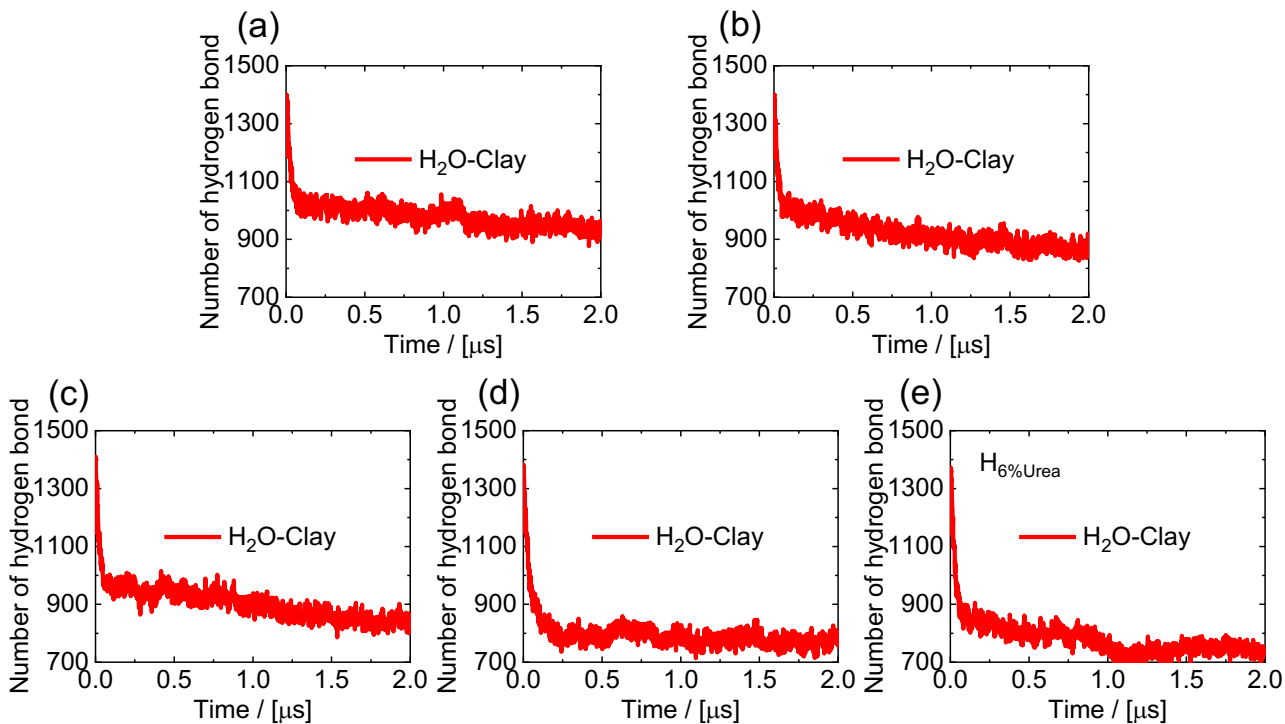


Figure S6. Evolution of the number of hydrogen bonds between the illite surface and H₂O molecules in the (a) H₀%Urea, (b) H_{1.6}%Urea, (c) H₃%Urea, (d) H_{4.5}%Urea, and (e) H₆%Urea systems.

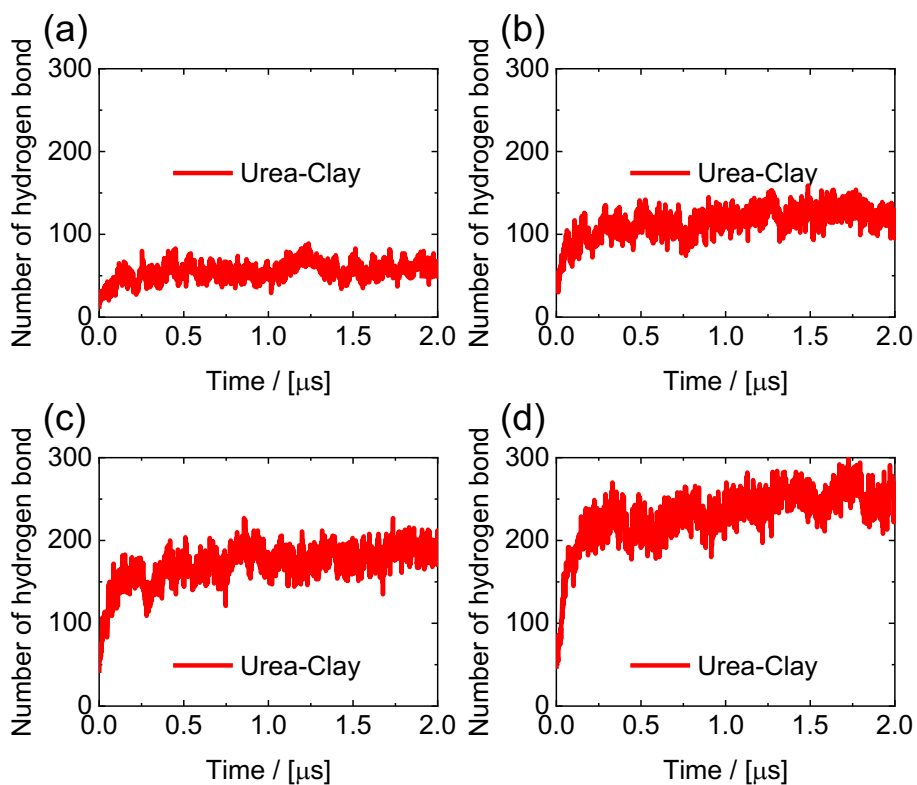


Figure S7. Evolution of the number of hydrogen bonds between the illite surface and urea molecules in the (a)

H_{1.6%}Urea, (b) H_{3%}Urea, (c) H_{4.5%}Urea, and (d) H_{6%}Urea systems.

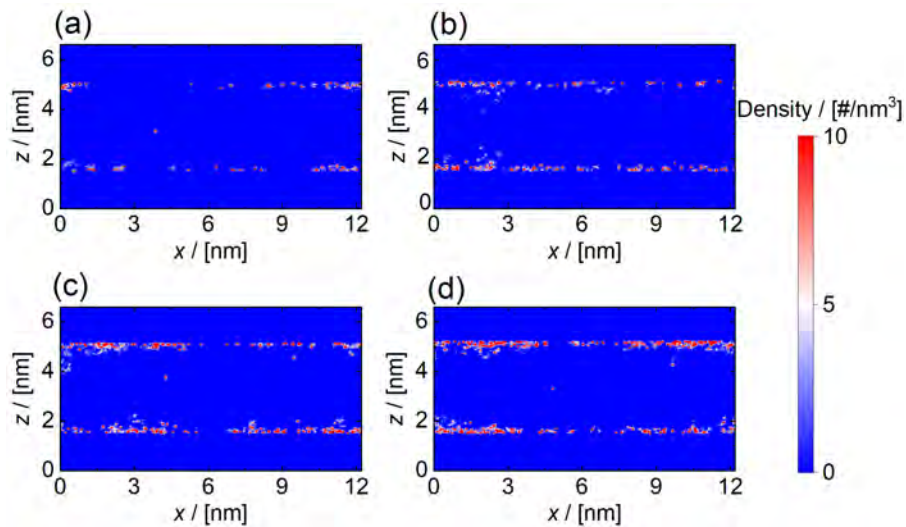


Figure S8. Number density distribution of urea molecules for the last 0.01 μ s in the (a) H_{1.6%}Urea, (b) H_{3%}Urea, (c) H_{4.5%}Urea, and (d) H_{6%}Urea systems.

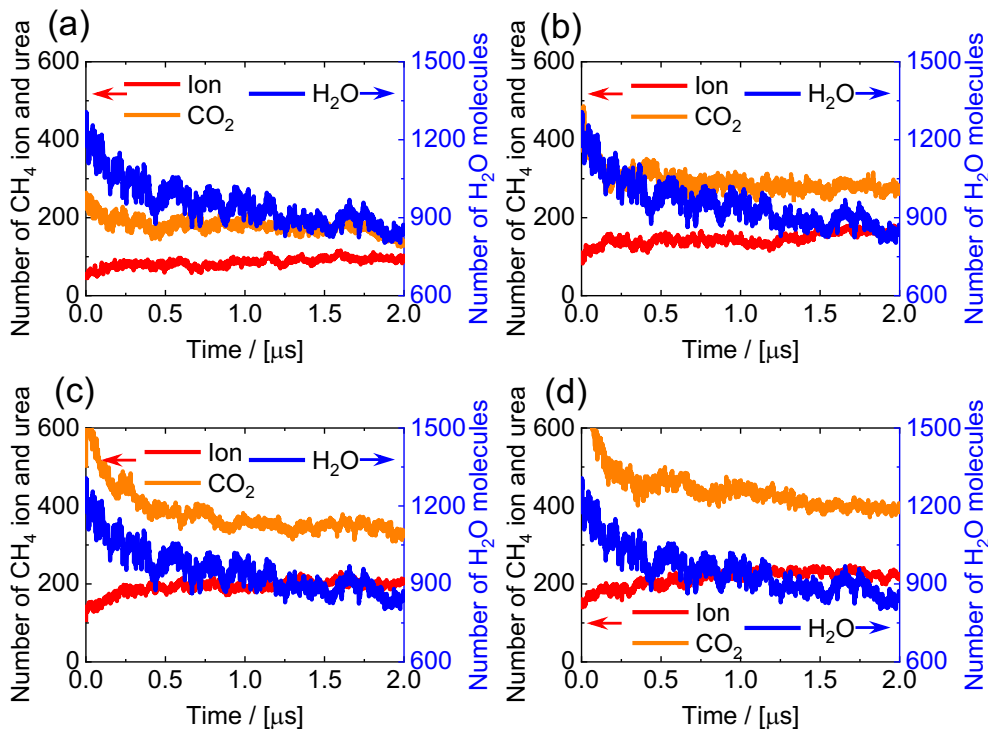


Figure S9. Evolution of the number of ions, CO₂, and H₂O molecules near the urea molecules within 5 Å in the (a) H_{1.6%}Urea, (b) H_{3%}Urea, (c) H_{4.5%}Urea, and (d) H_{6%}Urea systems.

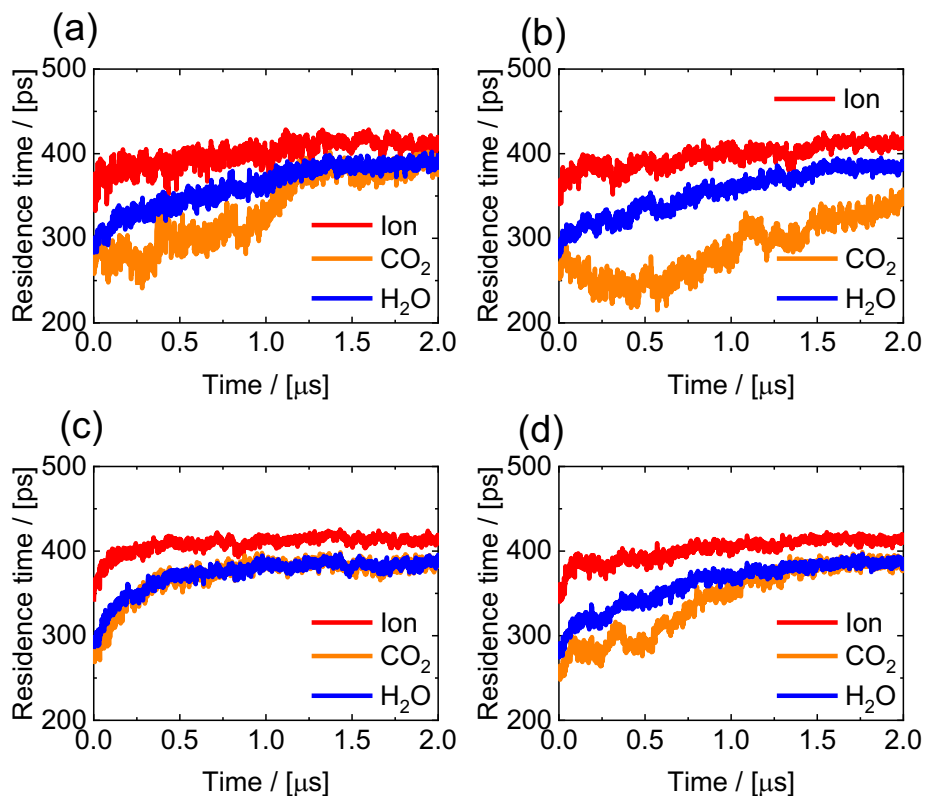


Figure S10. Evolution of the average residence time (τ_{Res}) for H₂O, ions and CO₂ molecules near the urea molecules in the (a) H_{1.6%}Urea, (b) H_{3%}Urea, (c) H_{4.5%}Urea, and (d) H_{6%}Urea systems.

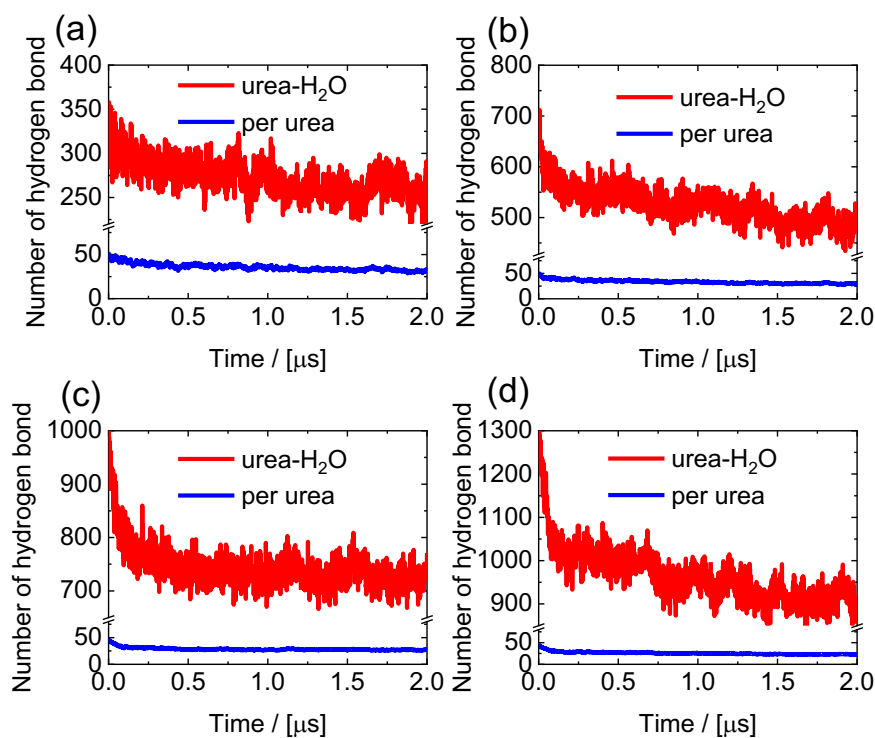


Figure S11. Evolution of the number of hydrogen bonds between urea and H₂O molecules and total hydrogen bonds near per urea molecule within 5 Å in the (a) H_{1.6%}Urea, (b) H_{3%}Urea, (c) H_{4.5%}Urea, and (d) H_{6%}Urea systems.

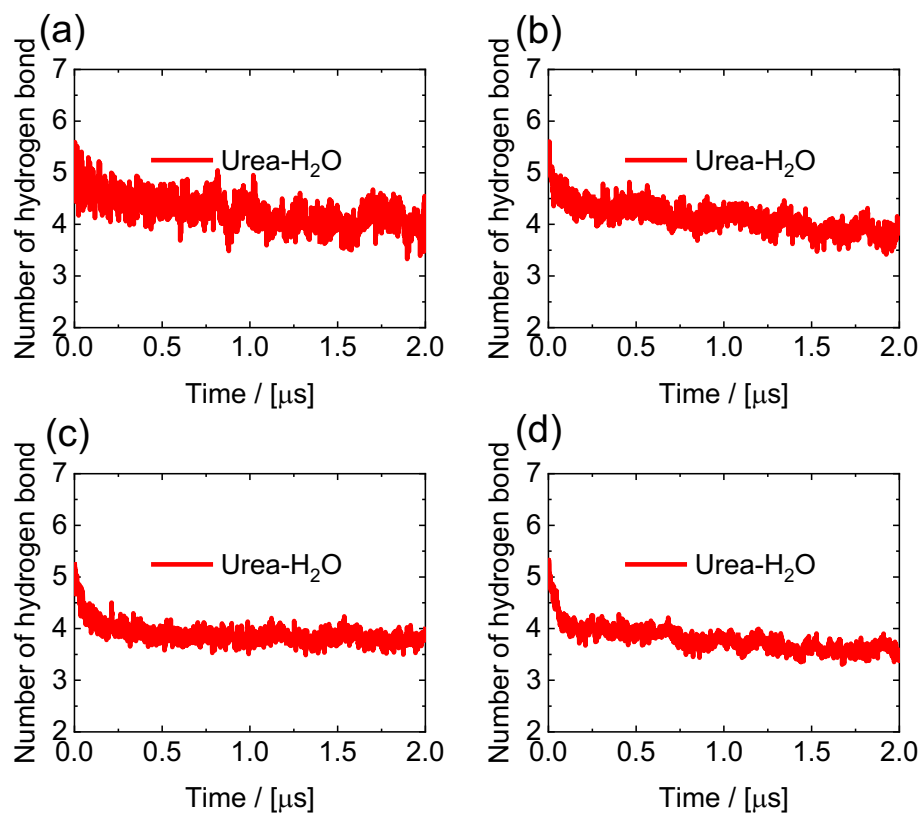


Figure S12. Evolution of the number of hydrogen bonds between per urea molecule and H₂O molecules in the (a) H_{1.6%}Urea, (b) H_{3%}Urea, (c) H_{4.5%}Urea, and (d) H_{6%}Urea systems.

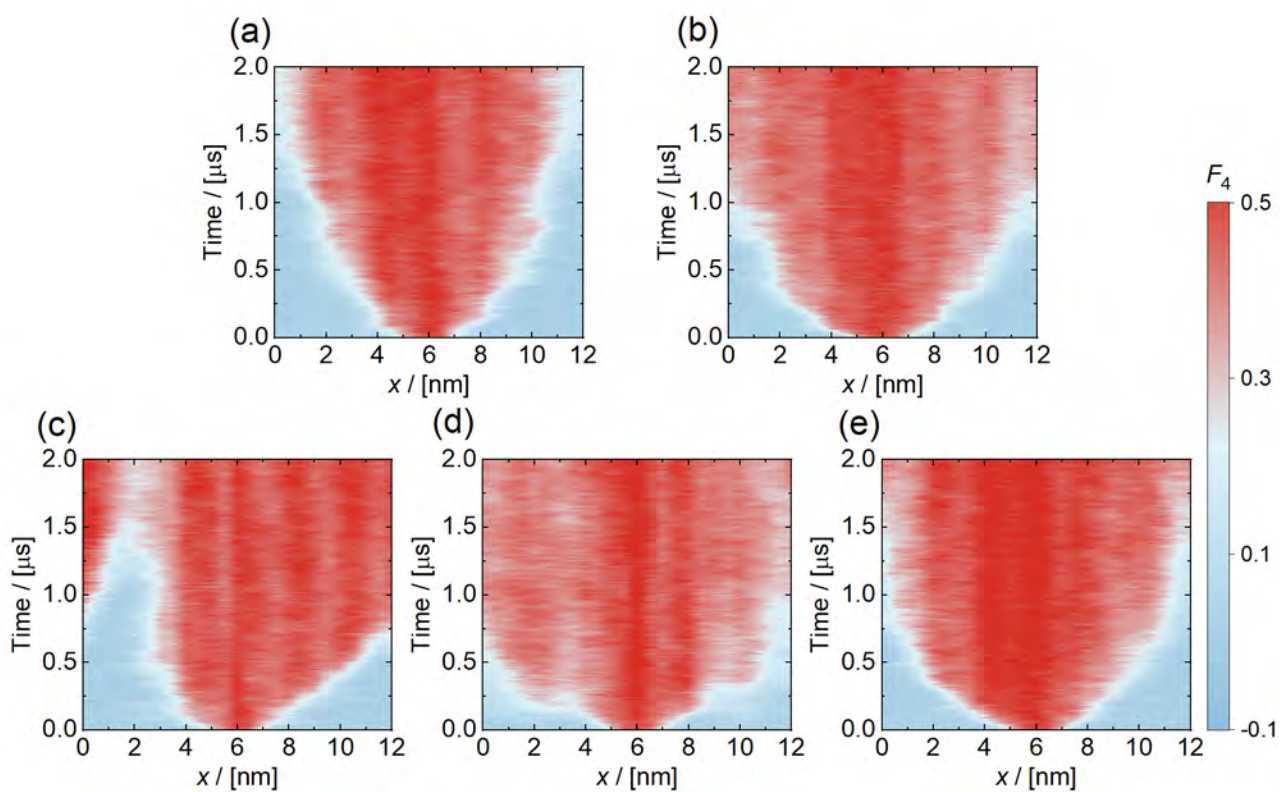


Figure S13. Evolution of the F_4 order parameter along the x-axis in the (a) $H_0\%Urea$, (b) $H_{1.6}\%Urea$, (c) $H_3\%Urea$, (d) $H_{4.5}\%Urea$, and (e) $H_6\%Urea$ systems.

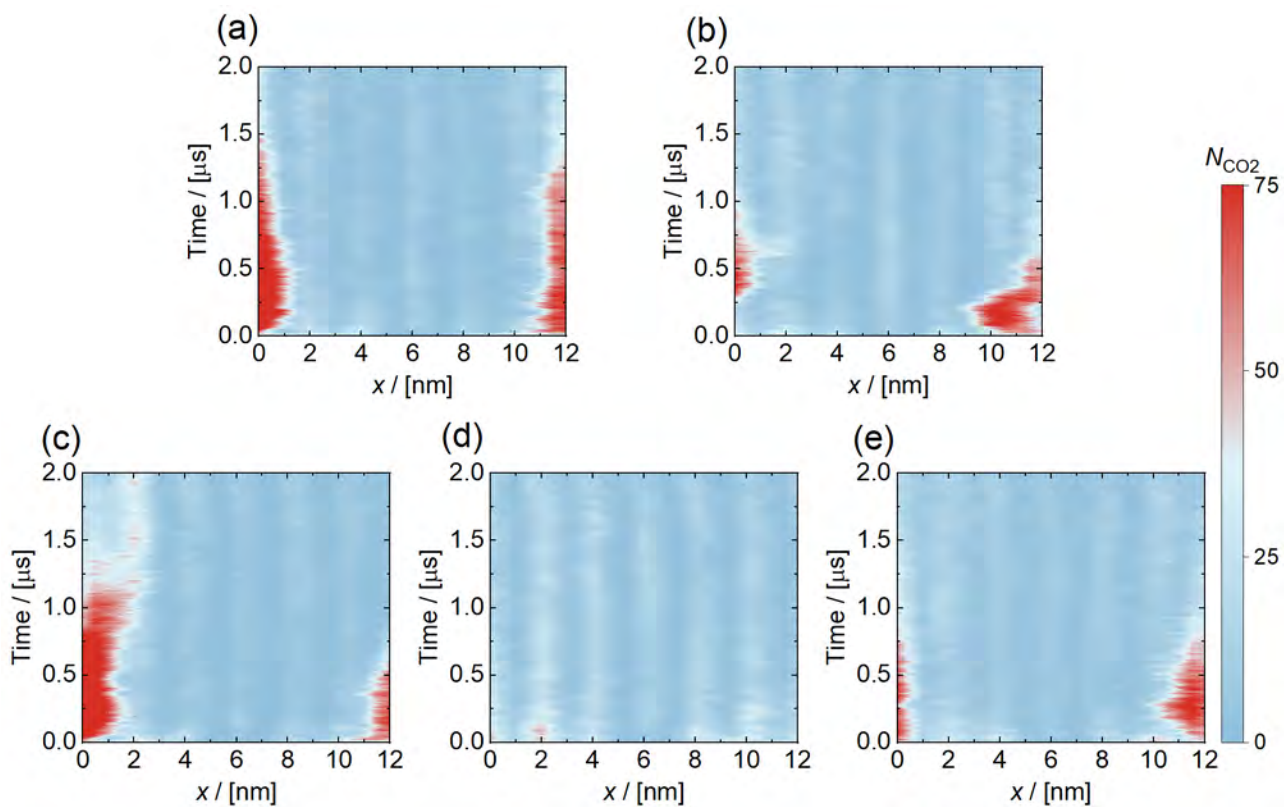


Figure S14. Evolution of the number of CO₂ molecules in the nanobubbles (N_{CO_2}) along the x-axis in the (a) H_{0%}Urea, (b) H_{1.6%}Urea, (c) H_{3%}Urea, (d) H_{4.5%}Urea, and (e) H_{6%}Urea systems.

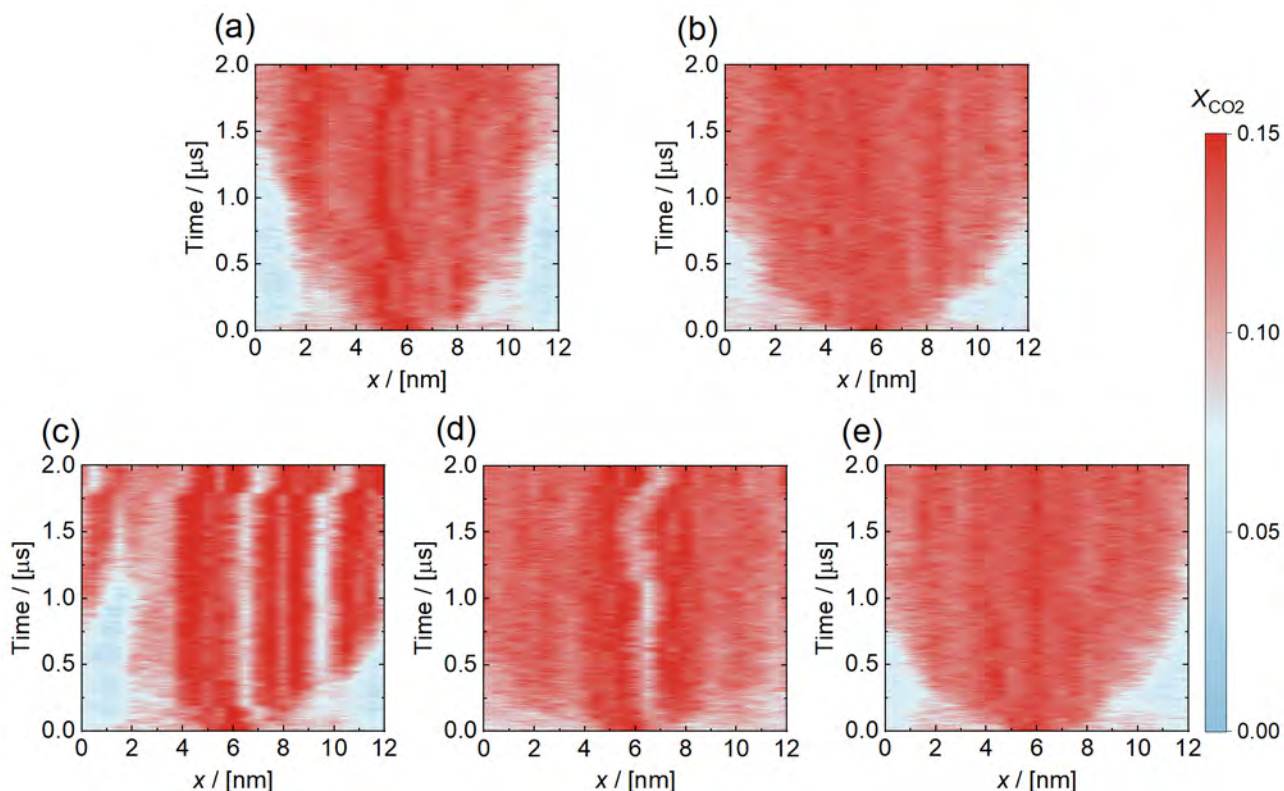


Figure S15. Evolution of the CO₂ mole fraction in water (x_{CO_2}) for solutions along the x-axis in the (a) H_{0%}Urea, (b) H_{1.6%}Urea, (c) H_{3%}Urea, (d) H_{4.5%}Urea, and (e) H_{6%}Urea systems.

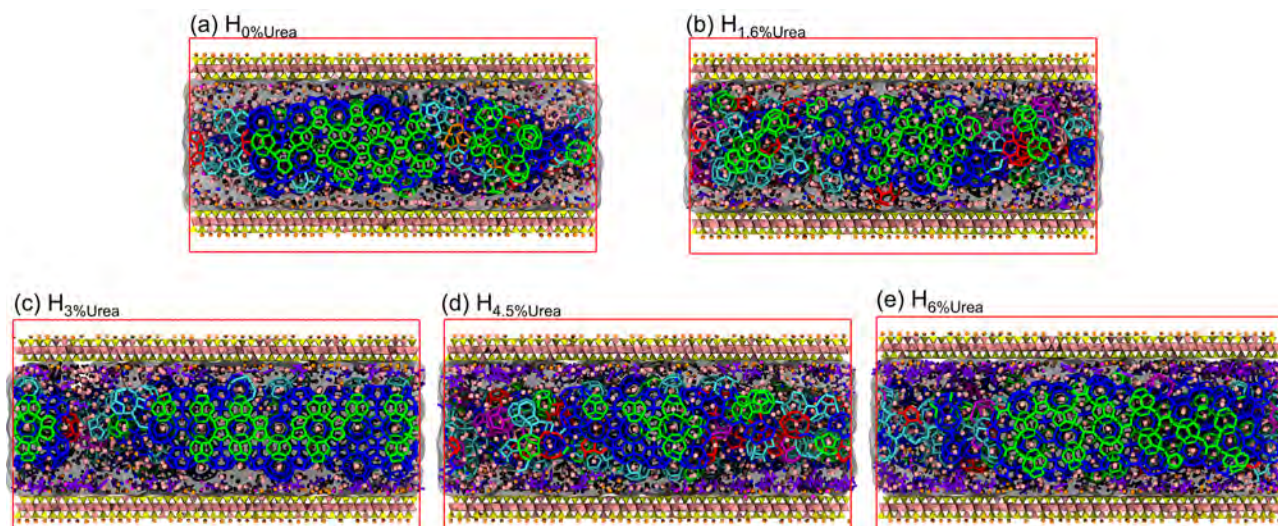


Figure S16. Growth snapshots of CO₂ hydrates formed at the end of the simulation (at 2.0 μs) for the z-x plane in the different simulations: (a) H_{0%}Urea, (b) H_{1.6%}Urea, (c) H_{3%}Urea, (d) H_{4.5%}Urea, and (e) H_{6%}Urea systems. Illite is displayed as polyhedral, *i.e.*, yellow (Si atom), and pink (Al atom). Pink, blue, magenta, orange, and violet represent CO₂, Cl⁻, Na⁺, K⁺, and urea, respectively. Hydrate cages are shown as sticks in various colors (green for 5¹², blue for 5¹²6², red for 5¹²6³, orange for 5¹²6⁴, cyan for 4¹5¹⁰6², purple for 4¹5¹⁰6³ and pink for 4¹5¹⁰6⁴).

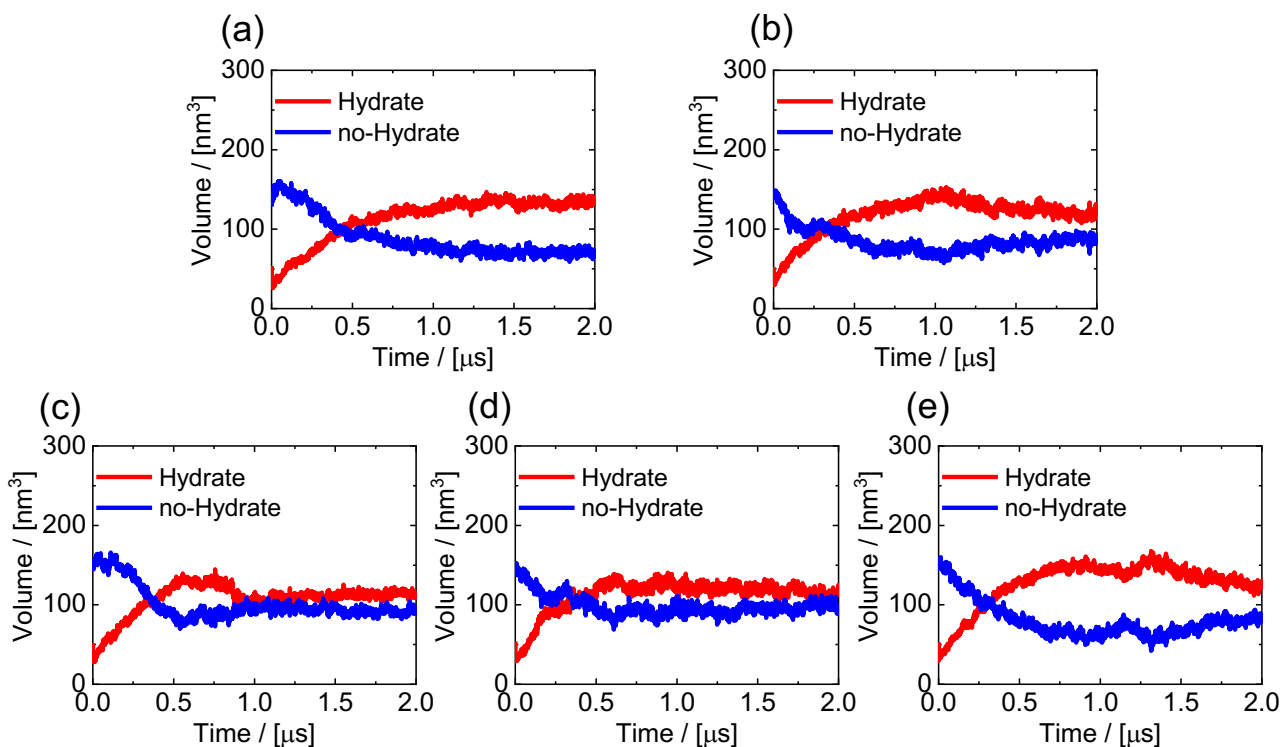


Figure S17. Evolution of the volume of the CO₂ hydrate solid in the (a) H_{0%}Urea, (b) H_{1.6%}Urea, (c) H_{3%}Urea, (d) H_{4.5%}Urea, and (e) H_{6%}Urea systems.

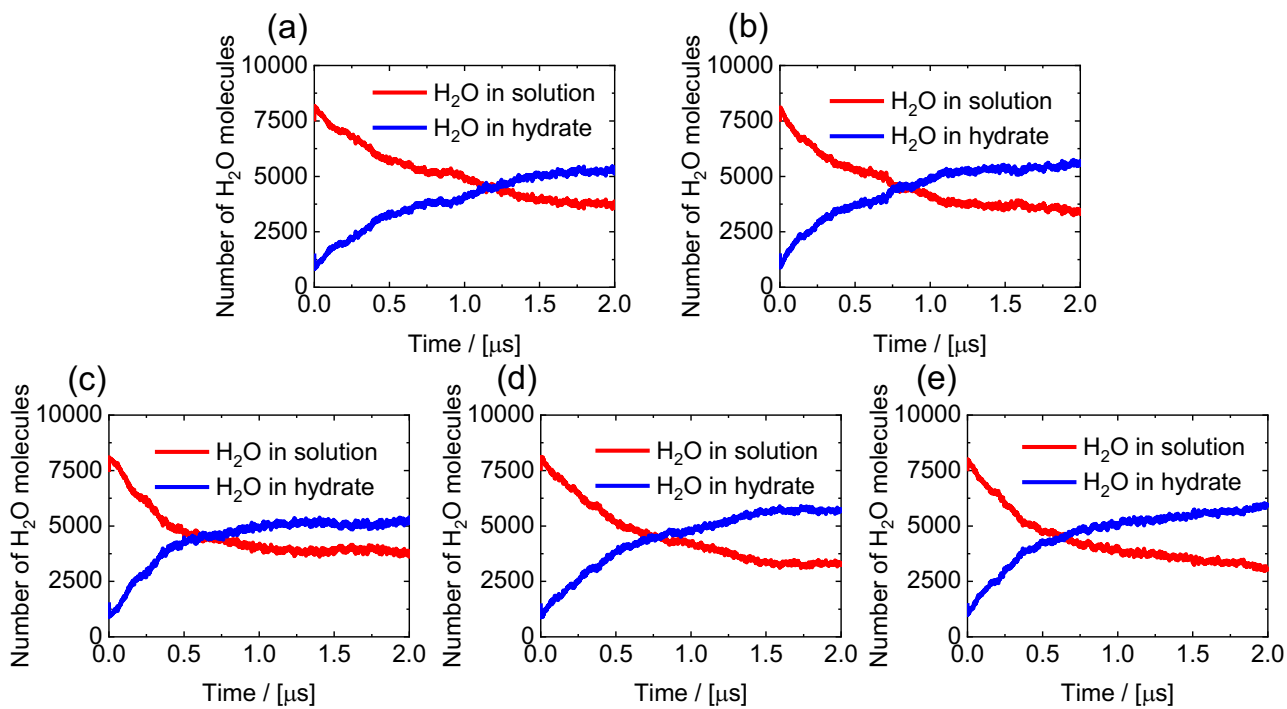


Figure S18. Evolution of the number of H₂O molecules in hydrate and solution in the (a) H_{0%}Urea, (b) H_{1.6%}Urea, (c)

H_{3%}Urea, (d) H_{4.5%}Urea, and (e) H_{6%}Urea systems.

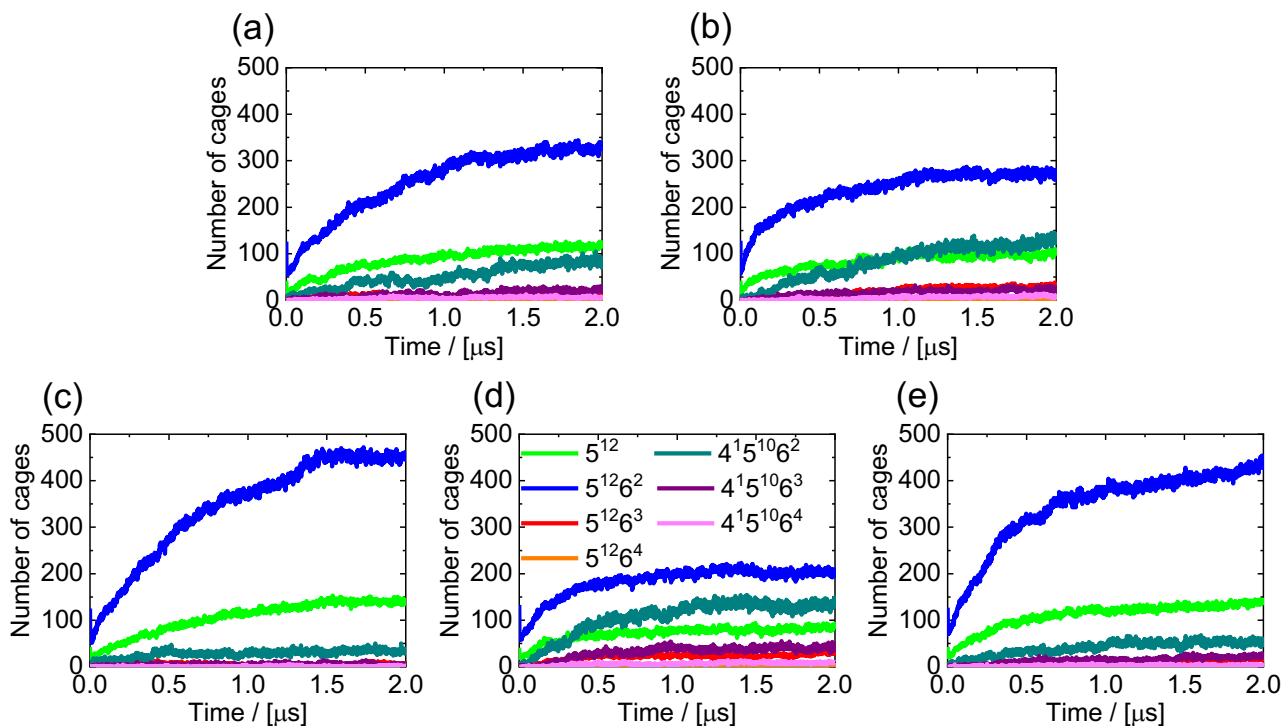


Figure S19. Evolution of the number of CO₂ hydrate cages in the (a) H_{0%}Urea, (b) H_{1.6%}Urea, (c) H_{3%}Urea, (d) H_{4.5%}Urea, and (e) H_{6%}Urea systems.

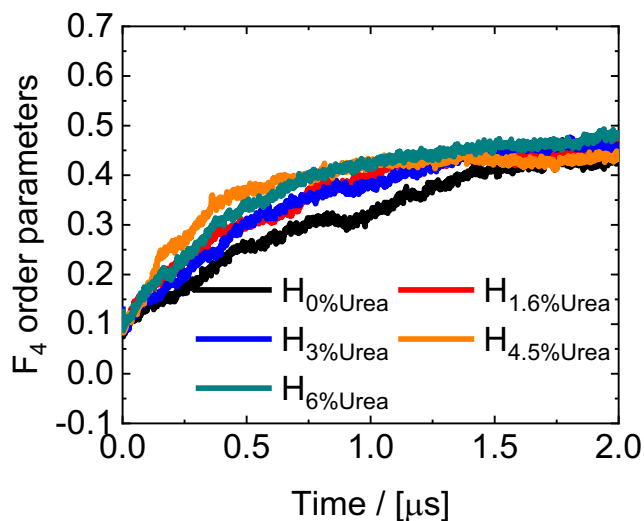


Figure S20. The evolution of the F_4 order parameter for the different five systems, *i.e.*, H_{0%}Urea, H_{1.6%}Urea, H_{3%}Urea, H_{4.5%}Urea, and H_{6%}Urea systems.

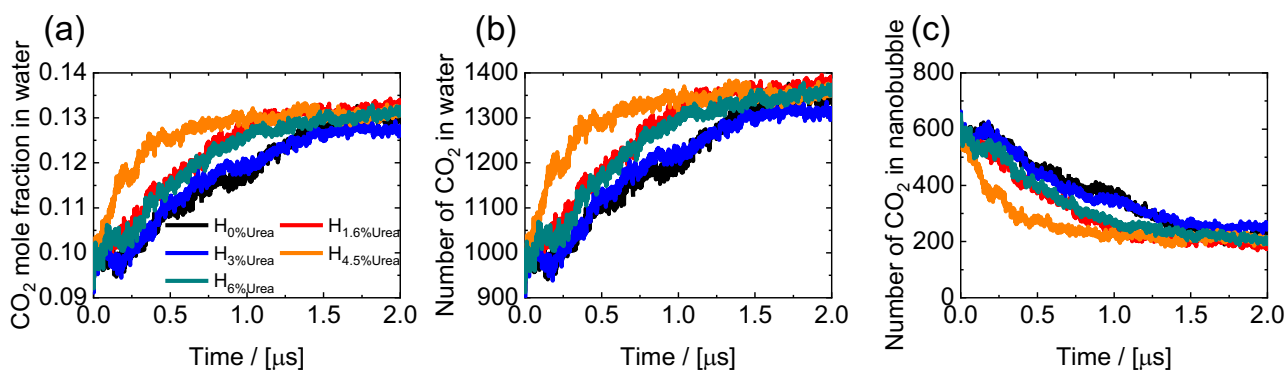


Figure S21. Evolution of (a) CO₂ mole fraction in water (x_{CO_2}), number of CO₂ molecules (b) in water, and (c) in nanobubble for the different five systems, *i.e.*, H_{0%}Urea, H_{1.6%}Urea, H_{3%}Urea, H_{4.5%}Urea, and H_{6%}Urea systems.

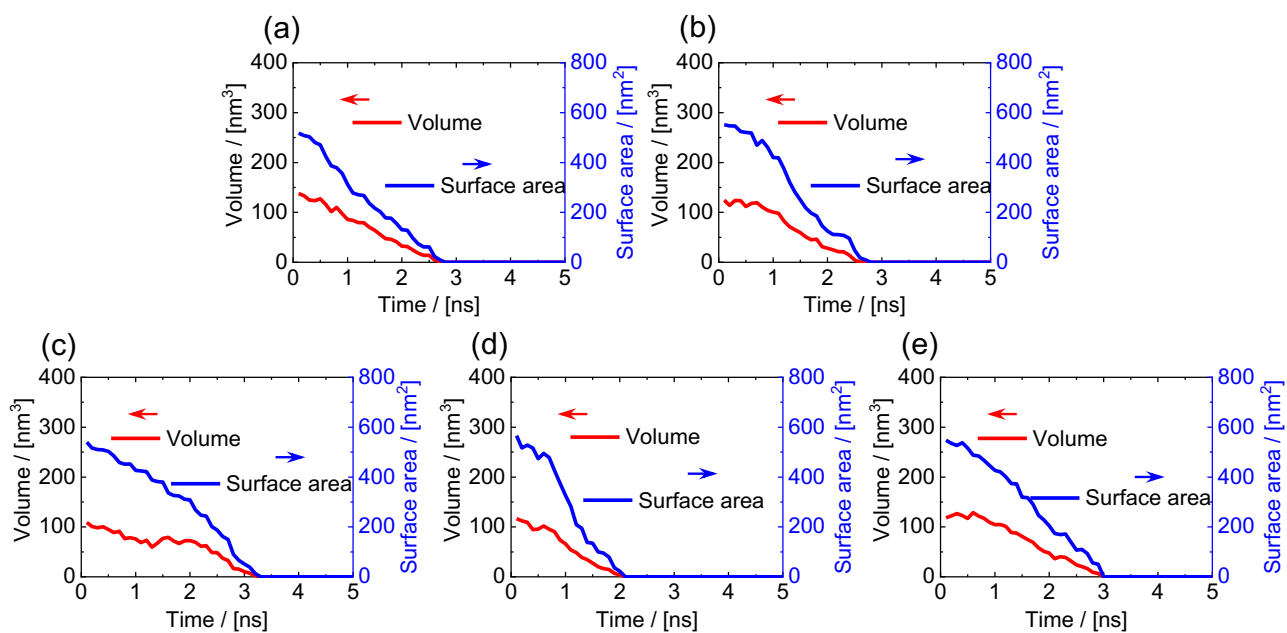


Figure S22. Evolution of the volume/surface area of CO₂ hydrate solid in the (a) H_{0%}Urea, (b) H_{1.6%}Urea, (c) H_{3%}Urea, (d) H_{4.5%}Urea, and (e) H_{6%}Urea systems.

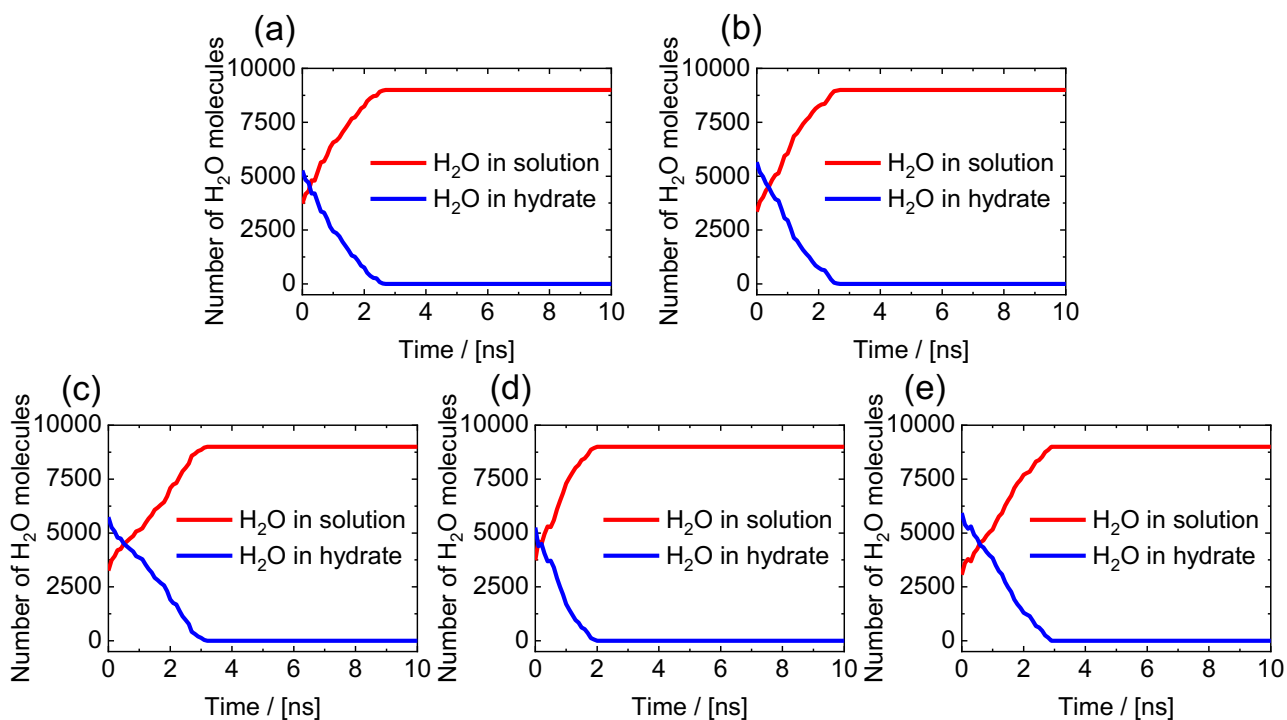


Figure S23. Evolution of the number of H₂O molecules in hydrate and solution in the (a) H_{0%}Urea, (b) H_{1.6%}Urea, (c) H_{3%}Urea, (d) H_{4.5%}Urea, and (e) H_{6%}Urea systems.

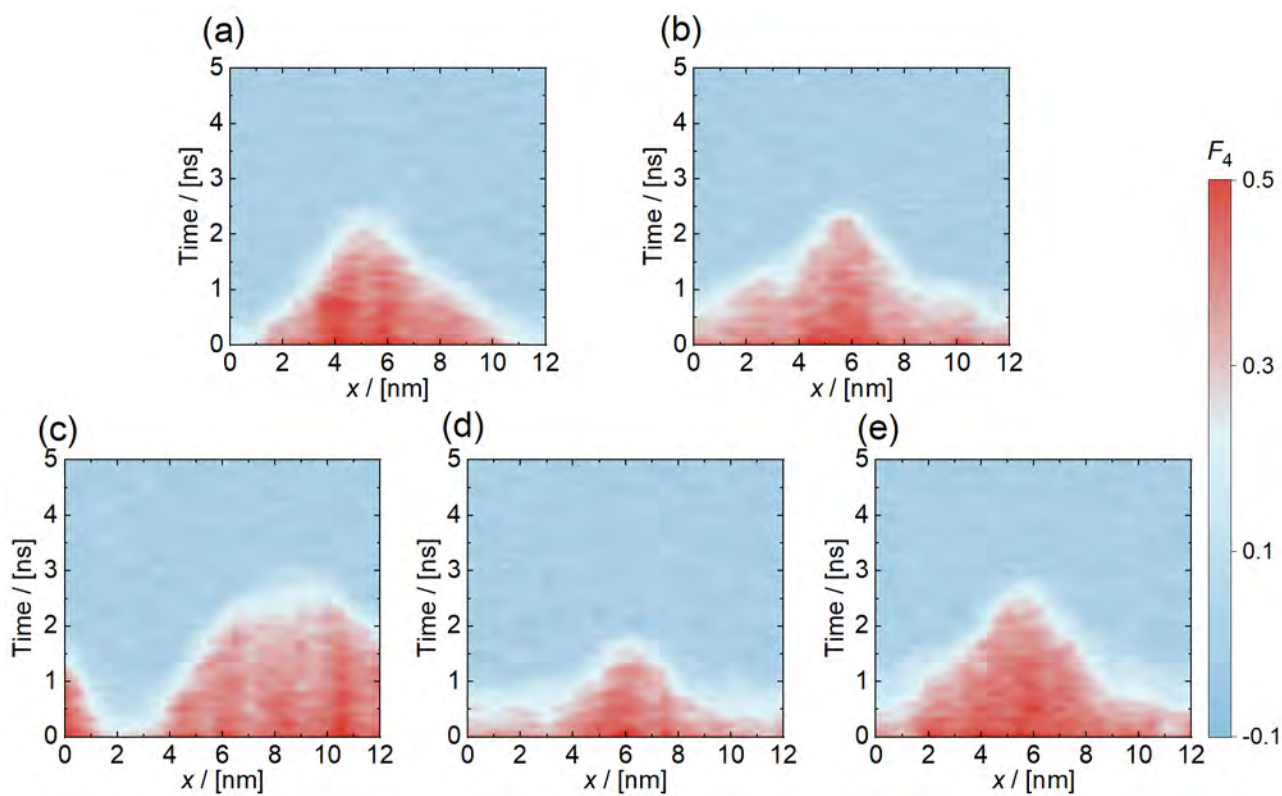


Figure S24. Evolution of the F_4 order parameter along the x-axis in the (a) H_{0%}Urea, (b) H_{1.6%}Urea, (c) H_{3%}Urea, (d)

H_{4.5%}Urea, and (e) H_{6%}Urea systems.

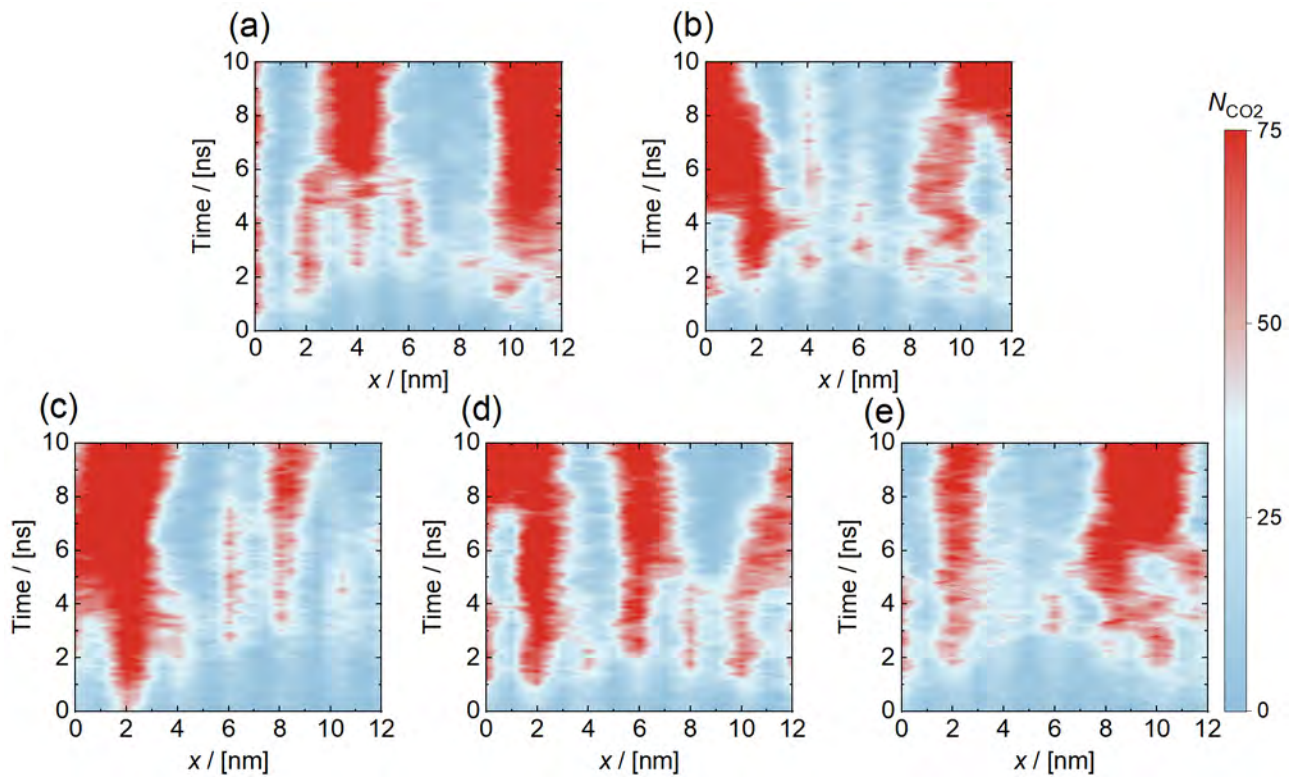


Figure S25. Evolution of the number of CO₂ in the nanobubbles (N_{CO_2}) along the x-axis in the (a) H_{0%}Urea, (b) H_{1.6%}Urea, (c) H_{3%}Urea, (d) H_{4.5%}Urea, and (e) H_{6%}Urea systems.

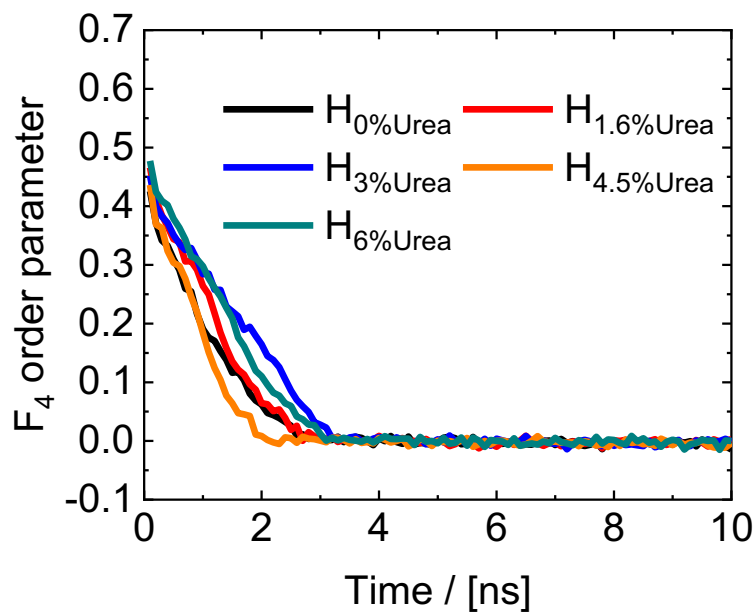


Figure S26. Evolution of the F_4 order parameter for the different five systems, *i.e.*, H_{0%}Urea, H_{1.6%}Urea, H_{3%}Urea,

H_{4.5%}Urea, and H_{6%}Urea systems.

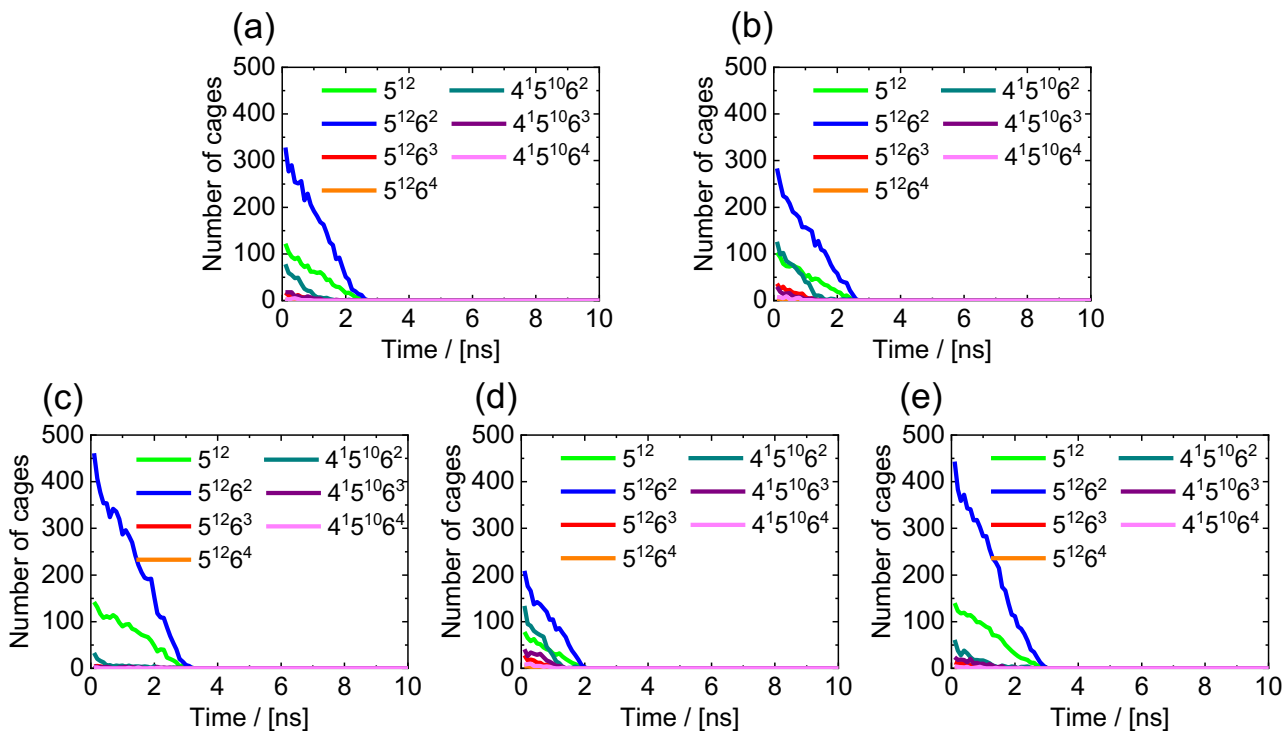


Figure S27. Evolution of the number of CO₂ hydrate cages in the (a) H_{0%}Urea, (b) H_{1.6%}Urea, (c) H_{3%}Urea, (d) H_{4.5%}Urea, and (e) H_{6%}Urea systems.

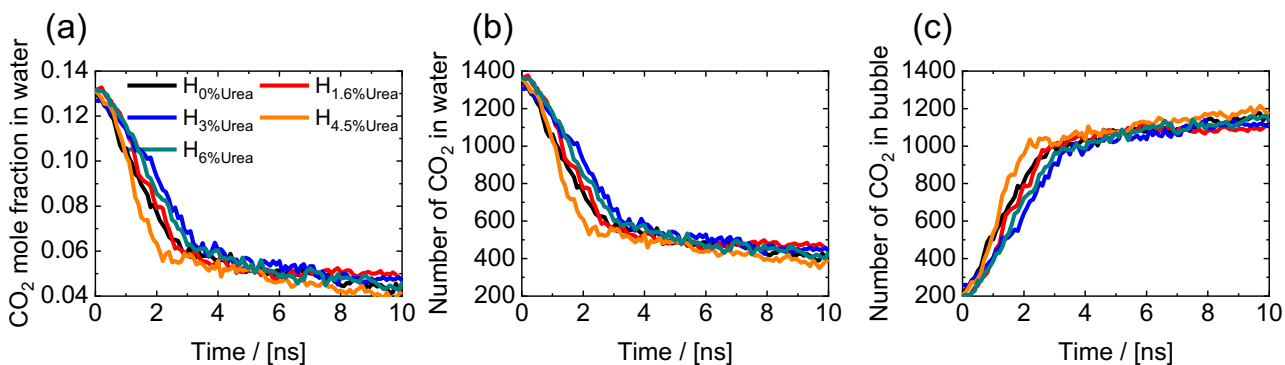


Figure S28. Evolution of (a) CO₂ mole fraction in water (x_{CO_2}), number of CO₂ molecules (b) in water, and (c) in nanobubble for the different five systems, *i.e.*, H_{0%}Urea, H_{1.6%}Urea, H_{3%}Urea, H_{4.5%}Urea, and H_{6%}Urea systems.

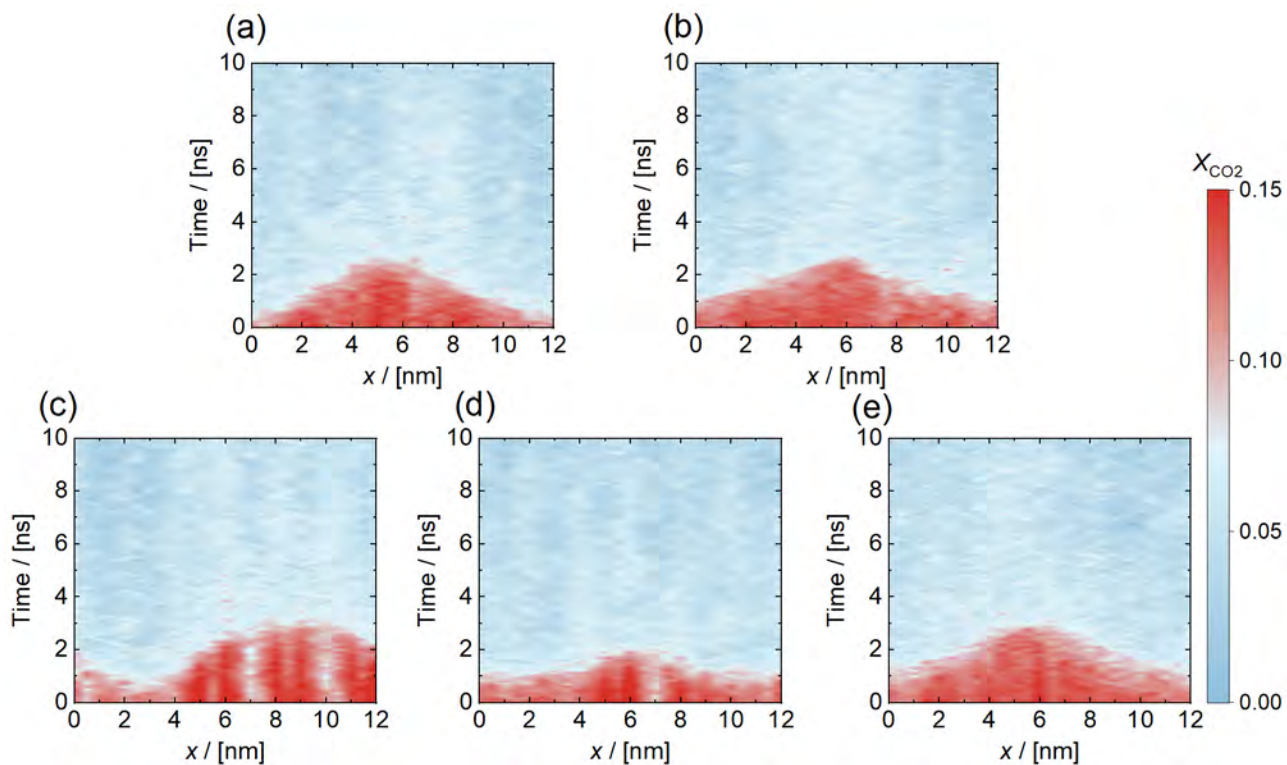


Figure S29. Evolution of the CO₂ mole fraction in water (x_{CO_2}) for solutions along the x-axis in the (a) H_{0%}Urea, (b) H_{1.6%}Urea, (c) H_{3%}Urea, (d) H_{4.5%}Urea, and (e) H_{6%}Urea systems.

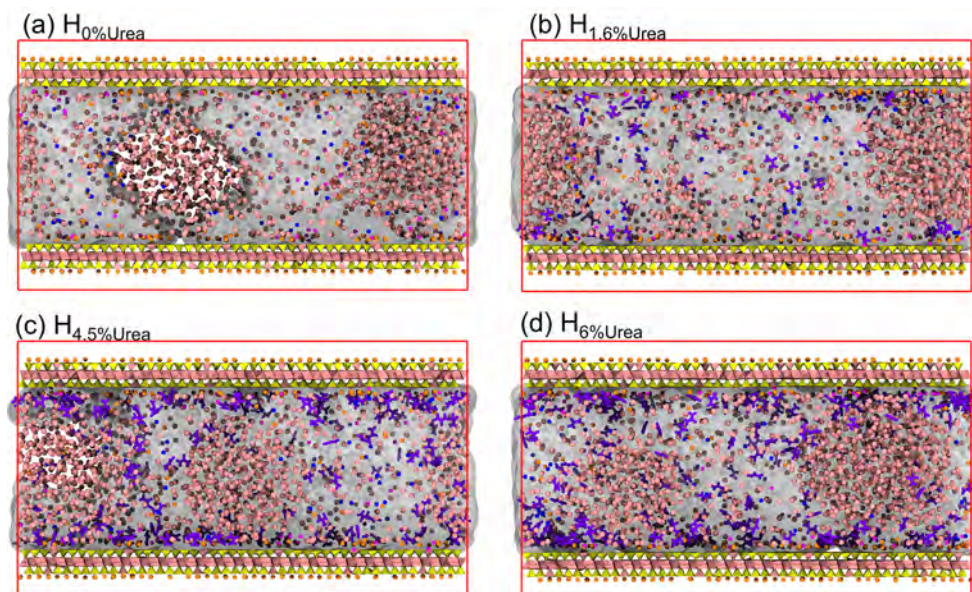


Figure S30. Dissociation snapshots at the end of the dissociation simulation (at the 10 ns) for the z-x plane in the different simulations: (a) H_{0%}Urea, (b) H_{1.6%}Urea, (c) H_{4.5%}Urea, and (d) H_{6%}Urea systems. Illite are displayed as polyhedral, *i.e.*, yellow (Si atom), and pink (Al atom). Pink, blue, magenta, orange, and violet represent CO₂, Cl⁻, Na⁺, K⁺, and urea, respectively.

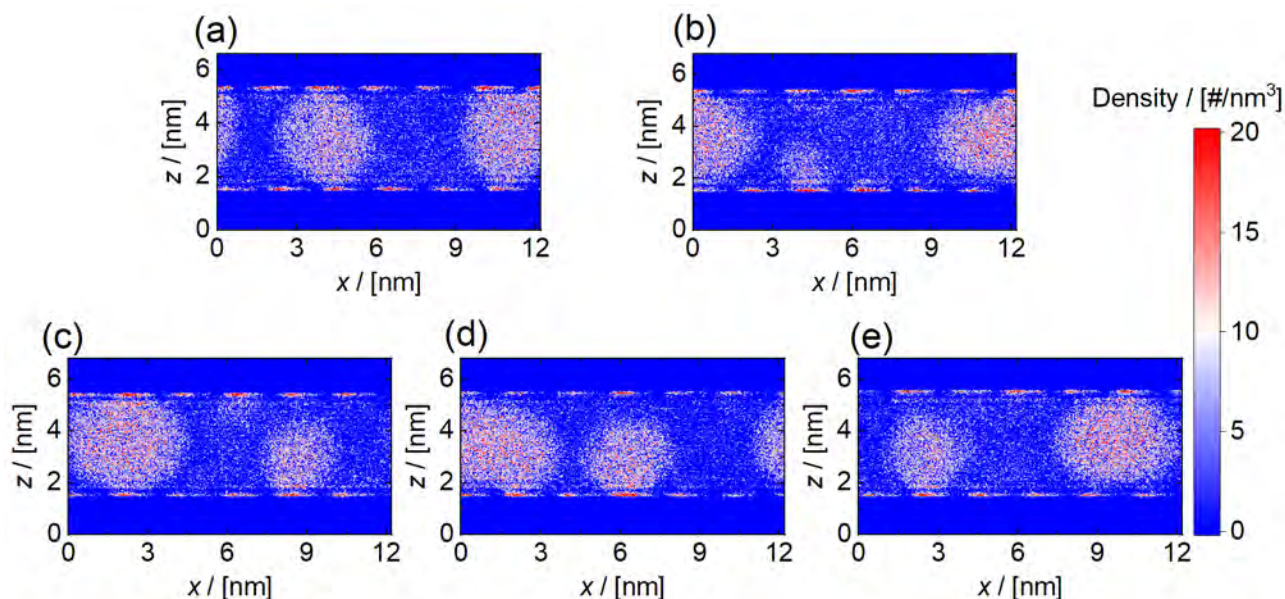


Figure S31. Number density distribution of CO₂ molecules for the last 2 ns of the dissociation process in the (a) H₀%Urea, (b) H_{1.6}%Urea, (c) H₃%Urea, (d) H_{4.5}%Urea, and (e) H₆%Urea systems.

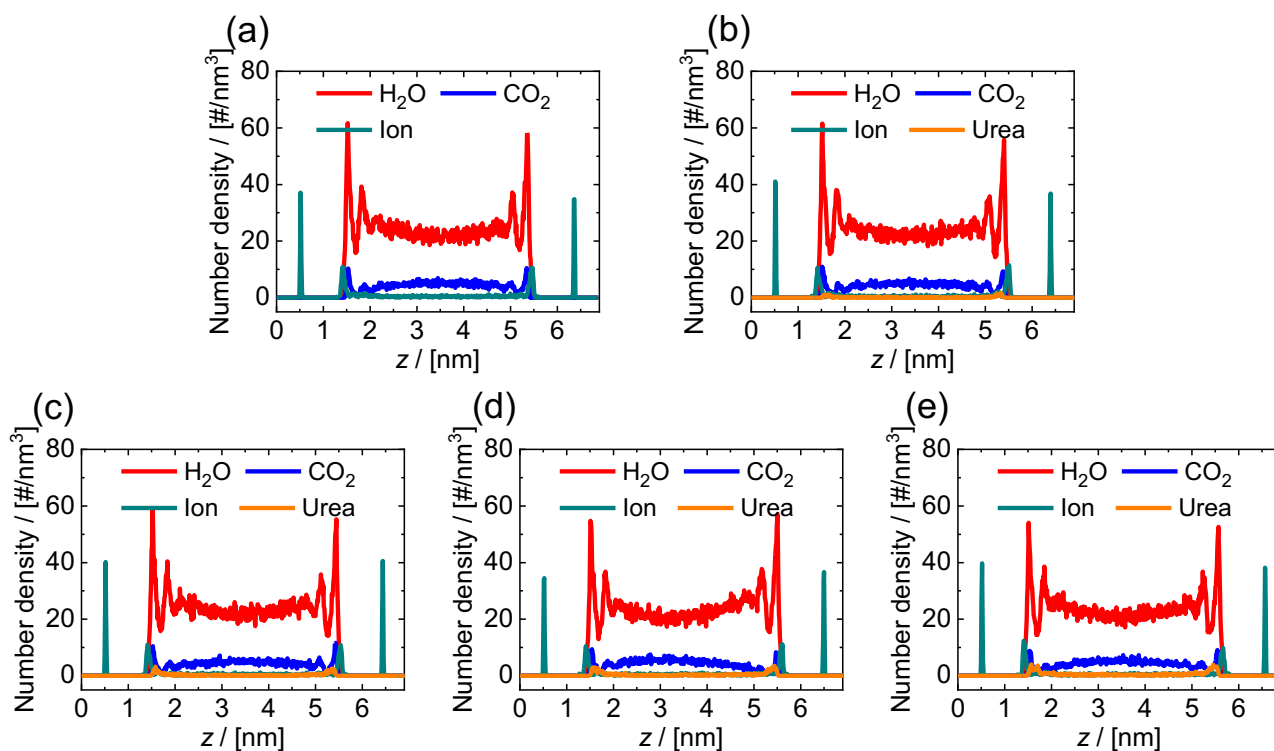


Figure S32. Number density distribution of H₂O, CO₂, ions, and urea molecules along the z-axis for 9 - 10 ns of the dissociation process for the (a) H₀%Urea, (b) H_{1.6}%Urea, (c) H₃%Urea, (d) H_{4.5}%Urea, and (e) H₆%Urea systems.

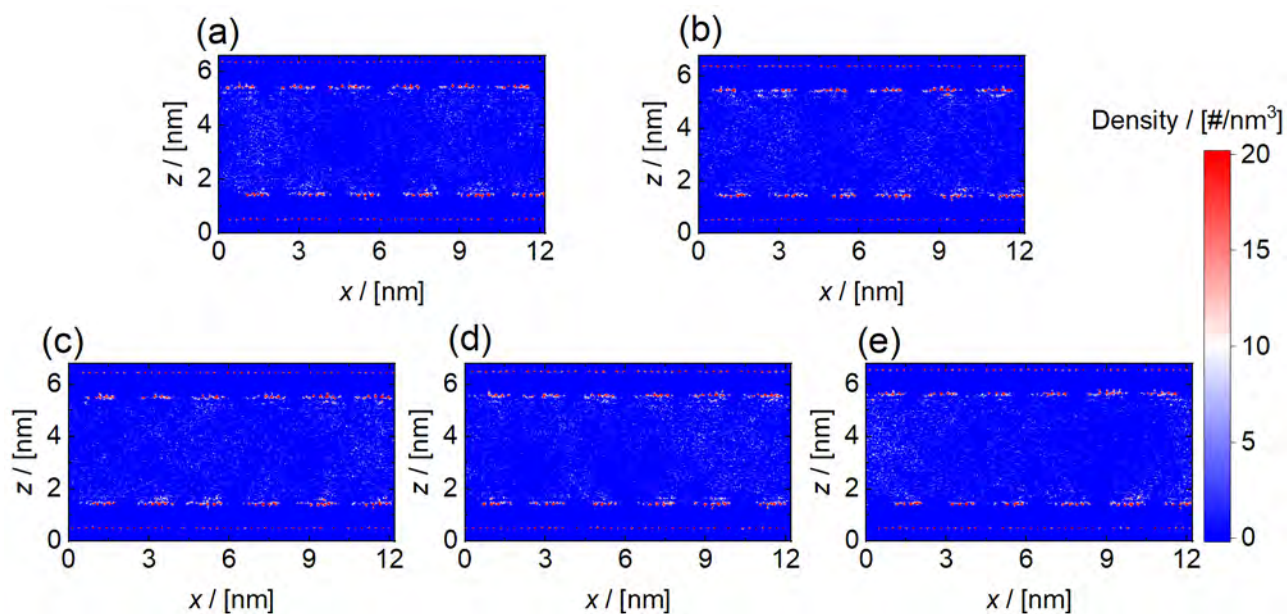


Figure S33. Number density distribution of ions for the last 2 ns of the dissociation process in the (a) $\text{H}_0\% \text{Urea}$, (b) $\text{H}_{1.6}\% \text{Urea}$, (c) $\text{H}_3\% \text{Urea}$, (d) $\text{H}_{4.5}\% \text{Urea}$, and (e) $\text{H}_6\% \text{Urea}$ systems.

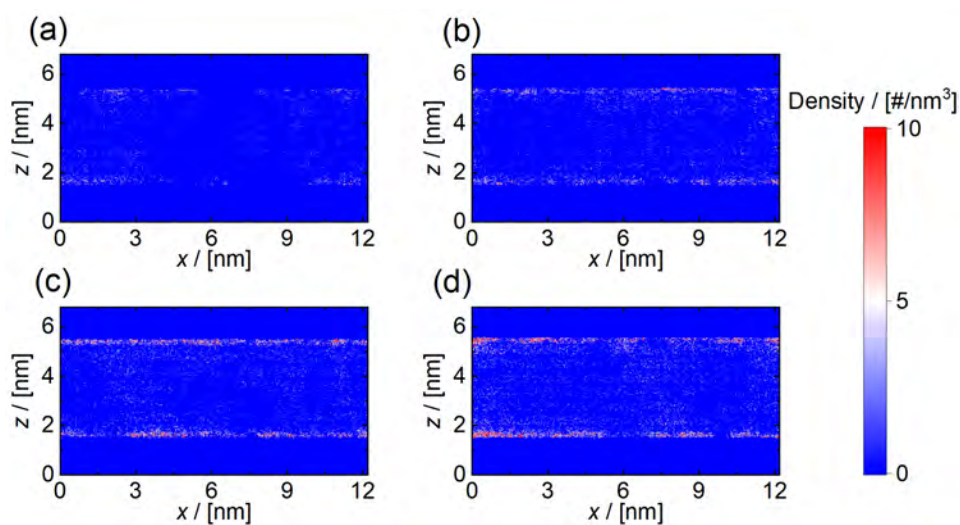


Figure S34. Number density distribution of urea molecules for the last 2 ns of the dissociation process in the (a) $\text{H}_{1.6}\% \text{Urea}$, (b) $\text{H}_3\% \text{Urea}$, (c) $\text{H}_{4.5}\% \text{Urea}$, and (d) $\text{H}_6\% \text{Urea}$ systems.

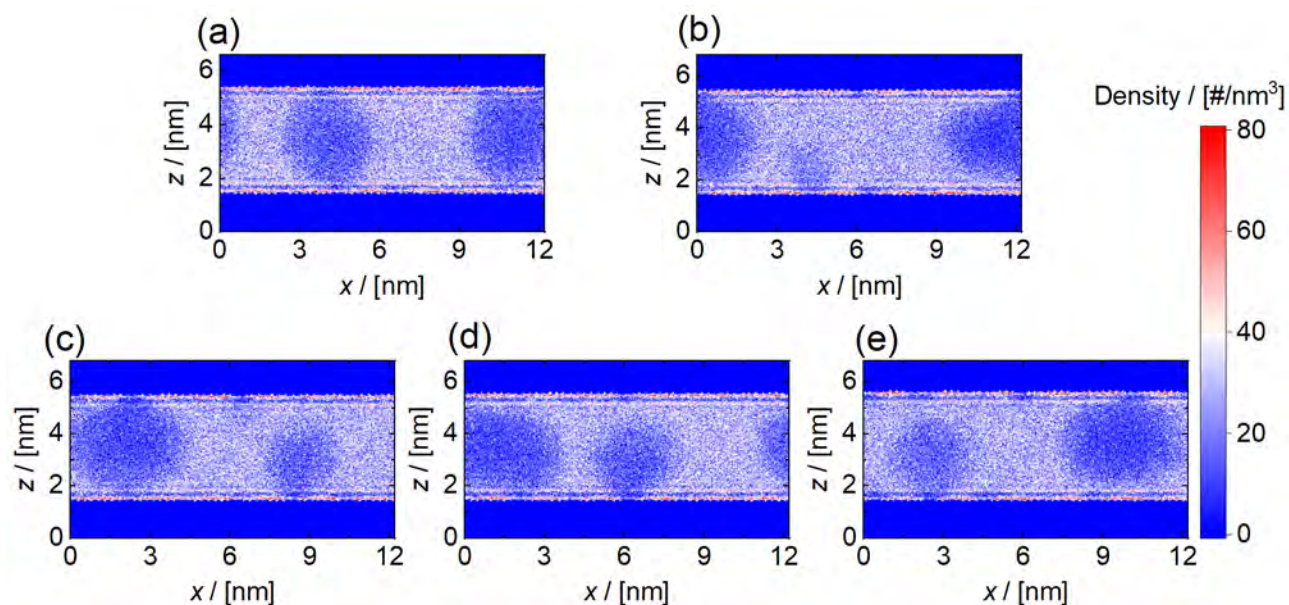


Figure S35. Number density distribution of H₂O molecules for the last 2 ns of the dissociation process in the (a) H₀%Urea, (b) H_{1.6}%Urea, (c) H₃%Urea, (d) H_{4.5}%Urea, and (e) H₆%Urea systems.

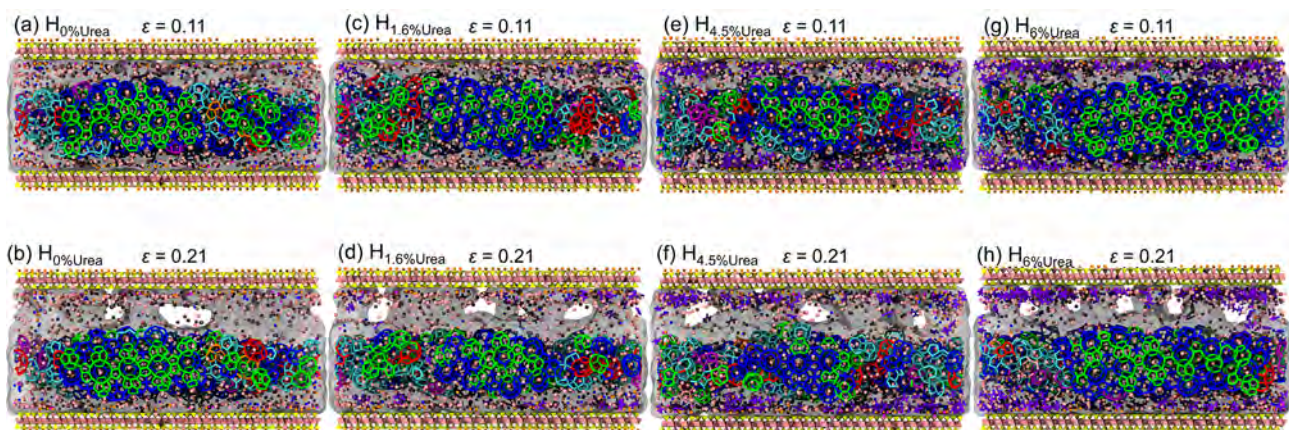


Figure S36. Snapshots of CO₂ hydrate-illite interface under tensile strain (ϵ) in the (a-b) H₀%Urea, (c-d) H_{1.6}%Urea, (e-f) H_{4.5}%Urea, and (g-h) H₆%Urea systems. Illite is displayed as polyhedral, *i.e.*, yellow (Si atom), and pink (Al atom). Pink, blue, magenta, orange, and violet represent CO₂, Cl⁻, Na⁺, K⁺, and urea, respectively. Hydrate cages are shown as sticks in various colors (green for 5¹², blue for 5¹²6², red for 5¹²6³, orange for 5¹²6⁴, cyan for 4¹5¹⁰6², purple for 4¹5¹⁰6³ and pink for 4¹5¹⁰6⁴).

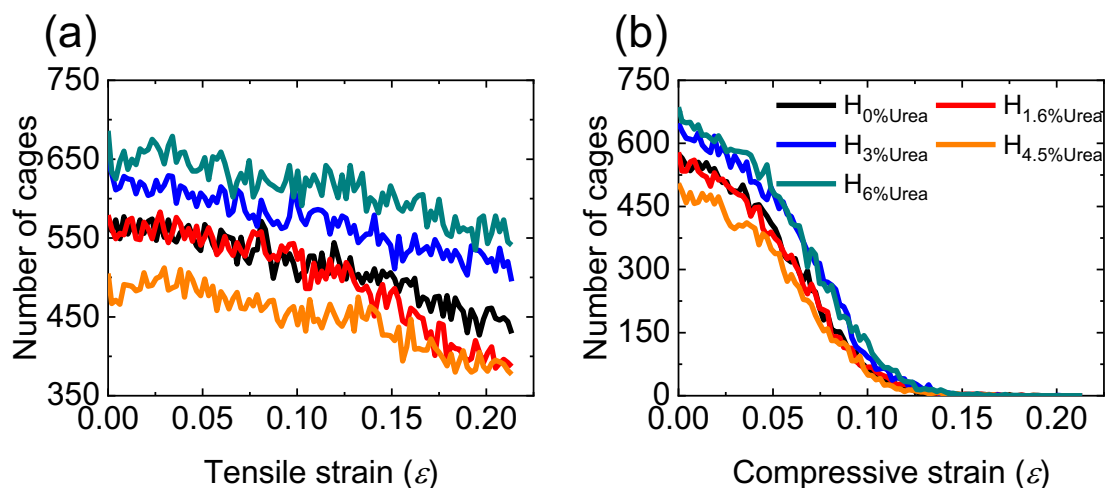


Figure S37. The evolution of the number of CO₂ hydrate cages under (a) tensile and (b) compressive strain (ϵ) for the different five systems, *i.e.*, H_{0%}Urea, H_{1.6%}Urea, H_{3%}Urea, H_{4.5%}Urea, and H_{6%}Urea systems.

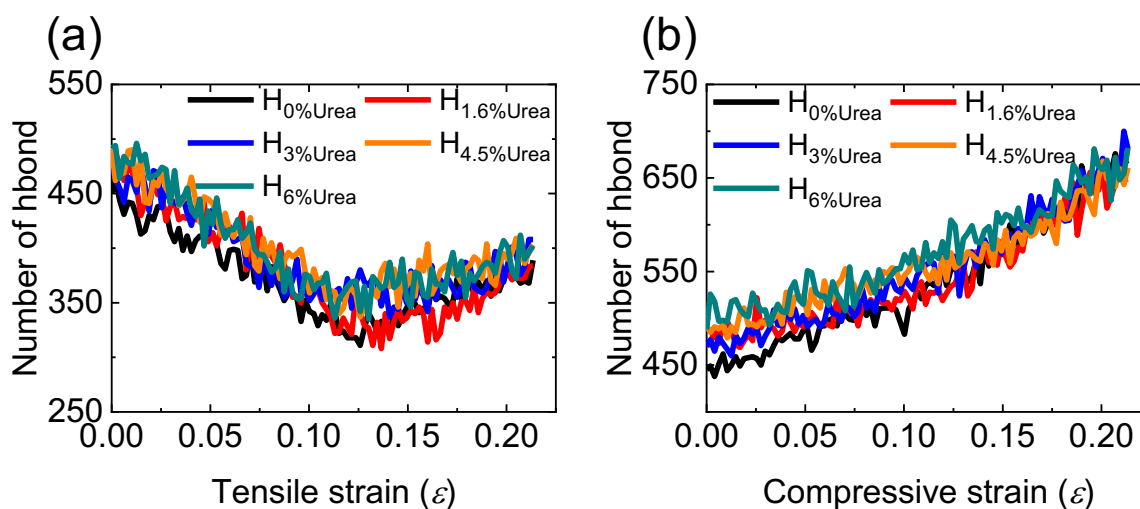


Figure S38. The evolution of the number of hydrogen bonds (hbonds) between urea/H₂O molecules and upper illite surface under (a) tensile and (b) compressive strain (ϵ) for the different five systems, *i.e.*, H_{0%}Urea, H_{1.6%}Urea, H_{3%}Urea, H_{4.5%}Urea, and H_{6%}Urea systems.

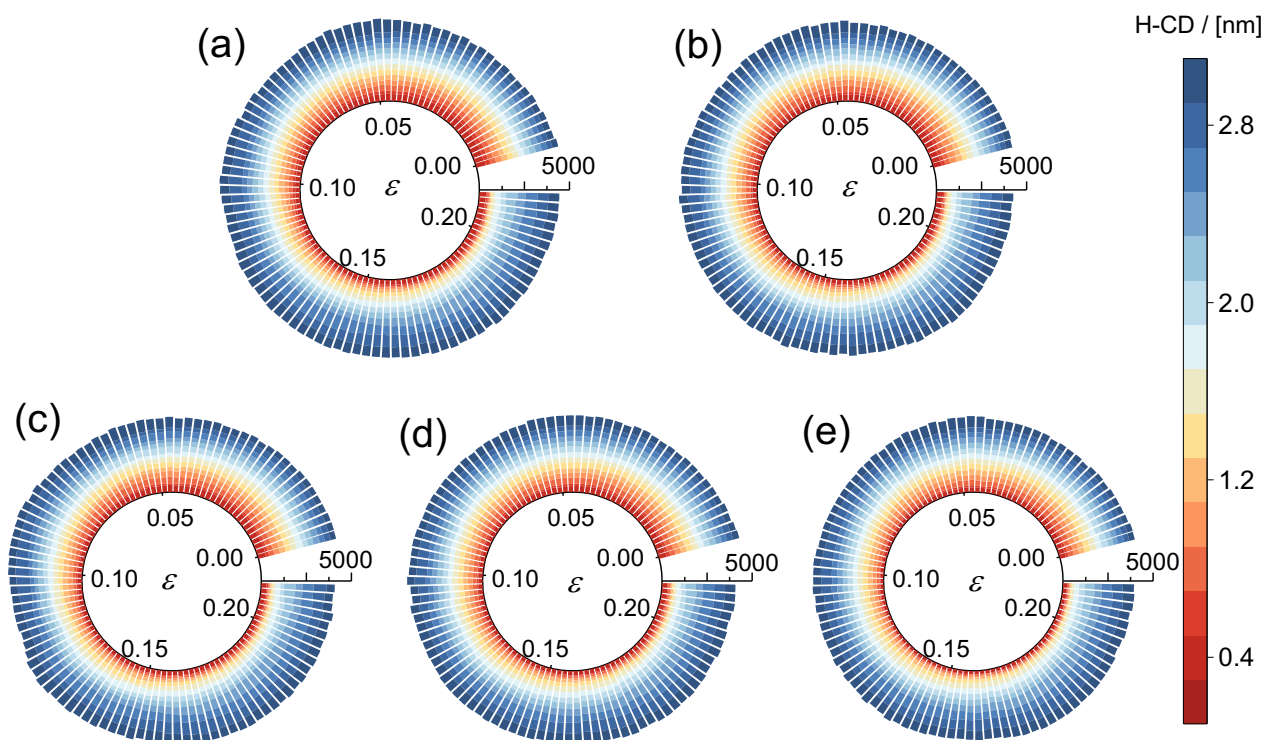


Figure S39. The number of hydrogen bonds at different distances from the upper illite surface (H-CD) under tensile strain (ϵ) in the (a) $H_{0\%Urea}$, (b) $H_{1.6\%Urea}$, (c) $H_{3\%Urea}$, (d) $H_{4.5\%Urea}$, and (e) $H_{6\%Urea}$ systems.

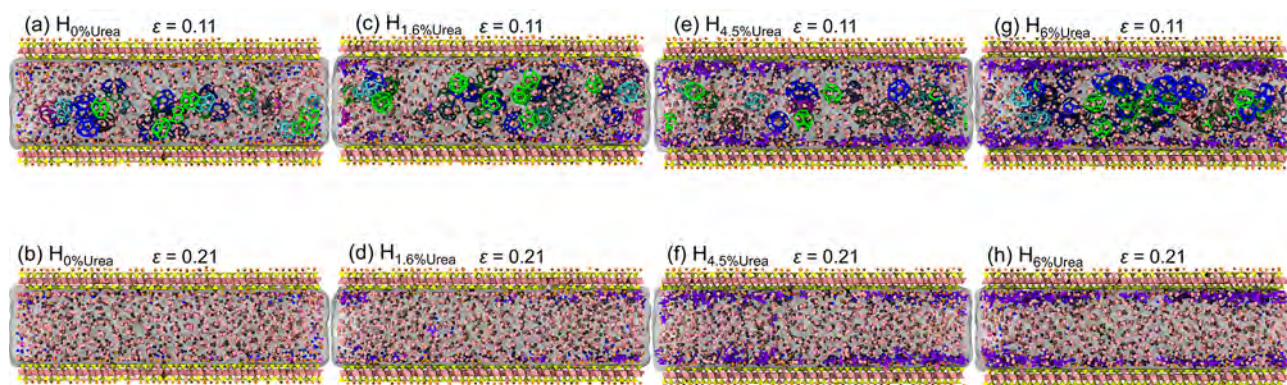


Figure S40. Snapshots of CO_2 hydrate-illite interface under compressive strain (ϵ) in the (a-b) $H_{0\%Urea}$, (c-d) $H_{1.6\%Urea}$, (e-f) $H_{4.5\%Urea}$, and (g-h) $H_{6\%Urea}$ systems. Illite are displayed as polyhedral, *i.e.*, yellow (Si atom), and pink (Al atom). Pink, blue, magenta, orange, and violet represent CO_2 , Cl^- , Na^+ , K^+ , and urea, respectively. Hydrate cages are shown as sticks in various colors (green for 5^{12} , blue for $5^{12}6^2$, red for $5^{12}6^3$, orange for $5^{12}6^4$, cyan for $4^{15}10^2$, purple for $4^{15}10^3$ and pink for $4^{15}10^4$).

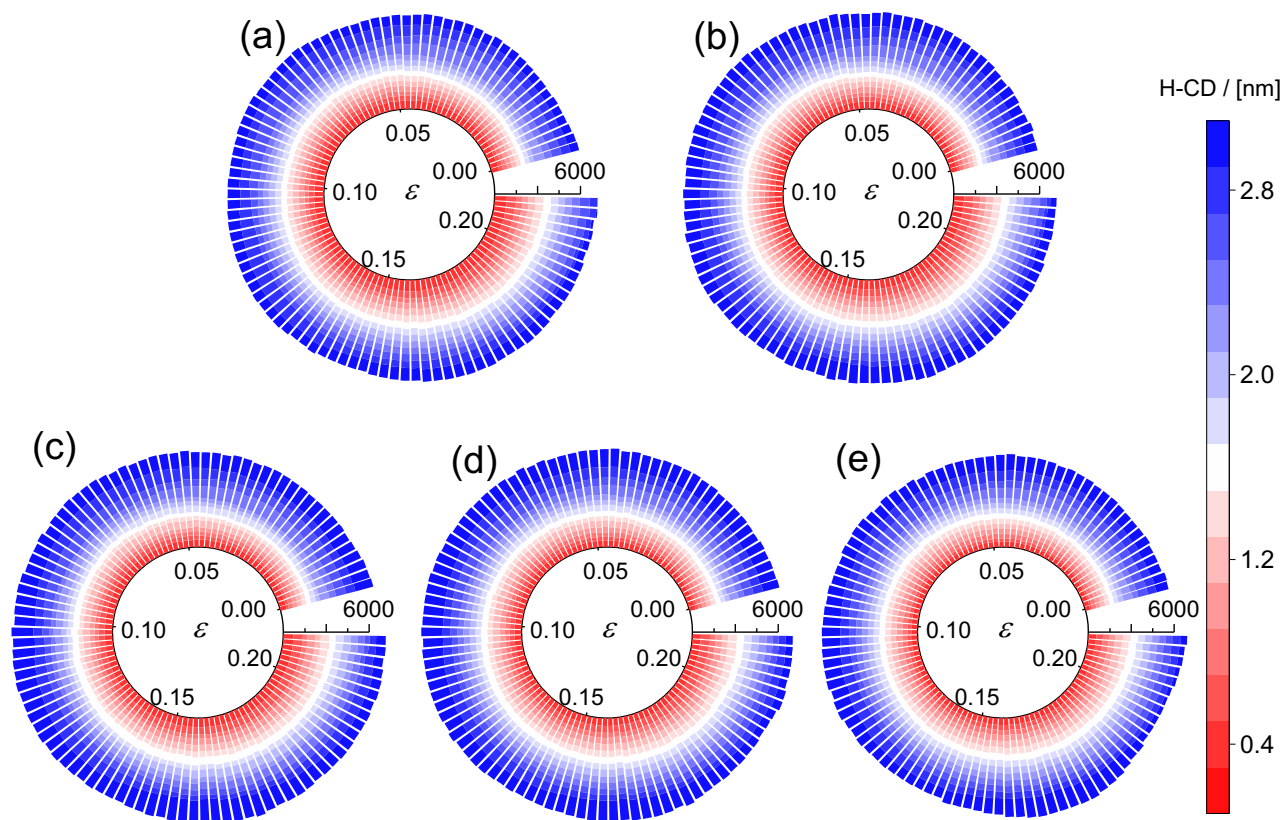


Figure S41. Number of hydrogen bonds at different distances from the upper illite surface under compressive strains (ϵ) in the (a) $H_0\%Urea$, (b) $H_{1.6}\%Urea$, (c) $H_3\%Urea$, (d) $H_{4.5}\%Urea$, and (e) $H_6\%Urea$ systems.

Supporting Videos

Video S1. Growth process of CO₂ hydrate in oceanic sediments for the H_{3%}Urea system. Illite is displayed as polyhedral, *i.e.*, yellow (Si atom) and pink (Al atom). Pink, blue, magenta, orange, and violet represent CO₂, Cl⁻, Na⁺, K⁺, and urea, respectively. Hydrate cages are shown as sticks in various colors (green for 5¹², blue for 5¹²6², red for 5¹²6³, orange for 5¹²6⁴, cyan for 4¹5¹⁰6², purple for 4¹5¹⁰6³ and pink for 4¹5¹⁰6⁴).

Video S2. Dissociation process of CO₂ hydrate in oceanic sediments for the H_{3%}Urea system. Illite is displayed as polyhedral, *i.e.*, yellow (Si atom) and pink (Al atom). Pink, blue, magenta, orange, and violet represent CO₂, Cl⁻, Na⁺, K⁺, and urea, respectively. Hydrate cages are shown as sticks in various colors (green for 5¹², blue for 5¹²6², red for 5¹²6³, orange for 5¹²6⁴, cyan for 4¹5¹⁰6², purple for 4¹5¹⁰6³ and pink for 4¹5¹⁰6⁴).

Video S3. Tension process of CO₂ hydrate-illite interface in oceanic sediments for the H_{3%}Urea system. Illite is displayed as polyhedral, *i.e.*, yellow (Si atom) and pink (Al atom). Pink, blue, magenta, orange, and violet represent CO₂, Cl⁻, Na⁺, K⁺, and urea, respectively. Hydrate cages are shown as sticks in various colors (green for 5¹², blue for 5¹²6², red for 5¹²6³, orange for 5¹²6⁴, cyan for 4¹5¹⁰6², purple for 4¹5¹⁰6³ and pink for 4¹5¹⁰6⁴).

Video S4. Compression process of CO₂ hydrate-illite interface in oceanic sediments for the H_{3%}Urea system. Illite is displayed as polyhedral, *i.e.*, yellow (Si atom) and pink (Al atom). Pink, blue, magenta, orange, and violet represent CO₂, Cl⁻, Na⁺, K⁺, and urea, respectively. Hydrate cages are shown as sticks in various colors (green for 5¹², blue for 5¹²6², red for 5¹²6³, orange for 5¹²6⁴, cyan for 4¹5¹⁰6², purple for 4¹5¹⁰6³ and pink for 4¹5¹⁰6⁴).

Supporting Reference

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