

Computer simulation evidence of the molecular mechanism of makeup removal using cleansing foam

~Is an "in silico formulator" superior to a human formulator?~

Yokoyama, Takahiro¹; Miwake, Hideki²; Hamaguchi, Masugu³; Nakatake, Ryouichi²; and Arai, Noriyoshi^{1*}

¹ Department of Mechanical Engineering, Keio University;

² Research Institute, Fancl Corporation ;

³ Kirin Central Research Institute, Kirin Holdings

Introduction:

Background

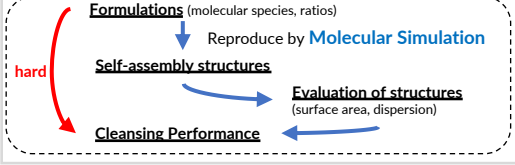
Surfactant-type cleansing agents: cleansing foams have excellent rinsing properties and eco-friendliness but also weak removability, since they are water-based [1]. In this context, improving the cleansing performance of cleansing foams is a high demand issue. However, cleansing foams are typically composed of many ingredients, making it difficult to find the best formulation from numerous combinations.

Molecular simulations can provide visualizations of the assembly structure [2,3] and the cleansing process at nano spatial-temporal scales, which are difficult to observe experimentally. To elucidate the molecular mechanism of cleansing will help reduce the high cost of formulation design.

Aim

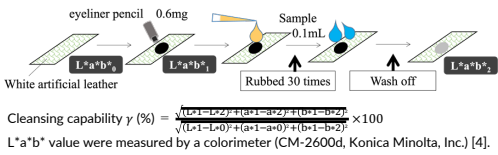
Clarifying the molecular mechanism of cleansing process by reproducing various formulations using molecular simulations and comparing their structures with cleansing capabilities.

Overview



Materials & Methods:

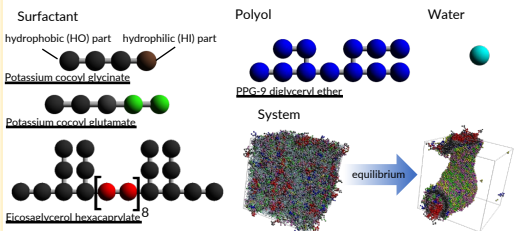
1. Evaluation of cleansing capability γ



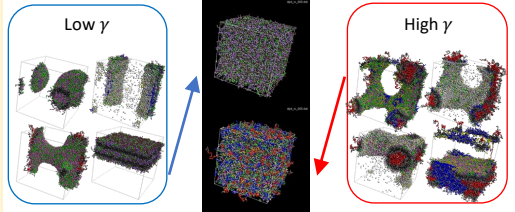
2. Dissipative Particle Dynamics (DPD)

$$m_i \frac{dv_i}{dt} = f_i = \sum_{j \neq i} -\frac{\partial U_{ij}}{\partial r_{ij}}$$

Coarse-grained simulation models

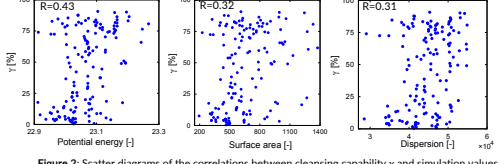


Self-assembled morphologies

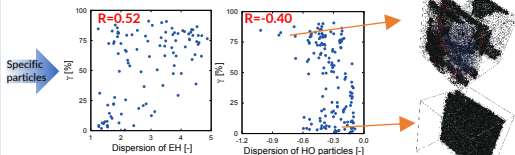


Results & Discussion:

Correlations between performance and morphology

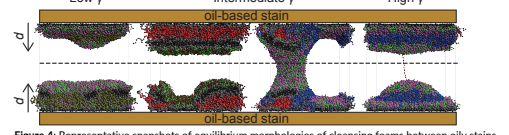


Using these indexes, we were able to grasp the change of structures quantitatively. The more complex the self-assembled structure, the higher the performance.

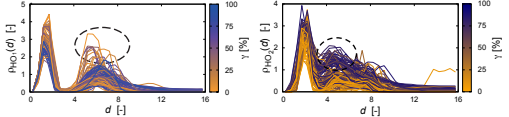


Not only the entire structures, but also the distributions of particles contribute to the cleansing performance, such as non-ionic surfactant (EH), and HO particles.

Self-assembled morphologies between oily stains



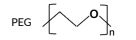
Divide HO group to strongly hydrophobic HO1 and weakly hydrophobic HO2 (HO1: alkyl chain, HO2: ether bond, ester bond, and amido bond)



The heights of the 2nd peak of $\rho_{HO1}(d)$ are higher for the formulations with low γ . The heights of the 2nd peak of $\rho_{HO2}(d)$ are higher for the formulations with high γ .

Future task

- 148 recipes consisting of 13 most frequently used molecules
- 64 additional recipes with 30 newly modeled molecules
- $R=0.67 \rightarrow R=0.46$
- Recipes containing molecules with PEG group.
- Analysis of formulations containing specific groups such as PEG (which belongs to HO2) remains a challenge.



Conclusions:

We investigated the mechanism of cleansing by simulating the self-assembled structures of cleansing foams.

- Various self-assembled structures were observed depending on the molecular species composing the system and their ratios.
- The complexity of a structure is one of the features of high-performance cleansing. However, the correlation between the performance of cleansing agents and their entire self-assembled structures are not strong enough.
- Focused on the inner distribution of aggregates, it was discovered that certain particles (molecules) such as EH or HO particles are crucial in cleansing process.
- The distribution of (HO1, HO2) particles when the cleansing foam is adsorbed on the dirt is also a significant clue to realize a high-performance cleansing.

References:

[1] K. Watanabe et al., *IFSCC Magazine*, **7**, 310–318 (2004).
 [2] R. Goetz et al., *J. Chem. Phys.*, **108**, 7397–7409 (1998).
 [3] A. Sarkar et al., *J. Am. Chem. Soc.*, **142**, 7606–7617 (2020).
 [4] T. Iwanaga et al., *Journal of Society of Cosmetic Chemists of Japan*, **39**, 186–194 (2005).