

Supplement Materials

Characteristic structural knowledge for morphological identification and classification in meso-scale simulations using principal component analysis

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Table S1. Parameters used in DPD simulation in dimensionless unit.

Parameter	
Particle number, N	24,000
Simulation box volume	8000
Coarse-grained degree, N_m	5
Spring constant, C_{ij}	4
Thermal energy, $k_B T$	1
Repulsion parameter of the same beads, a_{ii}	131.50
Repulsion parameter of the different beads, a_{ij} at temperature of 393 K	142.18
Time step, Δt	0.01
Equilibrium step number	1×10^6
Production step number	$\geq 1 \times 10^7$

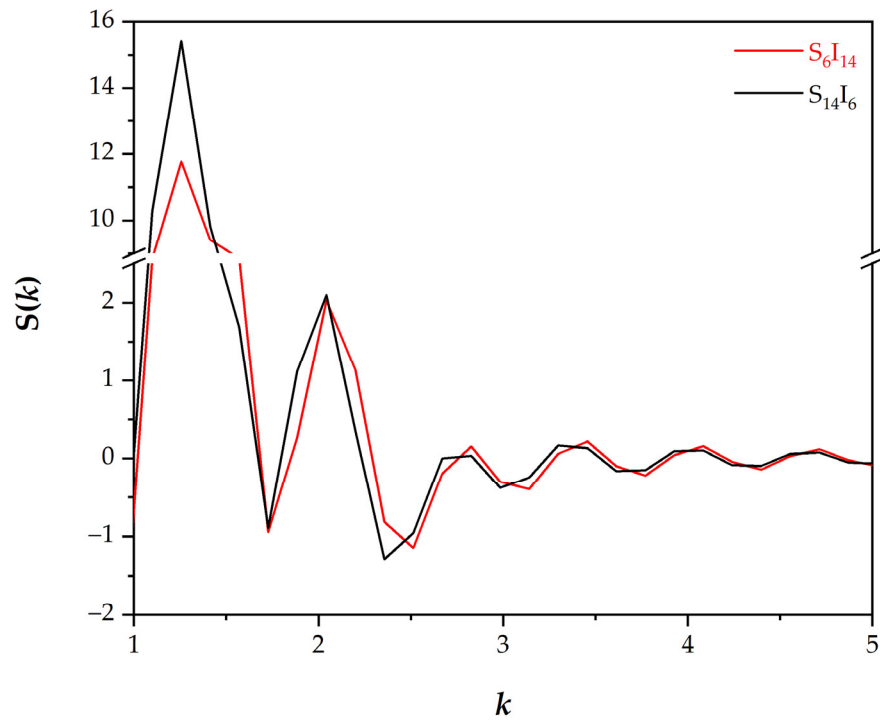


Figure S1. Structure factors $S(k)$ between chain-chain types of the gyroidal morphologies of S_6I_{14} and $S_{14}I_6$ were represented as red and black lines, respectively.

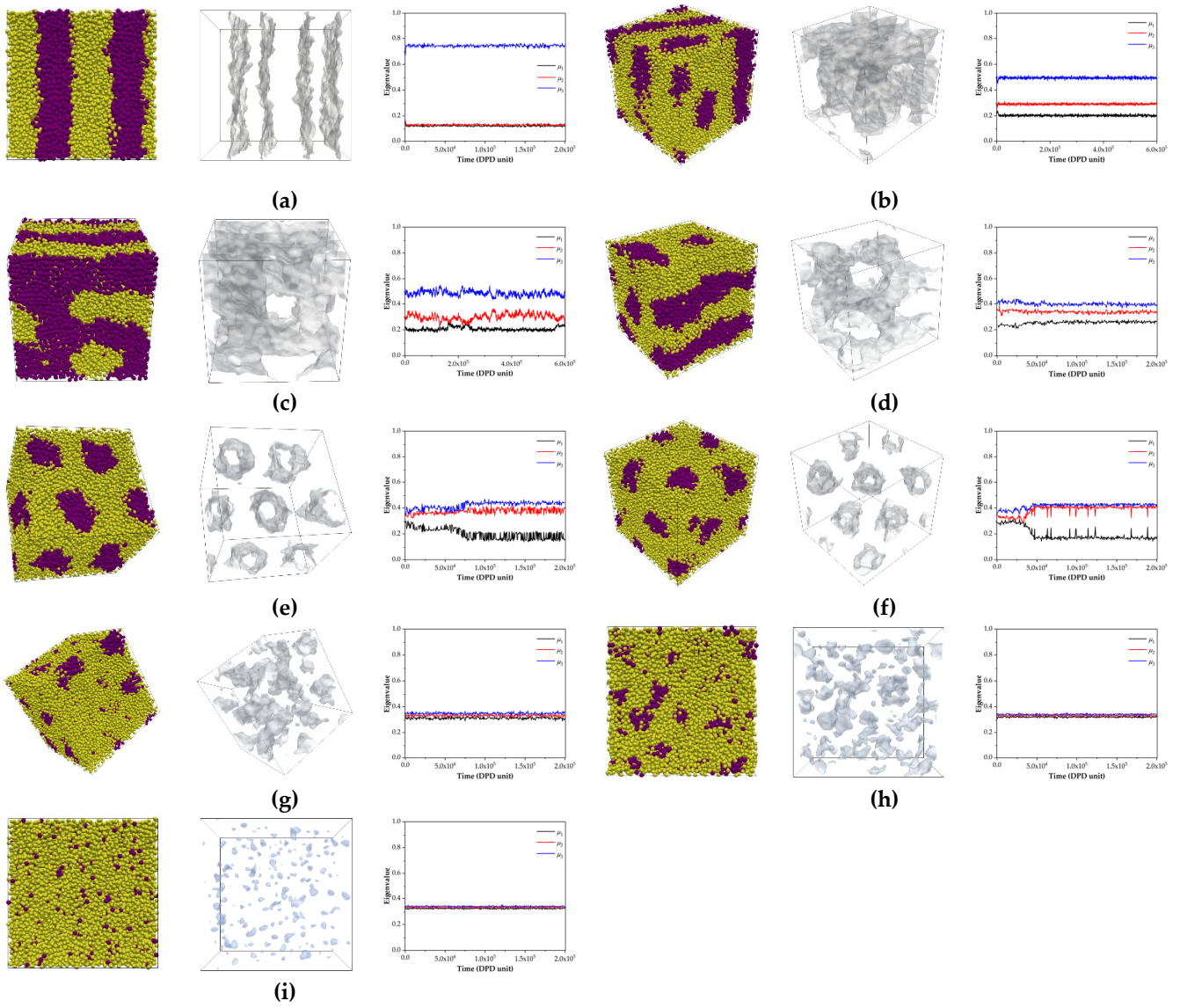


Figure S2. Panels in the composite figures illustrate apparent bead-arrangement, isosurface between distinct bead types and order parameter sorting from the left to right of (a) S11I9, (b) S12I8, (c) S13I7, (d) S14I6, (e) S15I5, (f) S16I4, (g) S17I3, (h) S18I2, and (i) S19I1, respectively. The order parameters were calculated using an executable, namely isosueface.exe, that is an implemented tool in DL_MESO software. Yellow and violet beads represent coarse-grained beads of styrene and isoprene, respectively.

Script S1: Structural knowledge analysis and PCA visualization
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Libraries

```
library(tidyverse)
library(ggforce)
library(factoextra)
library(ggrepel)
library(corrplot)
```

Data

```
data <- <name_of_excel_file>[<range_of_row>,<range_of_column>]
data
data.frame <- as.matrix.data.frame(data)
data.frame
```

Run the PCA

```
pca <- prcomp(data.frame,scale = TRUE)
pca
```

```
summary(pca)
```

```
eig.val <- get_eigenvalue(pca)
eig.val
```

```
options(ggrepel.max.overlaps = Inf)
```

```
fviz_eig(pca)
```

```
fviz_pca_ind(pca,title = "",geom = "point",col.ind = "cos2", gradient.cols = c("#00AFBB", "#E7B800", "#FC4E07"))
```

```
fviz_pca_var(pca,title = "",label = "none",col.var = "cos2",gradient.cols = c("#00AFBB", "#E7B800", "#FC4E07"))
```

```
res.var <- get_pca_var(pca)
```

```
res.var$coord
```

```
res.var$contrib
```

```
res.var$cos2
```

```
corrplot(res.var$cos2, is.corr = TRUE,)
```

#