Phase behaviour of block copolymer melts with arbitrary architecture

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Abstract

The Leibler theory [L. Leibler, Macromolecules \textbf{13} 1602 (1980)] for microphase separation in AB block copolymer melts is generalized for systems with arbitrary topology of molecules. A diagrammatic technique for calculation of the monomeric correlation functions is developed. The free energies of various mesophases are calculated within the second-harmonic approximation. Model highly-branched tree-like structures are considered as an example and their phase diagrams are obtained. The topology of molecules is found to influence the spinodal temperature and asymmetry of the phase diagrams, but not the types of phases and their order. We suggest that all model AB block-copolymer systems will exhibit the typical phase behaviour.
I. INTRODUCTION

One of the biggest challenges for theoretical polymer science is to predict the phase behaviour of a polymer system with given molecular structure. The required theory should explicitely contain molecular parameters and be able to predict thermodynamically stable phases (whatever they are) and temperatures of transitions among them. Apart from the theoretical interest this problem is of great industrial importance. Since the mechanical properties of the phases are studied quite well it is useful to predict the conditions at which these phases are stable. In the absence of such predictions the experimental search could take even years. In the present situation of a rapidly increasing number of novel materials even a crude theoretical model could help a great deal in direct applications.

Among the systems of interest are solutions of homopolymers, block copolymer melts, blends and so on. In this article we focus our attention on the A-B block copolymer melts. Such systems can only differ in the total content of one component, the length of molecules and the way the A and B blocks are arranged in one molecule. As a function of these parameters and temperature, the following phases were observed experimentally: the lamellar phase (LAM), the hexagonal phase (HEX), the body-centered cubic phase (BCC) and the bicontinuous gyroid phase (GYR) \[1,2\]. However, the structure of these phases strongly depends on temperature. Just below the order-disorder transition temperature the chains are slightly stretched and the mesophases consist of periodic patterns with only a small increase in the content of one component. This situation is called the weak-segregation regime. On contrary, the strong-segregation regime (very low temperatures) is characterized by domains which are almost pure in one component. Both regimes are limiting cases and therefore can be studied analytically. In the intermediate regime there is no small parameter and only numerical investigations are possible \[3\].

The first weak-segregation theory of AB melts is by Leibler \[4\] who considered a model Gaussian diblock copolymer. He was able to estimate the order-disorder transition temperature and predict the following sequence of transitions: \(\text{DIS} \rightarrow \text{BCC} \rightarrow \text{HEX} \rightarrow \text{LAM}\) with
decrease of temperature. Later, after the GYR phase was observed experimentally [3,4], Milner and Olmsted [7] used the second-harmonic approximation and predicted the GYR phase to be stable in between the HEX and LAM phases. The same conclusion was drawn in [3,5] and in the Landau-Brazovskii theory of Hamley and Podneks [9,10].

The method developed by Leibler was applied to other topologies of block copolymer molecules. Thus, Mayes and de la Cruz considered an ABA triblock molecule [11], de la Cruz and Sanchez studied star- and simple graft-copolymers [12]; Dobrynin and Erukhimovich developed a computer-aided tool to calculate phase diagrams for various topologies [13]. However, in all these studies the authors started with a fixed molecular topology. To consider another topology one needed to go through the whole calculation procedure again, from the very beginning. In this work we show how the Leibler approach can be used for an arbitrary topology of molecules. Section I repeats some main points of Leibler’s paper [4] and introduces a diagrammatic technique that helps to deal with molecules of arbitrary topology. The first step in this direction was made by Read [14], who invented a way to calculate structure factors of polymers of arbitrary complexity. We go further and show how to calculate higher vertex functions for such systems. Section II describes the way to calculate the free energies of different mesophases and discusses some of the approximations. We consider the face-centered cubic phase (FCC), the ordered bicontinuous double-diamond phase, the simple cubic (SC) and the square lattice (SL) phases in addition to the classical phases. In the Results and Discussion we apply the developed technique to a model dendrimer system which has recently attracted wide experimental [15] and theoretical [16] attention. The phase diagrams of various dendrimers are presented and their features are discussed. The results for many more systems that were not included in this publication are presented as supplementary materials on the Internet: http://rugmd4.chem.rug.nl/~morozov/research.html.

Finally, we summarize our work in the Conclusion and propose some possible developments of the method.
II. MEAN-FIELD THEORY

A mean-field theory of incompressible AB block copolymer melts was first formulated by Leibler [4]. In his approach a local deviation of the density of A component from its average value plays the role of an order parameter ψ(r). He found that the coefficients in the phenomenological free energy expansion

\[ F = F_0 + \frac{1}{2} \int \int \Gamma_2(q_1, q_2) \psi(q_1) \psi(q_2) + \frac{1}{3!} \int \int \int \Gamma_3(q_1, q_2, q_3) \psi(q_1) \psi(q_2) \psi(q_3) + \frac{1}{4!} \int \int \int \int \Gamma_4(q_1, q_2, q_3, q_4) \psi(q_1) \psi(q_2) \psi(q_3) \psi(q_4) + \cdots \]  

(2.1)

can be connected with microscopic properties of the system. This possibility is based on the assumption that in a melt a single chain behaves as almost ideal. This constitutes the so-called random phase approximation (RPA) which expresses the monomer density correlation functions in terms of the correlation functions of a Gaussian chain. For the vertex functions in (2.1) it gives [4]:

\[ \Gamma_2(q_1, q_2) = \delta(q_1 + q_2) \left[ \frac{S_{AA}(q_1) + S_{BB}(q_1)}{S_{AA}(q_1) S_{BB}(q_1) - S_{AB}^2(q_1)} - 2 \chi \right] \]
\[ \Gamma_3(q_1, q_2, q_3) = -G^{(3)}_{ijk}(q_1, q_2, q_3) \left[ \frac{S_{ik}^{-1}(q_1)}{S_{kk}^{-1}(q_1) - S_{ij}^{-1}(q_1)} \right] \]
\[ \times \left[ S_{jk}^{-1}(q_2) - S_{jB}^{-1}(q_2) \right] \left[ S_{kA}^{-1}(q_3) - S_{kB}^{-1}(q_3) \right] \]  

(2.2)

\[ \Gamma_4(q_1, q_2, q_3, q_4) = \gamma_{ijkl}(q_1, q_2, q_3, q_4) \left[ \frac{S_{kA}^{-1}(q_1)}{S_{ik}^{-1}(q_1) - S_{ij}^{-1}(q_1)} \right] \left[ \frac{S_{jA}^{-1}(q_2)}{S_{jk}^{-1}(q_2) - S_{jB}^{-1}(q_2)} \right] \]
\[ \times \left[ S_{kA}^{-1}(q_3) - S_{kB}^{-1}(q_3) \right] \left[ S_{jA}^{-1}(q_4) - S_{jB}^{-1}(q_4) \right] \]

where

\[ \gamma_{ijkl}(q_1, q_2, q_3, q_4) = -G^{(4)}_{ijkl}(q_1, q_2, q_3, q_4) \]
\[ + \int p S_{mn}^{-1}(p) \left[ G_i^{(3)}(q_1, q_2, p) G_j^{(3)}(q_1, q_3, q_4) + G_k^{(3)}(q_1, q_3, p) G_j^{(3)}(q_1, q_2, q_4) \right] \]
\[ + G_i^{(3)}(q_1, q_4, p) G_j^{(3)}(q_1, q_3, q_2) \]  

(2.3)

The summation over repeating indices is assumed. The second, the third and the fourth order correlation functions of the Gaussian chain are denoted by \( S_{ij} \), \( G_{ijkl}^{(3)} \) and \( G_{ijkl}^{(4)} \), respectively.
\(i, j, k, l = A, B\); \(S_{ij}^{-1}\) are the elements of the inverse matrix \(|S_{ij}|^{-1}\). The Flory-Huggins parameter describing the strength of interaction between A and B monomers is denoted by \(\chi\).

A general method of calculating the Gaussian correlation functions was developed in Appendix B of [4]. It is based on the fact that the Gaussian distribution is fully determined by its first two moments. As a result the probability to find several monomers at given positions on the chain is expressed as a product of the 2-point correlation functions of the consequently neighbouring monomers along the chain. For example,

\[
\begin{align*}
\mathbf{r}_1 \quad \mathbf{r}_2 \quad \mathbf{r}_3 & = P(\mathbf{r}_1, \mathbf{r}_2)P(\mathbf{r}_2, \mathbf{r}_3) \\
\end{align*}
\]

The correlation functions \(S_{ij}, G^{(3)}_{ijk}\) and \(G^{(4)}_{ijkl}\) are calculated by averaging the Fourier transforms of expressions like one in the previous example over all possible positions of monomers along the chain. At this point one needs to introduce some topology of the polymer molecule in order to perform such an averaging. In [4] this was done for a diblock molecule. A triblock molecule was considered in [11]. More sophisticated architectures were studied in [13]. We generalize this procedure for arbitrary topology of a polymer molecule.

In the following we use the graphical method formulated in Appendix A. A correlation function of a given order is calculated by drawing all possible (but topologically distinct) diagrams of this order. Thus,

\[
\begin{align*}
S_{ij} & = \quad + \\
G^{(3)}_{ijk} & = \quad + \\
\end{align*}
\]
This is just a schematic representation which shows that an $n$-th order diagram can contain up to $n$ blocks. The corresponding contributions to the correlation function are obtained by connecting these blocks in all possible ways. The topologically different connections are presented in Appendix A. The contributions to the correlation functions are calculated in Appendix B.

We want to note that not all diagrams contribute to the correlation functions of the system with given topology. One restriction is connected with the type of the correlation function we want to calculate. For example, the equation for $S_{AB}$ only contains diagrams with the first label placed on some A block and the second label placed on a B block. Therefore the first diagram in the general equation for $S_{ij}$ does not contribute to $S_{AB}$. The other factor reducing the number of diagrams is a molecular architecture. Thus, for a linear multiblock molecules all 'star' and 'fork' diagrams are impossible and should not be considered.

The final expressions for the correlation functions are obtained in the following way. Suppose we want to calculate $G_{i_1,i_2,...}$. To do this we need to place the first label on any block of the type $i_1$, the second label on any block of the type $i_2$ and so on. The resulting diagram will be one of the diagrams calculated in Appendix B. The corresponding expression is written down. Then we remove all labels and place them again avoiding previous configurations. This procedure is repeated until we covered all possible arrangements of the labels. The obtained expression is the equation for the required $G_{i_1,i_2,...}$.

Once we have calculated the correlation functions for a given system, its free energy can be obtained with the help of (2.1-2.3). In the next section this Landau-Ginzburg free energy will be a starting point for our analysis of the phase transitions.

It is obvious that the formulated approach is general and within this framework all
systems can be considered.

### III. CALCULATION OF PHASE DIAGRAM

In this section we show how to calculate a phase diagram of a system once its free energy functional (2.1) is known. Apart from some technical details this was developed elsewhere [17,18,22,23].

In the beginning we need to determine the order-disorder transition (ODT) temperature. The disordered phase can be described by the Landau-Ginzburg Hamiltonian with only a quadratic term left [17,23]. The probability to create a spatial density fluctuation \( \psi(q) \) is given by

\[
P[\psi] \sim \exp \left[ -\frac{1}{2} \int q \Gamma_2(q) \psi(q) \psi(-q) \right]
\]

The typical life-time of such a fluctuation is \( \tau \sim \frac{1}{\Gamma_2(q)} \). At some \( q_* \), temperature (or Flory-Huggins parameter) and total content of A monomers in the molecule \( \left( f = \frac{N_A}{N} \right) \), \( \Gamma_2(q_*) \) reaches zero. The probability to create such a mode tends to unity, while its life-time \( \tau_{q_*} \) diverges. It means that at this temperature the disordered phase becomes absolutely unstable with respect to the appearance of an ordered mesophase with typical domain size \( \sim q_*^{-1} \). Therefore, the set of equations:

\[
\Gamma_2(q_*, (\chi N)_s) = 0 \quad \left( \frac{\partial \Gamma_2(q, (\chi N)_s)}{\partial q} \right)_{q_*} = 0 \quad , \quad \left( \frac{\partial^2 \Gamma_2(q, (\chi N)_s)}{\partial q^2} \right)_{q_*} > 0
\]

determines the spinodal temperature (which is very close to the ODT value [1]) and the typical lengthscale \( q_* \).

As the next step we choose a set of mesophases which will possibly enter the phase diagram. Then, for each mesophase we represent the order parameter field \( \psi(q) \) in such a way that it possesses the symmetry of the mesophase. This representation necessarily contains some unknown parameters. Next \( \psi(q) \) should be plugged in the eq.(2.1) and the
resulting expression should be minimized with respect to the parameters. This procedure
gives the free energies of different mesophases as functions of temperature and composition.
The molecular properties enter via the coefficients in eq.(2.1). For the given temperature
(below the ODT) and composition $f$ the mesophase with the lowest free energy will be the
stable one. The phase diagram is drawn by varying temperature and $f$ over some ranges.

The way to ascribe a given symmetry to the order parameter is to expand it in the full
orthonormal set of functions with the given symmetry. The most popular choice is to use
an expansion in plane waves:

$$
\psi(q) = \sum_i A_i \left( e^{i\phi_i} \delta_{q,k_i} + e^{-i\phi_i} \delta_{q,-k_i} \right) \tag{3.3}
$$

where $\{k_i\}$ is a set of vectors defining a lattice of the given symmetry, $\phi_i$ and $A_i$ are the
arbitrary phases and amplitudes; $\delta_{i,j}$ denotes the Kronecker delta symbol. If the rotational
symmetry in the system is not broken we can rewrite (3.3) in the following way:

$$
\psi(q) = \sum_i a_i \sum_{j=1}^{m_i} \left( e^{i\phi_j^{(i)}} \delta_{q,k_j^{(i)}} + e^{-i\phi_j^{(i)}} \delta_{q,-k_j^{(i)}} \right) \tag{3.4}
$$

where the first sum runs over the Brillouin zones or coordination spheres in the reciprocal
space; $m_i$ is the number of vectors in the $i$-th zone. In practice we consider only the first few
terms in the series (3.4). The mathematical reason for that is to simplify the minimization
procedure. From the physical point of view, immediately below the transition temperature
the appearing mesophase is not well developed and we do not need too many details in
the representation (3.4). (As a rule, the length of the $q$-vectors increases with the number
of the zone. Therefore, the more terms are left in (3.4), the more detailed density profile
we get.) The next question is how many terms we should leave in (3.4). Unfortunately,
there is no quantitative criterion where to cut the density expansion. In the past, several
possibilities were considered. In the classical work [4] the expansion (3.4) was restricted
to $i = 1$. The so-called second-harmonic approximation ($i = 2$) [18] was formulated in
[24, 25, 28, 19] in order to study complicated bicontinuous phases. Up to several hundred
harmonics were used in the numerical method of Matsen [8] (however, the authors studied
the intermediate/strong segregation regimes and, therefore, our discussion is not directly applicable to their model). Comparison of their results shows that higher harmonics not only result in some non-significant shift of the phase boundaries, but also stabilize or destabilize several morphologies. In such a situation the number of harmonics is crucial.

In this paper we adopt the criterion derived in [10,9]. Instead of the mean-field theory Hamley and Podneks studied system with composition fluctuations. In their model the cut-off value for $q_n$ (the $q$-vector from the $n$th shell) appears naturally as an adaptation of the Ginzburg criterion:

$$\left( \frac{q_n}{q_*} \right)^2 - 1 \ll 1 \quad (3.5)$$

We apply this criterion to our mean-field theory in order to keep uniformity with the fluctuation approach (a similar criterion was used in [27]). There are several ways to support our choice. One type of argument is connected with the applicability region for our theory. In the RPA there is essentially one lengthscale associated with the gyration radius of molecules. It is intrinsically connected with the typical wavevector $q_*$. Therefore, the representation of the density profile by higher harmonics with $q \gg q_*$ is in strong contradiction with the used theory. (A similar idea was formulated in a different way in [28].) One can also speculate that near the spinodal the inequality (3.5) will always hold in the proper temperature range. The questions of applicability of (3.5) and its influence on the phase diagram will be addressed in future studies.

The cut-off (3.3) implies that only one harmonic should be used for LAM, HEX, BCC, SC and SL phases, while FCC, GYR and OBDD phases are represented by two-harmonic density profile. The basis vectors for LAM, HEX, BCC, SC and SL phases, as well as the first-shell vectors for FCC and OBDD (which coincides with the BCC first shell), are presented in [4,18]. For the rest we use (more detailed information can be found in [29]).
are all possible combinations of two, three and four $k$ energy (2.1). The integration is trivial since expression contains terms proportional to the vertex functions $\Gamma$. This function cannot be directly used for the arguments with the different moduli. However, in $\Delta$ we can approximate the $m$-th order vertex function in the form $\Gamma_m = \Gamma_2, \Gamma_3$ and $\Gamma_4$ which arguments are all possible combinations of two, three and four $k^{(i)}$ that add up to zero. For the phases represented by the two-harmonic density profile some of these terms will be the functions of the wave-vectors from the different shells. However, while calculating the vertex functions (Section 2 and Appendix A and B) we assumed everywhere that all wave-vectors have the same modulus $q_*$. Correspondingly, we obtained the $m$-th order vertex function in the form $\Gamma_m(q_1, \ldots, q_m) = \Gamma_m(x, h^{(m)}) \ (n_i = \frac{q_i}{q_1}, h^{(m)}$ and $x$ are defined in the Appendix A). This function cannot be directly used for the arguments with the different moduli. However, the following trick helps to avoid this problem.

The criterion (3.3) implies that $q_j = q_* + \Delta_j$, where $\Delta_j \ll q_*$. Therefore, to the first order in $\Delta$ we can approximate the $m$-th order vertex function of $q$-s with the different length by

$$\Gamma_m(q_1, \ldots, q_m) \approx \Gamma_m(q_1, \ldots, q_m) + \sum_{\alpha=1}^{m} \kappa_\alpha \Delta_\alpha,$$

where

$$\kappa_\alpha = \left. \frac{\partial \Gamma_m(q_1, \ldots, q_\alpha, \ldots, q_m)}{\partial q_\alpha} \right|_{q_\alpha = q_*},$$
Any vertex function is a symmetric function of its arguments. Therefore, \( \kappa_\alpha = \kappa \) is independent on \( \alpha \). Let us now put all \( \Delta_\alpha = \Delta \). This gives

\[
\Gamma_m(q_1, \ldots, q_m) \approx \Gamma_m(q_*n_1, \ldots, q_*n_m) + m\kappa\Delta
\]

But this expression should be identical to:

\[
\Gamma_m(x_* + \delta x, h^{(m)}) \approx \Gamma_m(x_*+h^{(m)}) + \frac{d\Gamma_m(x, h^{(m)})}{dx}\Big|_{x=x_*} \delta x
\]

where \( x = (q_* + \Delta)^2R_G^2 = x_*+\delta x \), \( \delta x \approx 2x_*\Delta \). Comparing (3.7) with (3.6) gives

\[
\kappa = \frac{2x_*}{mq_*} \frac{d\Gamma_m(x_*+h^{(m)})}{dx}\Big|_{x=x_*}
\]

Finally,

\[
\Gamma_m(q_1, \ldots, q_m) \approx \Gamma_m\left(\sum_{\alpha=1}^{m} \frac{\Delta_\alpha}{q_*}, h^{(m)}\right) = \Gamma_m(x_*+h^{(m)}) + \frac{2x_*}{m} \frac{d\Gamma_m(x_*+h^{(m)})}{dx}\Big|_{x=x_*} \sum_{\alpha=1}^{m} \frac{\Delta_\alpha}{q_*}
\]

The angle variables \( h^{(m)} \) are calculated from the vectors \( q_*n_i \). Here we assumed that the latter sum up to zero, which is only \textit{approximately} true. The approximation (3.8) allows us to take into account the small differences in the length of the wave-vectors from the first and the second shells.

The last step involves the minimization of the free energies with respect to the arbitrary phases \( \phi_j^{(i)} \). The algebra is labouring but straightforward. Denoting \( a_1 \) by \( a \), and \( a_2 \) by \( b \) we obtain

\[
F_{FCC} = 4a^2\Gamma_2(x_*) + 3b^2\Gamma_2\left(\frac{4}{3}x_*\right) + 12a^2b\Gamma_3\left(\frac{2}{\sqrt{3}} - 1\right) + \Xi_1^{(F)}a^4 + \frac{3}{4}\Xi_2^{(F)}b^4 + 4\Xi_3^{(F)}a^2b^2
\]

\[
F_{OBDD} = 6a^2\Gamma_2(x_*) + 4b^2\Gamma_2\left(\frac{3}{2}x_*\right) + 8a^3\Gamma_3(0) + \frac{3}{2}\Xi_1^{(O)}a^4 + \Xi_2^{(O)}b^4 + 2\Xi_3^{(O)}a^2b^2
\]

\[
F_{GYR} = 12a^2\Gamma_2(x_*) + 6b^2\Gamma_2\left(\frac{4}{3}x_*\right) + 8a^3\Gamma_3(0) - 12a^2b\Gamma_3\left(\frac{2}{\sqrt{3}} - 1\right)
\]

\[
+ 8b^2\Gamma_3\left(2\sqrt{3} - 3\right) + 3\Xi_1^{(G)}a^4 + \frac{3}{2}\Xi_2^{(G)}b^4 + 2\Xi_3^{(G)}a^3b + 2\Xi_4^{(G)}a^2b^2
\]

where the coefficients \( \Xi \) are given in Appendix C.
The one-harmonic free energies for the LAM, HEX, BCC, SC and SL phases are presented in [18]. As it was noticed in [30], we have two choices for the FCC free energy, which arise from the different solutions of the phase minimization equations. The first solution is presented in (3.9). The second (NFCC in the notation of [30]) was never found to be stable in our calculations.

We want to emphasize that the second harmonic approximation was used extensively by many authors [18,24–26,19,7]. However, almost all of them used the two-harmonic representation of the density profile for all phases. On contrary, we apply the criterion (3.5) and represent only the FCC, OBDD and GYR morphologies in the two-harmonic approximation. This approach was also tried in several studies [10,9,27]. The main difference with our work is that we not only keep the complicated angle dependence of the vertex functions, but also take into account their dependence on the wave vectors moduli.

IV. RESULTS AND DISCUSSION

In this section we apply the developed formalism to particular systems. As an example we will treat dendrimeric tree-like structures defined as follows. The origin of the molecule is a branching point with \( n_1 \) A-blocks of the same length \( f_1 \) attached to it. All \( n_1 \) A-blocks are topologically equivalent. They form a so-called first generation of the molecule. The end of each block is also a branching point with \( n_2 \) B-blocks of the same length \( f_2 \) attached to it. All \( n_1 n_2 \) B-blocks are topologically equivalent and form the second generation. The third generation again consists of A-blocks and so on till the \( g \)-th generation - the highest generation of the molecule. We introduce a \( g \)-dimensional topology vector \( \mathbf{n} = \{ n_1, n_2, \ldots, n_g \} \). The \( i \)-th element of this vector defines the number of blocks (A-blocks if \( i \) is odd, and B otherwise) originated from each block in the \((i-1)\)-th generation. The lengths of the blocks within one generation are necessarily the same but can be different in different generations. It is therefore useful to measure the lengths of the A-blocks in terms of the length of the first generation \( f_A \equiv f_1 \) (\( f_B \equiv f_2 \) for the B-blocks correspondingly). Thus, we
introduce the following $g$-dimensional vector $\tau = \{1, 1, \tau_3, \ldots, \tau_g\}$. The length of the $i$-th generation is given by

$$f_i = \tau_i (f_A \epsilon_i + f_B (1 - \epsilon_i)) \quad (4.1)$$

where

$$\epsilon_i = \begin{cases} 1 & i \text{ is odd} \\ 0 & \text{otherwise} \end{cases}$$

If we now introduce the total content of the A-component in the molecule $p$, then the following equations hold:

$$p = f_A \sum_{i=1}^{g} \epsilon_i \tau_i \prod_{j=1}^{i} n_j$$

$$1 - p = f_B \sum_{i=1}^{g} (1 - \epsilon_i) \tau_i \prod_{j=1}^{i} n_j \quad (4.2)$$

and for the given topology (fixed $g$, $n$ and $\tau$) and the overall composition $p$, $f_A$ and $f_B$ are determined in a unique way.

Next we want to be able to calculate the distance between two given blocks. We introduce the system of coordinates and number all blocks originated from some block on the $(i-1)$-th generation from 1 to $n_i$. Then the position of some block is given by a set of numbers, that define the trajectory from the origin of the molecule to this block, and will be denoted by a $g$-dimensional vector $R$. For the $i$-th generation block all components of $R$ higher than $i$ are equal to zero. (see Fig.1). The shortest distance between the ends of two blocks with positions $R^{(1)}$ and $R^{(2)}$ is given by

$$\Delta = \sum_{i=1}^{g} \tau_i (f_A \epsilon_i + f_B (1 - \epsilon_i)) \left(1 - \delta_{R_i^{(1)}-R_i^{(2)},0}\right) \left(2 - \delta_{R_i^{(1),0}-R_i^{(2),0}}\right) \quad (4.3)$$

We reformulate the framework of Sections 1 & 2 in the form of algorithm, depicted schematically as follows:

1) Input: $g$, $n$ and $\tau$

2) Vary $p$ and $\chi N$ in the ranges of interest and for each point:
- calculate the vertex functions
- calculate the free energies for the mesophases
- find the stable mesophase (the one with the lowest free energy)

3) Output: the phase diagram

The calculation of the vertex functions is performed in the fashion described in Section 1. For example, to calculate the third order correlation function \( G_{AAB} \) we need to move two labels over all A-blocks, and one label over all B-blocks. For each configuration of the labels we need to determine the type of diagram and then use the length of the blocks with the labels for \( f_\alpha, f_\beta, \ldots \) and the distances between those blocks for \( \Delta_1, \Delta_2, \ldots \) in the equations from Appendix B. The resulting number should be added to \( G_{AAB} \).

To distinguish among the different types of diagrams (line, fork, star) we introduce the concept of \textit{pathways}. A pathway from one block to another one consists of the positions of all blocks lying in-between those two (so, the pathway is a set of maximally \( 2(g-1) \) vectors). Now suppose we know that the labels (three or four of them) occupy three different blocks. According to Appendix A they can form a line- or a star-type diagram. In the first case one of the blocks belongs to the pathway between the others, while in the second case it does not. Checking this condition allows us to distinguish among different types of diagrams (In the 4-block case the number of conditions increases but the general idea remains the same).

The procedure of the free energy minimization with respect to the amplitudes \( a \) and \( b \) is the same as that of Leibler [4] and is not presented here. We proceed with the results for different dendrimers.

Figs.2-7 show the phase behaviour of various architectures. Despite the differences in topology, number of blocks and branching points, all systems behave in a similar way. Their phase diagrams are very close to those of the diblock [4], triblock [11], stars and graft copolymers [12,13]. The total number of blocks and branching points influences the spinodal temperature, while the ratio \( n_A/n_B \) and the number of branching points influences the asymmetry of phase diagrams. (A similar observation was made in [12] where the authors
connected the increase of complexity of molecules with the shift of the spinodal temperature.) Moreover, the relative sizes of the stability regions for different phases are almost the same for all systems, while the types of phases and their order is completely independent of the topology of molecules. Therefore, we conclude that all mean-field model systems with Gaussian chains will have phase diagrams similar to Figs.2-7. This is in agreement with the recently proposed constituting block copolymer hypothesis [31] which explains the phase behaviour of complicated molecules on the basis of the behaviour of their constitutive blocks. For example, in the case of the two-generation dendrimer from Fig.2, the A$_1$B$_2$ miktoarm-stars are the building blocks of the molecules and their behaviour qualitatively corresponds to the phase diagram of the \{2, 2\}-dendrimer. Since we can always construct a complicated molecule from simple ones, like diblock, triblock and so on, the resulting phase behaviour is going to be typical. However, unlike the general theory of Sections 2 and 3, our model system does not consider structures where the ends of the blocks are the branching points with both A and B blocks attached to them. (This is necessary, for example, to model comb copolymers.) Therefore we cannot be sure that such systems will also have typical phase diagrams. There is also evidence of the opposite: Dobrynin and Erukhimovich found that the theory fails for the homo-stars $A_nB_n$ with $n > 5$ and the corresponding phase diagrams contain regions where the phase behaviour cannot be predicted within the used approach [13]. Nevertheless, the other phase diagrams show the same typical behaviour.

We also want to notice the complete absence of the OBDD morphology. This is not surprising even for the \{4, 4, 4\}-dendrimers with the 4-fold symmetry of molecules. As it was shown by Matsen and Bates [3], the OBDD phase has larger interface in comparison with the GYR phase and could be stable only at very low temperatures. It is however screened by other morphologies (BCC and HEX).

Another interesting feature is a notably big region of stability of the bicontinuous gyroid phase in Fig.6. For such molecules the highly branched structure and the relative conformational freedom of the end A-B blocks favor the GYR phase to the LAM phase. This is not a feature of the \{3, 3, 3\}- and \{4, 4, 4\}-dendrimers, for example, because there the outer
shell of the molecule consists only of the A blocks. As the AB dendrimers were not studied experimentally yet, we recommend the systems like \{5, 1, 3, 1\} to produce relatively easy the bicontinuous gyroid phase.

At the end we want to mention that our tree-like model is capable of describing stars, polyblocks, umbrellas, and many other architectures. Because of space limitations those results are not presented here.

V. CONCLUSION

In the present paper we have shown how to calculate phase diagrams for systems with arbitrary topology. The developed method is a generalization of the work by Leibler [1]. The monomeric correlation functions are calculated with the help of a special diagrammatic technique which allows us to treat an arbitrary molecular topology. Apart from the assumption that the polymers obey Gaussian statistics even below the order-disorder transition temperature, the calculation of the vertex functions is approximation-free. The free energies for different mesophases are calculated within the second-harmonic approximation as it was formulated by Hamley and Podneks [10,1]. The influence of this approximation and its relationship with the random-phase approximation requires additional study.

The architecture of polymer molecules is known to influence the phase diagram only slightly [12]. This was once again confirmed by our studies of highly-branched model dendrimers. We found that the topology of molecules influences the spinodal temperature and asymmetry of the phase diagrams, but not the types of phases and their order. The typical phase diagrams are presented in the Results and Discussion. That model system, however, does not cover all possible architectures. Many more phase diagrams together with the C-code used to obtain them can be found on the Internet: [http://rugmd4.chem.rug.nl/~morozov/research.html](http://rugmd4.chem.rug.nl/~morozov/research.html)

Our method can be a starting point for a fluctuation theory of systems with arbitrary topology. This will require some straightforward generalization of the Hartree approximation
by Brazovskii [32,33] and Fredrickson and Helfand [34], since for some systems (see, for example, Figs.4 and 5) the 4-th order vertex function $\Gamma_4(h_1,h_2)$ exhibits a strong angle-dependence and, therefore, cannot be approximated by the constant value $\Gamma_4(0,0)$ or any other. One of the possible ways for such a generalization was proposed by Dobrynin and Erukhimovich [35].

VI. ACKNOWLEDGMENTS

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APPENDIX A: DIAGRAMMATIC TECHNIQUE

We consider a polymer molecule consisting of \( n_A \) blocks of type A and \( n_B \) blocks of type B. The total number of monomers in the molecule is \( N \). We shall use Greek letters to number the blocks: \( \alpha, \beta, \gamma, \delta = 1 \ldots n_A + n_B \). The length of the \( \alpha \)-th block is \( L_\alpha = f_\alpha N \). We use solid lines to denote blocks and dashed lines to denote arbitrary distance between blocks. The circles denote the labeled monomers. We use \( s, s', s'' \ldots \) for the positions of the labels on their blocks. A diagram with \( n \) labels will be called the \( n \)-th order diagram. The labels act as sources of momentum. A circle labeled with \( a, b, c \ldots \) emits \( q_a, q_b, q_c \ldots \) respectively. The total momentum of any diagram should be zero, therefore all external lines should carry zero momentum. We also introduce a convention (which does not affect our results) that all momenta flow from left to right along the chain. This can be illustrated by the following:

![Diagram](image)

A given number of blocks can form topologically different diagrams:

2: line 3: line star

4: line fork star

Necessarily, all momenta flowing in and out of each branching point should add up to zero.

Following \[4\] we assume that all wavevectors have the same length \( |q| = q_* \) and introduce the following angle variables:
for the 3rd order diagrams:  \( q_i^2 = q_i^2 h_i^{(3)} \),  \( h^{(3)} = (1, 1, h) \)

for the 4th order diagrams:  \( (q_i + q_j)^2 = q_i^2 h_{ij}^{(4)} \)

where

\[
\begin{pmatrix}
4 & h_1 & 4 - h_1 - h_2 & h_2 \\
4 - h_1 - h_2 & h_2 & 4 - h_1 - h_2 & h_1 \\
4 - h_1 - h_2 & h_2 & 4 & h_1 \\
h_1 & 4 & h_2 & 4 - h_1 - h_2 \\
\end{pmatrix}
\]

The matrix \( h^{(4)} \) has two important symmetry properties: \( h_{ij}^{(4)} = h_{ji}^{(4)} \) and \( h_{ij}^{(4)} = h_{kl}^{(4)} \) for \( i \neq j \neq k \neq l \). These properties will be extensively used in Appendix B.

For any internal line we subscribe the Gaussian propagator:

\[
P(l) = e^{-\frac{P^2}{2}} , \quad x = q_s^2 R_G^2
\]

where \( R_G \) is the radius of gyration of an ideal chain of \( N \) monomers. The product of the propagators is integrated over all possible positions of all labels on their blocks. The result should be then multiplied with the symmetry factor arising from all possible permutations of labels that result in topologically the same diagram and keep labels on the blocks they initially occupied. Finally, the normalization constant \( 1/N \) is added.

The following functions will appear as the building blocks for all diagrams:

\[
F_f = \frac{1 - e^{-fx}}{x}
\]

\[
\mathcal{F}_f(h) = \begin{bmatrix}
F_{fh} & F_f & f \\
h^2 & h^2 & h^2 \\
\end{bmatrix}
\]

\[
J_f(h) = \frac{1}{x^2} \begin{bmatrix}
1 - e^{-fx} & e^{-hx} \\
1 - e^{-fx} & 1 - e^{-fx} - fxe^{-fx} & f + e^{-fx} - 1 \\
\end{bmatrix}
\]

\[
D_f(h) = \frac{1}{x^2} \begin{bmatrix}
hfx + e^{-hx} - 1 & f + e^{-fx} - 1 & \frac{f^2x^2}{2} \\
\end{bmatrix}
\]

\[
T_1(f, h) = \frac{1}{x^4} \begin{bmatrix}
\frac{fx}{h} + e^{-fx} & \frac{fx}{h} + e^{-fx} & \frac{2h - 3}{(h - 1)^2} + \frac{e^{-fhx}}{h^2(h - 1)^2} & 2h + 1 \\
\frac{f^2x^2}{2} e^{-fx} & \frac{f^2x^2}{2} e^{-fx} + f - 3 + 3e^{-fx} & \frac{f^2x^2}{2} + 3 - 2f - (3 + fx)e^{-fx} \\
\end{bmatrix}
\]
\[ T_2(f, h) = \frac{1}{x^3} \left[ \frac{1}{h} e^{-f h x} - \frac{h - 2}{h(h - 1)^2} e^{-f h x} - \frac{f x}{h - 1} e^{-f h x} \right] \]
\[ 1 - e^{-f h x} - f x e^{-f h x} - \frac{f^2 x^2}{2} e^{-f h x} \]
\[ 2 e^{-f h x} + f x e^{-f h x} + f x - 2 \]

All functions of the angle variable \( h \) are defined in the following way:
\[
\ldots (h) = \begin{cases} 
  h \neq 0,1 & h = 1 
  \end{cases} \]
\[ h = 0 \]

Some of these functions were previously defined in [4,11,36]. Separate elements of the presented diagrammatic technique were used in [26,37–39].

**APPENDIX B: CALCULATION OF DIAGRAMS**

\[ S_{ij} = s'_a s'_b s = \quad \sum_{\gamma} \left( \begin{array}{c} 
  s' \\
  s \\
  N\Delta \\
\end{array} \right) \]

\[ I) = \frac{2}{N} \int_0^{L_\alpha} ds \int_0^s ds' P(s - s') = 2ND_f_a(1) \]

\[ II) = \frac{1}{N} \int_0^{L_\alpha} ds \int_0^{L_\beta} ds' P(s + s' + N\Delta) = NF_f_a F_f_\beta e^{-x\Delta} \]

\[ G_{ijk} = s''_a s'_b s = \quad \sum_{\gamma} \left( \begin{array}{c} 
  s'' \\
  s' \\
  s \\
\end{array} \right) \]

\[ I) = \frac{1}{N} \sum_{a \neq b \neq c=1}^3 \int_0^{L_\alpha} ds \int_0^s ds' \int_0^{s''} ds''' P \left( h_a^{(3)}(s - s') + h_c^{(3)}(s' - s'') \right) \]
\[ = 2N^2 \left[ 2g_2(f_a, h) + g_2(f_a, 1) \right] \]

\[ II) = \frac{1}{N} \int_0^{L_\alpha} ds \int_0^{L_\beta} ds' \int_0^{s'} ds'' \left[ P \left( h_c^{(3)}(s' - s'') + h_a^{(3)}(L_\beta - s' + N\Delta + s) \right) 
+ P \left( h_b^{(3)}(s' - s'') + h_a^{(3)}(L_\beta - s' + N\Delta + s) \right) \right] \]
\[ = N^2 e^{-h_a^{(3)}x\Delta} F_f_a(h_a^{(3)}) \left[ J_{f_\beta}(h) + J_{f_\beta}(h_a^{(3)}) \right] \]
\[ III = \frac{1}{N} \int_0^L ds \int_0^L ds' \int_0^L ds'' \mathcal{P} \left( h_c^{(3)} (s'' + N \Delta_1 + s') + h_a^{(3)} (L \beta - s' + N \Delta_2 + s) \right) \]
\[ = N^2 e^{-x_f h_c^{(3)}} e^{-x(h_c^{(3)} \Delta_1 + h_a^{(3)} \Delta_2)} \mathcal{F}_{f_a}(h_a^{(3)}) \mathcal{F}_{f_b}(h_b^{(3)}) - h_c^{(3)} \mathcal{F}_{f_a}(h_c^{(3)}) \]
\[ IV = \frac{1}{N} \int_0^L ds \int_0^L ds' \int_0^L ds'' \mathcal{P} \left( h_a^{(3)} (s + N \Delta_1) + h_b^{(3)} (s' + N \Delta_2) + h_c^{(3)} (s'' + N \Delta_3) \right) \]
\[ = N^2 \mathcal{F}_{f_a}(h_a^{(3)}) \mathcal{F}_{f_b}(h_b^{(3)}) \mathcal{F}_{f_c}(h_c^{(3)}) e^{-x(h_a^{(3)} \Delta_1 + h_b^{(3)} \Delta_2 + h_c^{(3)} \Delta_3)} \]

\[ G_{ijkl} = n_{\alpha} n_{\beta} n_{\gamma} n_{\delta} \]

\[ I = \frac{2}{N} \sum_{p \neq q=1}^{4} \int_0^L ds \int_0^L ds' \int_0^L ds'' \int_0^L ds''' \mathcal{P} \left( s''' - s'' + h_p^{(4)} (s' - s'') + s - s' \right) \]
\[ = 8N^3 \sum_{p=2}^{4} T_1(f, h_p^{(4)}) \]
$$II = \frac{2}{N} \sum_{p=1, p \neq a}^{4} \int_{0}^{L_{a}} ds \int_{0}^{L_{\beta}} ds' \int_{0}^{s'} ds'' \int_{0}^{s''} ds'''$$

$$\mathcal{P} \left( s'' - s''' + h_{ap}^{(4)}(s' - s'') + L_{\beta} - s' + N\Delta + s \right)$$

$$= 2N^{2} e^{-x\Delta} F_{fa} \sum_{p=1, p \neq a}^{4} T_{2}(f_{\beta}, h_{ap}^{(4)})$$

$$III = \frac{4}{N} \int_{0}^{L_{a}} ds \int_{0}^{s} ds' \int_{0}^{L_{\beta}} ds'' \int_{0}^{s''} ds'''$$

$$\mathcal{P} \left( s'' - s''' + h_{ap}^{(4)}(L_{\beta} - s'' + N\Delta + s') + s - s' \right)$$

$$= 4N^{2} e^{-hx\Delta} J_{fa}(h_{ab}^{(4)}) J_{f_{\beta}}(h_{ab}^{(4)})$$

$$IV = \frac{2}{N} \int_{0}^{L_{a}} ds \int_{0}^{s} ds' \int_{0}^{L_{\beta}} ds'' \int_{0}^{s''} ds'''$$

$$\mathcal{P} \left( s'' + N\Delta_{1} + s'' + h_{ap}^{(4)}(L_{\beta} - s'' + N\Delta_{2} + s') + s - s' \right)$$

$$= 2N^{2} e^{-x(\Delta_{1}+h_{ab}^{(4)}\Delta_{2})} F_{fa} e^{-x f_{\beta}} \mathcal{F}_{f_{\beta}}(h_{ab}^{(4)} - 1) J_{fa}(h_{ab}^{(4)})$$

$$V = \frac{1}{N} \sum_{p=1, p \neq a,b}^{4} \int_{0}^{L_{a}} ds \int_{0}^{L_{\beta}} ds' \int_{0}^{s'} ds'' \int_{0}^{s''} ds'''$$

$$\mathcal{P} \left( s'' + N\Delta_{1} + s'' + h_{ap}^{(4)}(s' - s'') + L_{\beta} - s' + N\Delta_{2} + s \right)$$

$$= N^{2} e^{-x(\Delta_{1}+\Delta_{2})} F_{fa} F_{f_{\gamma}} e^{-f_{\beta x}} \sum_{p=1, p \neq a,b}^{4} D_{f_{\beta}}(h_{ap}^{(4)} - 1)$$

$$VI = \frac{2}{N} \int_{0}^{L_{a}} ds \int_{0}^{L_{\beta}} ds' \int_{0}^{L_{\gamma}} ds'' \int_{0}^{s''} ds'''$$

$$\mathcal{P} \left( s + N\Delta_{1} + s'' + N\Delta_{2} + h_{ab}^{(4)}(N\Delta_{3} + s''') + s'' - s''' \right)$$

$$= 2N^{2} e^{-x(\Delta_{1}+\Delta_{2}+h_{ab}^{(4)}\Delta_{3})} F_{fa} F_{f_{\beta}} J_{f_{\gamma}}(h_{ab}^{(4)})$$

$$VII = \frac{1}{N} \int_{0}^{L_{a}} ds \int_{0}^{L_{\beta}} ds' \int_{0}^{L_{\gamma}} ds'' \int_{0}^{L_{\gamma}} ds'''$$

$$\mathcal{P} \left( s'' + N\Delta_{1} + s'' + h_{ab}^{(4)}(L_{\gamma} - s'' + N\Delta_{2} + s') + L_{\beta} - s' + N\Delta_{3} + s \right)$$

$$= N^{2} e^{-x(\Delta_{1}+h_{ab}^{(4)}\Delta_{2}+\Delta_{3})} F_{fa} \mathcal{F}_{f_{\beta}}(h_{ab}^{(4)} - 1) e^{-x(f_{\beta}+f_{\gamma})} \mathcal{F}_{f_{\gamma}}(h_{ab}^{(4)} - 1) F_{f_{\beta}}$$

$$VIII = \frac{1}{N} \int_{0}^{L_{a}} ds \int_{0}^{L_{\beta}} ds' \int_{0}^{L_{\gamma}} ds'' \int_{0}^{L_{\gamma}} ds'''$$

$$\mathcal{P} \left( s + N\Delta_{4} + s'' + N\Delta_{3} + s''' + N\Delta_{1} + s'' + h_{ab}^{(4)}(L_{\gamma} - s'' + N\Delta_{2}) \right)$$

$$= N^{2} e^{-x(\Delta_{1}+h_{ab}^{(4)}\Delta_{2}+\Delta_{3}+\Delta_{4})} F_{fa} F_{f_{\beta}} e^{-f_{\gamma x}} \mathcal{F}_{f_{\gamma}}(h_{ab}^{(4)} - 1) F_{f_{\gamma}}$$
\[ IX = \frac{1}{N} \int_0^{L_\alpha} ds \int_0^{L_3} ds' \int_0^{L_7} ds'' \int_0^{L_5} ds''' \]
\[ \mathcal{P} \left( s + N\Delta_1 + s' + N\Delta_3 + h_{ab}^{(4)} N\Delta_5 + s'' + N\Delta_2 + s''' + N\Delta_1 \right) \]
\[ = N^3 F_{f_a} F_{f_b} F_{f_7} F_{f_3} e^{-x(\Delta_1 + \Delta_2 + \Delta_3 + \Delta_4 + h_{ab}^{(4)} \Delta_5)} \]

**APPENDIX C: COEFFICIENTS \( \Xi \)**

We use the following notation: \( \tilde{\Gamma}_4 \left( \sum_{\alpha=1}^4 \Delta_\alpha, q_1, h_2 \right) = \tilde{\Gamma}_4 \left( \sum_{\alpha=1}^4 \Delta_\alpha, h^{(4)} \right) \).

**FCC:**

\[ \Xi^{(F)}_1 = 6\tilde{\Gamma}_4 \left( 0, 0, \frac{4}{3} \right) + \tilde{\Gamma}_4 \left( 0, 0, 0 \right) + 2\tilde{\Gamma}_4 \left( 0, \frac{4}{3}, \frac{4}{3} \right) \]
\[ \Xi^{(F)}_2 = 4\tilde{\Gamma}_4 \left( 4Q, 0, 2 \right) + \tilde{\Gamma}_4 \left( 4Q, 0, 0 \right) \]
\[ \Xi^{(F)}_3 = 3\tilde{\Gamma}_4 \left( 2Q, 0, z \right) + \tilde{\Gamma}_4 \left( 2Q, 2, z \right) + \tilde{\Gamma}_4 \left( 2Q, z, z \right) + \tilde{\Gamma}_4 \left( 2Q, \frac{8}{3}, z \right) \]
\[ z = 2 - \frac{2}{\sqrt{3}}, \quad Q = \frac{2}{\sqrt{3}} - 1 \]

**OBDD:**

\[ \Xi^{(O)}_1 = 2\tilde{\Gamma}_4 \left( 0, 0, 2 \right) + 8\tilde{\Gamma}_4 \left( 0, 0, 1 \right) + \tilde{\Gamma}_4 \left( 0, 0, 0 \right) + 4\tilde{\Gamma}_4 \left( 0, 1, 1 \right) \]
\[ \Xi^{(O)}_2 = \tilde{\Gamma}_4 \left( 4Q, 0, 0 \right) + 2\tilde{\Gamma}_4 \left( 4Q, \frac{4}{3}, \frac{4}{3} \right) + 6\tilde{\Gamma}_4 \left( 4Q, 0, \frac{4}{3} \right) \]
\[ \Xi^{(O)}_3 = 4\tilde{\Gamma}_4 \left( 2Q, 0, 2 \right) + 6\tilde{\Gamma}_4 \left( 2Q, 0, z \right) - 2\tilde{\Gamma}_4 \left( 2Q, 2, \frac{2}{3} \right) - 6\tilde{\Gamma}_4 \left( 2Q, 2, z \right) + \tilde{\Gamma}_4 \left( 2Q, z, z \right) \]
\[ + \tilde{\Gamma}_4 \left( 2Q, 4, z \right) + \tilde{\Gamma}_4 \left( 2Q, \frac{8}{3}, z \right) - 2\tilde{\Gamma}_4 \left( 2Q, \frac{4}{3}, z \right) \]
\[ z = 2 - 2\sqrt{\frac{2}{3}}, \quad Q = \sqrt{\frac{3}{2}} - 1 \]

**GYR:**

\[ \Xi^{(G)}_1 = \tilde{\Gamma}_4 \left( 0, 0, 0 \right) + 4\tilde{\Gamma}_4 \left( 0, 0, \frac{1}{3} \right) + 4\tilde{\Gamma}_4 \left( 0, 0, 1 \right) + 8\tilde{\Gamma}_4 \left( 0, 0, \frac{5}{3} \right) - 4\tilde{\Gamma}_4 \left( 0, \frac{1}{3}, \frac{2}{3} \right) \]
\[ - 4\tilde{\Gamma}_4 \left( 0, \frac{5}{3}, \frac{5}{3} \right) + 2\tilde{\Gamma}_4 \left( 0, 0, \frac{4}{3} \right) + 2\tilde{\Gamma}_4 \left( 0, \frac{2}{3}, \frac{2}{3} \right) + 4\tilde{\Gamma}_4 \left( 0, 0, \frac{2}{3} \right) \]
\[ \Xi^{(G)}_2 = 8\tilde{\Gamma}_4 \left( 4Q, 0, 1 \right) + 2\tilde{\Gamma}_4 \left( 4Q, 0, 2 \right) + 4\tilde{\Gamma}_4 \left( 4Q, 1, 1 \right) + \tilde{\Gamma}_4 \left( 4Q, 0, 0 \right) \]
$$
\Xi_3^{(G)} = 2\tilde{\Gamma}_4 \left( Q, 0, \frac{1}{3} \right) - 2\tilde{\Gamma}_4 \left( Q, 0, \frac{5}{3} \right) + \tilde{\Gamma}_4 \left( Q, \frac{1}{3}, \frac{1}{3} \right) + \tilde{\Gamma}_4 \left( Q, \frac{2}{3}, \frac{5}{3} \right) + 2\tilde{\Gamma}_4 \left( Q, 1, \frac{4}{3} \right) - 6\tilde{\Gamma}_4 \left( Q, 2, \frac{1}{3} \right) - 2\tilde{\Gamma}_4 \left( Q, 2, \frac{1}{\sqrt{3}} \right) - 2\tilde{\Gamma}_4 \left( Q, 2, \sqrt{3} \right) - 2\tilde{\Gamma}_4 \left( Q, \frac{1}{3}, \frac{4}{3} \right) + 2\tilde{\Gamma}_4 \left( Q, \frac{5}{3}, \frac{1}{3} \right) + \frac{2}{\sqrt{3}} - 2\tilde{\Gamma}_4 \left( Q, \frac{1}{3}, \frac{5}{3} + \frac{1}{3} \right) + 2\tilde{\Gamma}_4 \left( Q, 1, \frac{1}{3} + \frac{1}{\sqrt{3}} \right) + 2\tilde{\Gamma}_4 \left( Q, \frac{5}{3}, \frac{1}{3} + \frac{1}{\sqrt{3}} \right) - 2\tilde{\Gamma}_4 \left( Q, \frac{7}{3}, \frac{1}{\sqrt{3}} - \frac{1}{3} \right) + 2\tilde{\Gamma}_4 \left( Q, \sqrt{3}, 2 - \frac{1}{\sqrt{3}} \right) + \tilde{\Gamma}_4 \left( Q, \frac{2}{\sqrt{3}}, 2 - \frac{1}{\sqrt{3}} \right) - 2\tilde{\Gamma}_4 \left( Q, 2 - \sqrt{3}, \frac{4}{\sqrt{3}} \right) - 2\tilde{\Gamma}_4 \left( Q, \frac{1}{\sqrt{3}}, 2 - \sqrt{3} \right) + \tilde{\Gamma}_4 \left( Q, 2\sqrt{3}, 2 - \sqrt{3} \right)
$$

$$
\Xi_4^{(G)} = 6\tilde{\Gamma}_4 \left( 2Q, 0, 2 \right) - 2\tilde{\Gamma}_4 \left( 2Q, 2, \frac{1}{\sqrt{3}} \right) - 2\tilde{\Gamma}_4 \left( 2Q, 2, \sqrt{3} \right) - 2\tilde{\Gamma}_4 \left( 2Q, \frac{8}{3}, 2 - \frac{1}{\sqrt{3}} \right) - 2\tilde{\Gamma}_4 \left( 2Q, \frac{8}{3}, 2 - \sqrt{3} \right) - 2\tilde{\Gamma}_4 \left( 2Q, 1, \frac{1}{2} + \frac{1}{\sqrt{3}} \right) - 2\tilde{\Gamma}_4 \left( 2Q, \frac{4}{3}, 2 + \frac{1}{\sqrt{3}} \right) + 12\tilde{\Gamma}_4 \left( 2Q, 0, 2 + \frac{1}{\sqrt{3}} \right) + 6\tilde{\Gamma}_4 \left( 2Q, 0, 2 + \frac{2}{\sqrt{3}} \right) - 2\tilde{\Gamma}_4 \left( 2Q, 1, 1 + \sqrt{3} \right) + 2\tilde{\Gamma}_4 \left( 2Q, \frac{4}{3}, \frac{2}{3} + \sqrt{3} \right) + 2\tilde{\Gamma}_4 \left( 2Q, 3, \sqrt{3} - 1 \right) + 2\tilde{\Gamma}_4 \left( 2Q, 4, 2 - \sqrt{3} \right) - 4\tilde{\Gamma}_4 \left( 2Q, 2 - \sqrt{3}, \frac{4}{\sqrt{3}} \right) - 4\tilde{\Gamma}_4 \left( 2Q, 2 - \sqrt{3}, \frac{2}{\sqrt{3}} \right) - 2\tilde{\Gamma}_4 \left( 2Q, 2 - \sqrt{3}, 2\sqrt{3} \right) + 12\tilde{\Gamma}_4 \left( 2Q, 0, 2 - \sqrt{3} \right)
$$

$$Q = \frac{2}{\sqrt{3}} - 1$$
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FIG. 1. Model dendrimeric tree-like structure. The empty circle denotes the origin of the molecule. The filled circles denote the labeled blocks. a) Here we illustrate the introduced system of coordinates. The positions of the labels A, B and C are $R^{(A)} = \{1, 2, 2\}$, $R^{(B)} = \{2, 0, 0\}$ and $R^{(C)} = \{2, 3, 2\}$, respectively. b) The bold lines denote the pathways between the labeled blocks. The pathways are: $PW(A,B)=\{\{1,2,0\},\{1,3,0\}\}$ and $PW(C,D)=\{\{2,2,0\},\{2,0,0\},\{3,0,0\},\{3,2,0\}\}$. 

FIGURES
FIG. 2. Phase diagram for the $g = 2$, $n = \{2, 2\}$ and $\tau = \{1, 1\}$ dendrimer. The DIS-FCC line on the small graph does not touch the FCC-BCC line because of the finite step size.
FIG. 3. Phase diagram for the $g = 2$, $n = \{10, 4\}$ and $\tau = \{1, 1\}$ dendrimer.
FIG. 4. Phase diagram for the $g = 3$, $n = \{3,3,3\}$ and $\tau = \{1,1,1\}$ dendrimer.
FIG. 5. Phase diagram for the $g = 3$, $n = \{4, 4, 4\}$ and $\tau = \{1, 1, 1\}$ dendrimer. The cartoon of the dendrimer is not in scale: all lengths of the A-blocks should be the same.
FIG. 6. Phase diagram for the $g = 4$, $n = \{5, 1, 3, 1\}$ and $\tau = \{1, 1, 1\}$ dendrimer.
FIG. 7. Phase diagram for the $g = 5$, $n = \{1, 1, 1, 1, 1\}$ and $\tau = \{1, 1, 1, 1, 1\}$ dendrimer (pentablock).