

Stable Frank-Kasper phases of self-assembled, soft matter spheres

Abhiram Reddy^a, Michael B. Buckley^b, Akash Arora^c, Frank S. Bates^c, Kevin D. Dorfman^c, and Gregory M. Grason^{a,1}

^aDepartment of Polymer Science and Engineering, University of Massachusetts, Amherst, MA 01003; ^bDepartment of Physics, University of Massachusetts, Amherst, MA 01003; ^cDepartment of Chemical Engineering and Materials Science, University of Minnesota, Minneapolis, MN 55455

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Single molecular species can self-assemble into Frank Kasper (FK) phases, finite approximants of dodecagonal quasicrystals, defying intuitive notions that thermodynamic ground states are maximally symmetric. FK phases are speculated to emerge as the minimal-distortional packings of space-filling spherical domains, but a precise quantitation of this distortion and how it affects assembly thermodynamics remains ambiguous. We use two complementary approaches to demonstrate that the principles driving FK lattice formation in diblock copolymers emerge directly from the strong-stretching theory of spherical domains, in which minimal inter-block area competes with minimal stretching of space-filling chains. The relative stability of FK lattices is studied first using a diblock foam model with unconstrained particle volumes and shapes, which correctly predicts not only the equilibrium σ lattice, but also the unequal volumes of the equilibrium domains. We then provide a molecular interpretation for these results via self-consistent field theory, illuminating how molecular stiffness increases the sensitivity of the intra-domain chain configurations and the asymmetry of local domain packing. These findings shed new light on the role of volume exchange on the formation of distinct FK phases in copolymers, and suggest a paradigm for formation of FK phases in soft matter systems in which unequal domain volumes are selected by the thermodynamic competition between distinct measures of shape asymmetry.

self-assembly | Frank-Kasper phases | optimal lattices | block copolymers

Spherical assemblies occur in nearly every class of supramolecular soft matter, from lyotropic liquid crystals and surfactants, to amphiphilic copolymers (1). In concentrated or neat systems, self-assembled spherical domains behave as giant “mesoatoms,” adopting periodically-ordered crystalline arrangements. While superficially similar to lattices formed in atomic or colloidal systems – which are stabilized largely by bonding or translational entropy – the periodic order in soft materials is governed by distinctly different principles because lattice formation occurs in thermodynamic equilibrium with the formation of the “mesoatoms” from the constituent molecules themselves. Thus, the equilibrium sizes and shapes of “mesoatoms” are inextricably coupled to the lattice symmetry, and vice versa.

In this article, we address the emergence of non-canonical, Frank Kasper (FK) lattices in soft materials, characterized by complex and large-unit cells yet formed by assembly of a single molecular component. Initially constructed as models of metallic alloys (2, 3), FK lattices are a family of periodic packings (4, 5) whose sites are tetrahedrally-close packed (i.e. sitting on the vertices of nearly equilateral tetrahedra, the densest local arrangement of equal radius spheres) and can be decomposed into polyhedral (e.g. Voronoi or Wigner Seitz) cells surrounding each site containing 12, 14, 15 or 16 faces.

Known as the FK polyhedra, these cells (Z12, Z14, Z15 and Z16) possess variable volume and envelope spheres of distinct radii. Hence, FK lattices are natural candidates to describe ordered, locally-dense packings of spherical elements of different radii such as atomic alloys (3, 5) or binary nanoparticle superlattices (6). Once considered anomalous in soft matter systems, the past decade has seen an explosion in the observation of FK lattices in a diverse range of sphere-forming assemblies. These include (A15, σ) liquid-crystalline dendrimers (7, 8), linear (σ , A15) tetrablock (9, 10), (σ , A15, C14, C15) diblock (11–13) and (A15) linear-dendron (14) block copolymer melts, (A15) amphiphilic nanotetrahedra (15, 16), (A15, σ , C14, C15) concentrated ionic surfactants (17, 18) and (C14) monodisperse, functionalized nanoparticles (19). The central puzzle surrounding the formation of FK lattices in these diverse systems is understanding why single-components assemble into phases composed of highly heterogeneous molecular environments.

A common element distinct to FK formation in soft systems is the thermodynamic cost of *asphericity* imposed by incompatibility between uniform density and packing of perfectly spherical objects (Fig. 1). In soft assemblies, the ideally spherically symmetric domains are warped into lower-symmetry, polyhedral shapes which fill space without gaps. Intuitively, one expects that the minimal free energy state is the one for which the quasi-spherical domains (qSD) remain “most spherical.” The most commonly invoked notion of *sphericity* in this context is the dimensionless cell area A to volume V ratio,

Significance Statement

Formation of complex Frank-Kasper phases in soft matter systems confounds intuitive notions that equilibrium states achieve maximal symmetry, owing to an unavoidable conflict between shape and volume asymmetry in space-filling packings of spherical domains. Here we show the structure and thermodynamics of these complex phases can be understood from the generalization of two classic problems in discrete geometry, the Kelvin and Quantizer problems. We find that self-organized asymmetry of Frank-Kasper phases in diblock copolymers emerges from the optimal relaxation of cellular domains to unequal volumes in order to simultaneously minimize area and maximize compactness of cells, highlighting an important connection between crystal structures in condensed matter and optimal lattices in discrete geometry.

A.R., M.B.B., A.A., F.S.B., K.D.D. and G.M.G. designed research; M.B.B. developed computational method (foam model); A.R., M.B.B. and A.A. performed research; F.S.B., K.D.D. and G.M.G. supervised research; A.R., A.A. and G.M.G. analyzed computational data; and A.R., M.B.B., F.S.B., K.D.D. and G.M.G. wrote the paper.

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¹To whom correspondence should be addressed. E-mail: grason@mail.pse.umass.edu

125 $\mathcal{A} \equiv A/(36\pi V^2)^{1/3}$, which has a lower bound of 1 achieved by
 126 perfect spheres. The cellular partitions of FK lattices play a
 127 key role in the mathematical modeling of dry foams, known
 128 as the *Kelvin problem* (20–23), which seeks minimal area of
 129 partitions of space into equal volume cells. * Based on the fact
 130 that the lowest-area, equal-volume cellular partition known
 131 to date, the Weaire-Phelan foam (20), derives from the FK
 132 lattice A15, Zihlerl and Kamien proposed that this lattice is
 133 generically favored thermodynamically in so-called “fuzzy col-
 134 loid” models (24, 25), an argument subsequently adapted to
 135 sphere phases of block copolymers (26, 27). Recently, Lee,
 136 Leighton and Bates reasoned that average “sphericity” could
 137 be increased (i.e. decreased mean \mathcal{A}) below the Weaire-Phelan
 138 structure if the equal-volume constraint for distinct cells is re-
 139 laxed, as would occur for molecular exchange between distinct
 140 qSD, presumably, in equilibrium with the thermodynamic
 141 cost for deviations from a preferred aggregation number (11).
 142 Based on the Voronoi partitions, which have unequal volumes
 143 for FK lattices, σ was argued to have lower mean dimensionless
 144 area than A15, and thus should be stable over that lattice
 145 according to the sphericity argument, consistent with obser-
 146 vations of a σ lattice in diblock copolymer melts (13) and
 147 self-consistent field theory of conformationally and architec-
 148 turally asymmetric diblocks (28).

149 While the role of volume asymmetry has been implicated
 150 previously in the formation FK lattices by soft qSD assem-
 151 blies (29), critical questions remain unanswered. First, what
 152 are the relevant measures of sphericity optimized by the as-
 153 sembly thermodynamics? Second, how do these in concert
 154 determine the optimal balance between shape asymmetry
 155 (non-spherical domains) and volume asymmetry (molecular
 156 partitioning among domains) for a given qSD lattice? Finally,
 157 how does this balance select the equilibrium lattice and de-
 158 termine the scale of thermodynamic separation between the
 159 many competing FK lattices? We address these questions in
 160 the context of what we call the *diblock foam model* (DFM),
 161 which describes the thermodynamic competition between in-
 162 terdomain surface energy and chain stretching. For optimal
 163 mean qSD size, the DFM quantifies the thermodynamic cost
 164 of asphericity in terms of a geometric mean of reduced cell
 165 area and dimensionless radius of gyration of the cells, and
 166 thus, integrates elements of both the *Kelvin* and *lattice Quan-*
 167 *tizer* problems (30). These geometric proxies for inter-block
 168 repulsion and intra-molecular stretching in qSD exhibit qual-
 169 itatively different dependencies on cell shape, a factor that
 170 we show, based on this model and self-consistent field theory
 171 (SCFT) analysis, to be critical to the volume partitioning
 172 among distinct qSD and optimal lattice selection.

173 Amongst the various classes of FK-forming soft matter (7–
 174 9, 11–15, 17, 19), we posit that diblock copolymers represent
 175 the optimal starting point for investigating the selection of low
 176 symmetry FK phases by soft matter spheres. Diblock copoly-
 177 mers are a relatively simple chemical system, consisting of two
 178 flexible chains bonded together at their ends, and there exist
 179 robust theoretical methods for studying their phase behavior
 180 in the context of universal physical models (31, 32). The funda-
 181 mental mechanisms underlying assembly of diblock copolymers
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183 *The tetrahedral coordination of FK lattice implies that their partitions closely approximate the geo-
 184 metric constraints of Plateau borders, and are therefore near to minimal-area partitions. In ad-
 185 dition to A15, at least two more partitions of FK lattices, σ and H, have also been shown previ-
 186 ously (21, 23) to beat the area of optimum originally conjectured by Kelvin, the BCC partition. In
 Supplemental Section 2B, we report that the FK lattice P also belongs to this rarified category.

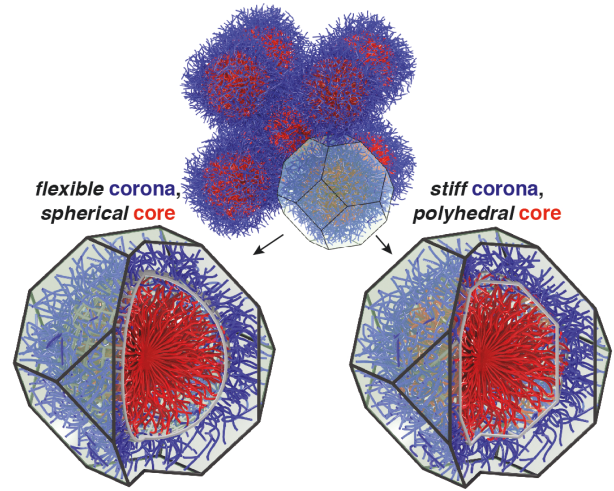


Fig. 1. Chain packing of spherical diblock copolymer domains of the BCC lattice (top), with corresponding limits of weakly-coupled (bottom left) and strongly-coupled (bottom right) of core domain shape of polyhedral (truncated-octahedron) cell symmetry.

209 that we elucidate here furnish the foundation for subsequent
 210 investigations of other soft matter systems, where these basic
 211 principles are conflated with additional phenomena emerging
 212 from electrostatics, hydrophobic interactions, and detailed
 213 packing of the complicated (non-Gaussian) configurations of
 214 their constituents.

Diblock Foam Model of FK Lattice Selection

215 We adopt what we call the *diblock foam model* (DFM), first
 216 developed by Milner and Olmsted, in which the free energy
 217 of competing arrangements is reduced to purely geometric
 218 measures of the cellular volumes enclosing the qSD (33, 34).
 219 To a first approximation, these cells are the polyhedral Voronoi
 220 cells for a given point packing, whose faces represent coronal
 221 brushes flattened by contact with neighboring qSD coronae.
 222 The model is based on strong-stretching theory (SST) of di-
 223 block copolymer melts, in which inter-block repulsions drive
 224 separation into sharply divided core and coronal domains and
 225 the chains are well-extended. We also consider the case of large
 226 elastic asymmetry between core and coronal blocks, which it-
 227 self derives from asymmetry of the block architecture or the
 228 segment sizes. This corresponds to the polyhedral-interface
 229 limit (35), in which the core/coronal interface in each qSD
 230 adopts a perfect, affinely shrunk copy of the cell shape (see
 231 Fig. 1, bottom right). Polyhedral warping of the interface
 232 is favored when the stiffness of the coronal blocks, which fa-
 233 vors a more uniform extension from the interface to the outer
 234 cell wall, dominates over entropic stiffness of core blocks and
 235 inter-block surface energy, which both favor round interfaces.

236 In this limit, the free energy per chain (26, 34), $F(X)$, of a
 237 given lattice packing X derives from two contributions,
 238

$$F(X) = \gamma \frac{\mathcal{A}(X)}{R_0} + \frac{\kappa}{2} \mathcal{I}(X) R_0^2, \quad [1]$$

239 where γ and κ are coefficients fixed by the chain properties
 240 (i.e., block lengths, segment lengths, inter-block repulsion),
 241 and R_0 is the radius of a sphere of equal volume to the mean
 242 volume of cells, or $(4\pi/3)R_0^3 = n_X^{-1} \sum_{i=1}^{n_X} V_i$, where V_i is i^{th}
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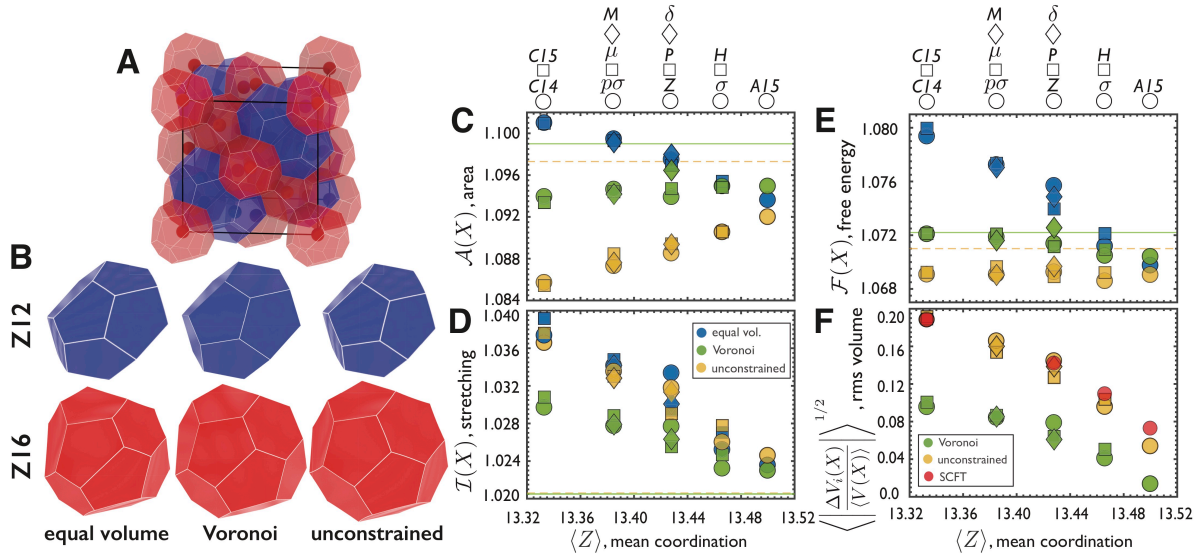


Fig. 2. A DFM structure for the cubic repeat of C15 is shown in (A), qSD centers shown within the Z12 and Z16 cells, red and blue, respectively. In (B), equilibrium shapes for three cell geometries studied, in which slight curvature of cell faces and edges is visible for the relaxed shape cases. Results of the DFM are shown for 11 competing FK phases (labeled above), plotted as function of mean coordination, or average number of cell faces $\langle Z \rangle$: mean dimensionless area (C); mean dimensionless stretching (D); mean free energy (E); and rms volume variation among cells relative mean volume $\langle V(X) \rangle$ (F). In (C)-(E) points are labeled according to the legend in (D) and the dashed and solid lines shows unconstrained and Voronoi results respectively for BCC. In (F), variable volume cell results are compared to qSD volumes extracted from SCFT at $\chi N = 40$, $f = 0.25$ and $\epsilon = 2$ as described in the text.

cell volume of n_X total cells in X (see Supplemental Sec. 1A for details). The first term represents the enthalpy of core-corona contact, and hence is proportional to the (per volume) interfacial area, which itself is proportional to the cell area A_i , measured by the dimensionless (mean) cell area, $\mathcal{A}(X) = (n_X^{-1} \sum_{i=1}^{n_X} A_i) / (4\pi R_0^2)$. The second term represents the entropic costs of extending polymeric blocks (here modeled as Gaussian chains) in radial trajectories within qSD. This cost grows with the square of domain size and depends on qSD shape through the dimensionless square radius of gyration, or stretching moment $\mathcal{I}(X) = (n_X^{-1} \sum_{i=1}^{n_X} I_i) / (4\pi R_0^5/5)$ where $I_i = \int_{V_i} d^3\mathbf{x} |\mathbf{x} - \mathbf{x}_i|^2$ is the second-moment volume of the i^{th} cell, whose center lies at \mathbf{x}_i . Optimizing mean cell size (R_0) yields the minimal free energy of lattice X , relative to the perfect sphere free energy $F_0 = \frac{3}{2}(\gamma^2 \kappa)^{1/3}$,

$$\mathcal{F}(X) \equiv \min_{R_0} [F(X)] / F_0 = [\mathcal{A}^2(X) \mathcal{I}(X)]^{1/3}. \quad [2]$$

This geometric mean favors simultaneously low values of dimensionless area and stretching[†]. While minimal area partitions (at constant volume) are associated Kelvin's foam problem, lattice partitions that optimize I_i (at fixed density) are the object of the *Quantizer problem* (30), which has applications in computer science and signal processing (36).

The Milner and Olmsted model has been studied for flat-faced Voronoi cells of FCC, BCC and A15 (27, 34), showing that the latter FK lattice of sphere-forming diblocks is favored over the former two canonical packings in the polyhedral interface limit. Here, we analyze a vastly expanded class of 11 FK lattices, possessing up to 56 qSD per periodic repeat. Most critically, we employ a Surface Evolver (37) based approach that minimizes $\mathcal{F}(X)$ over arbitrary volumes and shapes of constituent cells in the DFM structure (see Supplementary Sec. 2 for detailed method and tabulated results).

[†] Assembly thermodynamics depends on the dimensionless ratios of structure-averaged area and stretching of cells and volume, as opposed to averages of dimensionless cell area and stretching.

To assess the importance of relaxing volume and shape, consider the three distinct ensembles of qSD cells, shown for C15 in Fig. 2A,B. We have computed results for *equal-volume*, *relaxed-shape* cells, which cannot exchange mass, and *centroidal Voronoi cells*, which have fixed flat-face shapes but unequal volumes (also fixed). The former ensemble neglects the possibility of mass exchange between micelles, while the second optimizes stretching (36) but is suboptimal in terms of cell area[‡]. Neither model is realistic but they provide useful points of comparison to the unconstrained, *relaxed-volume and shape* cells, which strictly minimize $\mathcal{F}(X)$ for given X . Fig. 2C shows that allowing both volume and shape to relax leads to a complete inversion of the trend of $\mathcal{A}(X)$ with $\langle Z \rangle$. Importantly, there is also a near degeneracy for the free energy of FK structures in Fig. 2E, which all lie within 0.08% in $\mathcal{F}(X)$ (as compared to the relatively large $\approx 1\%$ spread for equal-volume qSD). These results confirm the critical role of volume exchange among asymmetric qSD in the thermodynamics of lattice formation (11, 12). Among these nearly degenerate, fully unconstrained DFM structures, the σ phase overtakes A15 (minimal for fixed, equal volume) as the minimal energy phase (with next lowest energy for P), consistent with its observation upon annealing (11, 13) as well as recent SCFT studies of conformationally asymmetric diblocks (28). Notably, however, in the *relaxed-volume and shape* DFM, σ possesses neither the minimal area (C14), nor minimal stretching (BCC). Rather, its predicted stability results from the optimal compromise between these competing measures of domain asphericity.

The interplay between area and stretching underlies the emergent asymmetry in equilibrium qSD volumes. Comparing the equal-volume to unconstrained DFM results in Fig. 2C and D shows that volume relaxation has a far more significant

[‡] Centroidal Voronoi cells have generating points at the centers of volume of the cell, and hence, for a give X minimize the mean-square distance of all points to their corresponding central point (see Supplemental Sec. 2A for additional details)

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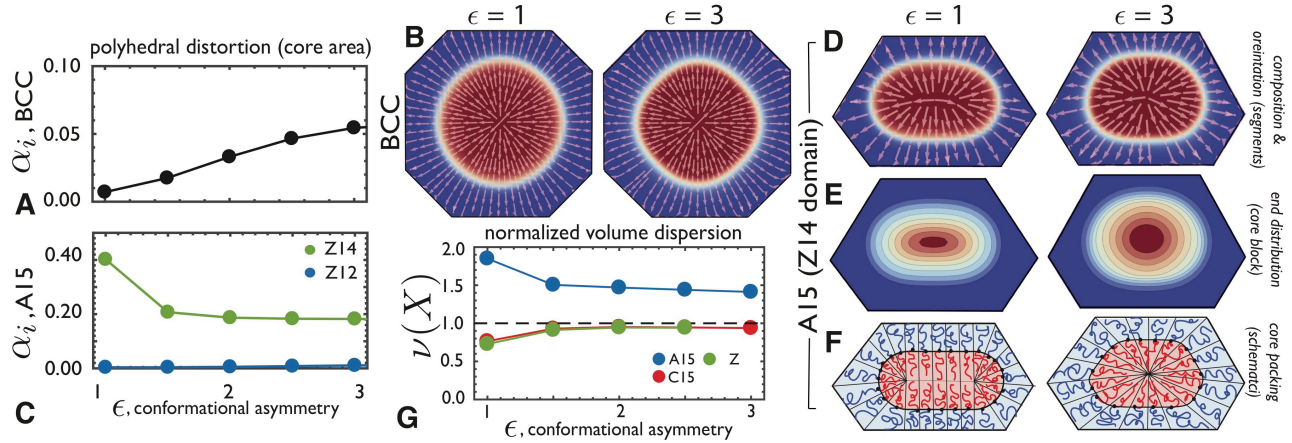


Fig. 3. The polyhedral warping of the A/B interface, measured by α_i from SCFT profiles of $\chi N = 40$ and $f = 0.29$ diblocks, of BCC qSD is plotted vs. conformational asymmetry $\epsilon = a_A/a_B$ in (A). Corresponding 2D cross sections (normal to $[100]$ through center of primitive cell) of qSD within the truncated-octahedral cells of BCC are shown in (B), with composition varying from red in A-rich regions to blue in B-rich regions (A/B interface is white). Also shown in vectors are the mean orientation of A-block segments (polar order parameter) (38). In (C), the areal distortion of Z14 and Z12 qSD from SCFT predictions of A15 are shown (same composition and segregation strength as BCC), with corresponding section of the Z14 (cut normal to $[100]$ through face of primitive cell, see Supplemental Fig. 16) qSD shown in (D) as in (B). Additionally, spatial distribution of the A-block (core forming) chain ends are shown in (E), varying from zero density (blue) to maximal density (red) within the cores. Schematics illustrating respective *discoidal* and *polyhedral* qSD packing are shown in (F). In (G), the volume dispersion (normalized by the DFM prediction) is plotted vs. conformational asymmetry.

effect on relaxation of $\mathcal{A}(X)$ than $\mathcal{I}(X)$, which changes little by comparison. Relaxation proceeds for all structures by inflating cells with relatively larger area, and shrinking smaller-area cells, restrained only by stretching cost creating highly unequal domain sizes (Supplemental Fig. 1). Volume exchange for lattices with large proportions of lower area Z12 cells (e.g., C14 and C15) achieve relatively large ($\approx 2\%$) drops in $\mathcal{A}(X)$ when compared to the high- $\langle Z \rangle$ end of the spectrum (e.g., $\approx 0.2\%$ for A15).

Cell volume asymmetry in equilibrated DFM structures pushes well beyond that of the “natural” geometry of Voronoi cells, which is strictly optimal for stretching, but not for its product with the square of dimensionless area. Fig. 2F shows that both unconstrained and Voronoi models of qSD cell geometry exhibit an increase volume dispersion with decreasing mean coordination (or, with increasing fraction of Z12s). However, optimal unconstrained DFM cells are nearly twice as polydisperse in volume as the Voronoi distribution. This massive volume asymmetry among qSD (up to $\approx 19\%$ variance for C14 and C15) is driven by dramatic reduction in inter-block contact area, a drive that is ultimately limited by the thermodynamic balance with the entropic (stretching) costs of filling space with qSD of unequal size. These results imply that structures with a larger equilibrium volume dispersion (such as the lower- $\langle Z \rangle$ C14 and C15) structures are more susceptible to the effects of thermal processing that selectively promote or inhibit chain exchange among equilibrating spheres (12) than phases such as A15, which relax free energy relatively little through volume equilibration.

Previous SCFT studies (12, 28) have shown that the canonical BCC sphere phase is overtaken by a stable σ lattice when the elastic asymmetry, embodied by ratios of statistical segment lengths $\epsilon \equiv a_A/a_B \gtrsim 1.5$. DFM not only correctly predicts σ as the dominantly stable sphere phase, but also does a remarkable job of predicting the relative hierarchy among metastable FK competitors. This is evident in Supplemental Fig. 16A-C, where we compare the free energies, scaled enthalpies and entropies for σ , Z, C14, C15 and A15

predicted by the unconstrained cell DFM to AB diblock SCFT calculations using methods described in ref. (32) at somewhat strong segregation conditions $\chi N = 40$, where χ is the Flory-Huggins parameter for A/B contact and N is the degree of polymerization. DFM correctly predicts the narrow 0.01% scale of free-energy splitting between these competitors for $\epsilon = 2$ diblocks in the composition range $f \leq 0.25$, where f is the volume fraction of the minority block. Moreover, DFM predicts their ranking relative to σ with the exception of Z, which DFM predicts to be nearly degenerate with C15. The accuracy of DFM extends beyond thermodynamics to structure, most notably the volume asymmetry in Fig. 2F.

Molecular Mechanism of Aspherical Domain Formation

To probe the molecular mechanism that underlies the selection of FK lattices in block copolymers, we analyze two order parameters that quantify the respective asymmetric shapes and volumes of qSD, computed from the volumes enclosing A-rich cores in SCFT composition profiles of diblocks at $\chi N = 40$, $f = 0.29$ and for variable conformational asymmetry (see Supplementary Section 3B). The first parameter,

$$\alpha_i = \frac{\mathcal{A}_i^{A/B} - 1}{\mathcal{A}^{\text{poly}} - 1}, \quad [3]$$

measures the degree of polyhedral warping of the core in terms of the dimensionless area $\mathcal{A}_i^{A/B}$ of the A/B interface of the i^{th} domain relative to a sphere, where $\mathcal{A}_i^{\text{poly}}$ is the dimensionless area predicted for the perfectly polyhedral interface of the corresponding cell from the unconstrained DFM: $\alpha_i = 0$ for spherical interfaces; and $\alpha_i = 1$ for interfaces that adopt the polyhedral shapes of the DFM cells. We define a second parameter, $\nu(X)$, that measures asymmetry of unequal volumes enclosed within A/B interfaces predicted by SCFT, relative to the volume asymmetry predicted by polyhedral cells of DFM for the same structure X

$$\nu(X) = \frac{\left\langle \left| \frac{\Delta V_i(X)}{\langle V(X) \rangle} \right|^2 \right\rangle_{A/B}^{1/2}}{\left\langle \left| \frac{\Delta V_i(X)}{\langle V(X) \rangle} \right|^2 \right\rangle_{\text{poly}}^{1/2}} \quad [4]$$

497 where $\Delta V_i(X) = V_i - \langle V(X) \rangle$ is the volume deviation of the
 498 i^{th} domain relative to the average in X , and values of $\nu(X)$
 499 greater (less) than 1 indicate that qSD in SCFT are more
 500 (less) polydisperse predicted by relaxed DFM cells.

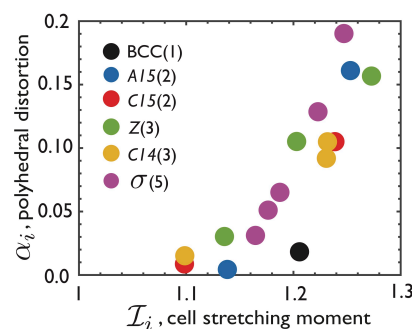
501 It has been argued previously (27) that the polyhedral warp-
 502 ing, or faceting, of core-corona interfaces should increase with
 503 ϵ , which controls the ratio of corona- to core-block stiffness,
 504 due to the relatively lower entropic cost of more uniformly
 505 stretched coronae achieved by polyhedral interfaces. This ex-
 506 pectation is consistent with the observed monotonic increase
 507 of α from 0 at $\epsilon = 1$ to the saturated value of $\alpha \approx 0.05$ for
 508 $\epsilon \gtrsim 2 - 3$ for the qSD in BCC plotted in Fig. 3A[§]. As shown
 509 in Supplemental Fig. 17, the polyhedral warp of the interface
 510 grows also with increasing f , due to the increased proxim-
 511 ity of the qSD cell boundary to the interface and relatively
 512 shorter coronal blocks at larger core fractions. While clearly
 513 far from a sharply faceted shape, the increase in core shape
 514 anisotropy is obvious from 2D cuts through of the qSD shown
 515 in Fig. 3B, showing a visible warp of A/B interface towards
 516 the truncated-octahedral shape of the BCC cell at $\epsilon = 3$.

517 For the FK phases, which are composed of distinct-
 518 symmetry qSD, areal distortion exhibits a markedly different
 519 dependence on increased coronal/core stiffness, as illustrated
 520 by the plots of α_{12} and α_{14} vs. ϵ for A15 in Fig. 3C. Z12
 521 domains exhibit a monotonic, albeit modest, increase in dis-
 522 tortion with ϵ . Surprisingly, for the Z14 domains, the excess
 523 area drops from its maximal value of $\alpha_{14} \simeq 0.4$ in the con-
 524 formationally symmetric case for $\epsilon = 1$ down to a lower, yet
 525 significant plateau value of $\alpha_{14} \simeq 0.2$, roughly twice the areal
 526 distortion for BCC.

527 The origin of this counterintuitive drop in dimensionless
 528 area of the Z14 cells with increased outer block stiffness is
 529 illustrated in Fig. 3D, which compares 2D sections of the
 530 Z14 qSD of A15 at $\epsilon = 1$ and $\epsilon = 3$. While the shape for
 531 larger outer-block stiffness ($\epsilon = 3$) is consistent with a quasi-
 532 faceted interface that copies the polyhedral cell (with rounded
 533 edges) of the Z14 domain, the conformationally symmetric
 534 case ($\epsilon = 1$) is neither faceted nor spherical. It instead adopts
 535 oblate, or *discoidal* shape. The contrast in core shape is fur-
 536 ther reflected in the sub-interface (vector) orientational order
 537 parameter of A-segments (38) and the spatial distribution of
 538 A-block chain ends, also shown in Fig. 3D,E. For larger ϵ , the
 539 preference for more uniform coronal block stretching drives the
 540 quasi-polyhedral domain shape, with radial chain trajectories
 541 extending from the center of the domain, a point at which
 542 core block ends are concentrated. In contrast, for the case of
 543 $\epsilon = 1$, the stiffness of the core blocks is sufficient to resist
 544 deformations away from uniform core thickness. Occupying
 545 the somewhat flattened Z14 cell with a qSD of uniform core
 546 thickness then leads to the discoidal shape, in which chain
 547 ends spread laterally in a quasi-lamellar core rimmed by a
 548 quasi-toroidal packing at its circumference. The preference for
 549 uniform core thickness within the relatively oblate Z14 cell,
 550 which gives rise to a larger area discoidal interface for $\epsilon = 1$,
 551 ultimately gives way to the quasi-polyhedral qSD shape, and
 552 corresponding radial chain stretching, with increased outer
 553 block stiffness for $\epsilon \gtrsim 2$ (see schematic in Fig. 3F).

554 Supplemental Fig.18 shows evidence of this same *discoidal*
 555 \rightarrow *polyhedral* transition qSD within the most oblate cells of

557 [§]While this extends beyond what is realized with most flexible linear diblocks, bulky side chains
 558 including bottlebrush configurations and miktoarm polymers would make the upper limit accessible.



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Fig. 4. Correlation between polyhedral warping of core shapes (α_i) within symmetry-distinct qSD extract from SCFT at $\chi N = 40$, $f = 0.25$ and $\epsilon = 2$ and the degree of frustration of chain stretching in the correspond cell, quantified by the (cell-wise) dimensionless stretching moment, \mathcal{I}_i .

other FK phases, C15 and Z, leading to a corresponding drop
 in excess area α_i from $\epsilon = 1$ to $\epsilon \approx 2$ for those cells. In
 Fig. 3G we find this intra-domain shape transition with in-
 creasing corona-/core-block stiffness is coupled to a transition
 in volume asymmetry among qSD. Discoidal domains of the
 conformationally symmetric diblocks ($\epsilon = 1$) realize a volume
 dispersion that is strongly divergent from the polyhedral geom-
 etry in the DFM, including both greater ($\nu(X) > 1$, for A15)
 and lesser ($\nu(X) < 1$ for Z, C15) dispersity. Yet, in the limit
 of $\epsilon \gtrsim 2$, relatively stiffer coronal blocks pull the cores into
 radial-stretching, quasi-polyhedral shapes. This transition to
 more compact cores, in turn, results into volume redistribut-
 ing among equilibrium qSD tending to the $\nu(X) \rightarrow 1$ limit,
 consistent with agreement between asymmetric volumes of
 DFM and SCFT shown in Fig. 2F.

Notwithstanding the broad agreement between SCFT and
 DFM predictions, the degree of polyhedral warping of qSD
 shape is both arguably modest (i.e., $\alpha \lesssim 0.3$ for $\epsilon \gg 1$ for
 this χN and f) and highly variable in the FK structures, sug-
 gesting a heterogeneous degree of shape frustration among
 cells. Moreover, the *discoidal* \rightarrow *polyhedral* transition occurs
 only in high- α qSD, whereas low- α cells (e.g., Z12 cells of
 A15) maintain radial stretching and a monotonic dependence
 on ϵ . What controls the variability of coupling between cell
 geometry and polyhedral distortion? Fig. 4 shows the correla-
 tion between α_i for qSD extracted from SCFT at $\chi N = 40$,
 $f = 0.25$ and $\epsilon = 2$ (i.e. in the quasi-polyhedral shape regime)
 plotted as a function the dimensionless stretching \mathcal{I}_i for the
 corresponding cells from the DFM. The generically increasing
 trend of α_i with \mathcal{I}_i for cell geometries across competing phases
 argues that the variable degree of shape frustration within
 distinct qSD, and its consequent impact on qSD core shape,
 is regulated by the constraints of asymmetric chain-stretching
 in polyhedral cells. In other words, the ultimate degree of as-
 phericity of core distortion of qSD (measured by dimensionless
 area), is in fact, controlled by the local asphericity in radial
 stretching required by space-filling chain packing (measured
 by dimensionless radius of gyration).

Concluding Remarks

We anticipate that the emergence of optimal FK lattice struc-
 ture and thermodynamics via a balance of competing measures
 of domain asymmetry highlighted here for high molecular
 weight diblock copolymers will extend to other copolymer

621 systems where these phases have been observed, including
 622 architecturally asymmetric copolymers, linear multiblocks,
 623 low molecular weight/high- χ systems and blends. In par-
 624 ticular, lower molecular weight polymers drive the system
 625 closer towards the strong segregation limit and away from
 626 the mean-field limit. Each of these materials exhibit different
 627 molecular mechanisms through which the relative stiffness of
 628 the coronal domain transmits the asymmetry of the local qSD
 629 packing into the core shape. For example, the observation
 630 of polygonal/polyhedral warping of outer zones of core-shell
 631 domains of linear multiblock polymers (39) provides a plau-
 632 sible mechanism to stabilize the σ phase observed in linear
 633 tetrablocks (9). On the other hand, accurately modeling the
 634 formation of σ by low molecular weight conformationally asym-
 635 metric diblocks (11, 13), likely requires a non-Gaussian (finite
 636 extensibility) model of chain stretching, but one which never-
 637 theless, like the dimensionless radius of gyration $\mathcal{I}(X)$, favors
 638 compact domains and competes against the minimal area prefer-
 639 ence for unequal domain volumes. Beyond copolymers, we
 640 speculate further that additional intra- and inter-molecular
 641 mechanisms play the role of balancing the drive for minimal
 642 domain area in the formation FK phases, from giant nanotetra-
 643 hedra (15, 16) to ionic surfactants (17, 18). We speculate that
 644 the DFM studied here may anticipate a much broader class of
 645 “generalized foam models” that integrate two or more measures
 646 of cellular shape, and which may be useful as minimal models
 647 for a wider range of tessellated architectures (e.g. living tissue)
 648 (41).

649 The present results for the DFM also shed new light on
 650 the non-equilibrium pathways for stabilizing metastable FK
 651 competitors, as has been demonstrated for conformational
 652 asymmetric linear diblocks quenched from high-temperature
 653 disorder sphere phases to low temperature metastable A15,
 654 C14 and C15 phases (12, 40). The low temperature quench
 655 is suspected to freeze out the inter-domain chain exchange
 656 needed to achieve the equilibrium σ state, thus the kinetically-
 657 trapped quenched state inherits the volume distribution of high
 658 temperature micelle liquid state. The DFM suggests a new
 659 way to analyze the stability of FK states when domain volumes
 660 are out of equilibrium, suggesting the observation of C14 and
 661 C15 may be selected among the low-temperature kinetically
 662 trapped arrangements because it inherits a volume distribution
 663 that is both smaller in average cell size and possibly more
 664 polydisperse than the equilibrium state at the low temperature,
 665 and hence, a better fit to the “aggregation fingerprint” of low-
 666 $\langle Z \rangle$ packings.

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Supporting Information: Stable Frank Kasper phases of self-assembled, soft matter spheres

Abhiram Reddy^a, Michael B. Buckley^b, Akash Arora^c, Frank S. Bates^c, Kevin D. Dorfman^c, and Gregory M. Gason^a

^aDepartment of Polymer Science and Engineering, University of Massachusetts, Amherst, 01003; ^bDepartment of Physics, University of Massachusetts, Amherst, 01003; ^cDepartment of Chemical Engineering and Materials Science, University of Minnesota, Minneapolis, MN 55455

1. Diblock Foam Model

A. Polyhedral Interface Limit of Strongly-Segregated Diblock Sphere Lattices. We briefly overview the strong-segregation theory (SST) calculation of Milner and Olmsted (1, 2) for spherical domains in *polyhedral interface limit* (PIL), also known as the *straight path ansatz*. We further show how the free energy of competing sphere packings is computed from purely geometric measures of the cellular volumes that enclose distinct spheres (3, 4), which forms the basis for the Diblock Foam Model (DFM).

Here, we focus on the case of AB linear diblocks with conformational asymmetry, but the theory can be generalized to other architectures like miktoarm stars (4). We consider a chain with total segment number $N = N_A + N_B$, with $f = N_A/N$ the fraction of the A-block. Segments are taken to have equal volumes ρ_0^{-1} and potentially unequal statistical segment lengths, a_A and a_B , for the respective blocks. The ratio of segment lengths defines the conformational asymmetry $\epsilon \equiv a_A/a_B$. Within SST, the total free energy $F(X)$ (in units of $k_B T$) of a periodic repeat spherical assembly of lattice packing X decomposes into two terms

$$F(X) = F_{int} + F_{st}, \quad [1]$$

which represents the respective costs of inter-block repulsions at a core/coronal interfaces and the entropic cost of stretching of (Gaussian) chains from random walk configurations.

The first term $F_{int} = \Sigma A_{int}$ simply derives from the product of total area of core/coronal contact, A_{int} , times Σ to give the surface area energy between phase separated A and B domains (5),

$$\Sigma = \rho_0 a \sqrt{\frac{\chi}{6}} \left(\frac{2}{3} \frac{\epsilon^{3/2} - \epsilon^{-3/2}}{\epsilon - \epsilon^{-1}} \right), \quad [2]$$

where χ is the Flory-Huggins parameter for AB repulsion and $a \equiv \sqrt{a_A a_B}$ is the geometric mean of segment lengths*. For the i th cell of X , the core/corona interface is an affinely shrunk copy of the outer cell that encloses a fraction f of the total cell volume. Hence, the area of the core interface of i th domain is $f^{2/3} A_i$, where A_i is the cell area, and $A_{int} = f^{2/3} \sum_{i=1}^{n_X} A_i$, where n_X is the number of domains (and cells) per periodic repeat.

The entropic contribution from chain stretching for domain α in cell i (denoted as volume $V_{\alpha,i}$) can be evaluated using the SST entropy derived from the ‘‘parabolic brush’’ theory (6), which can be expressed as (7),

$$F_{st,i}^{(\alpha)} = \frac{3\pi^2 \rho_0}{8N_A^2 a_A^2} \int_{V_{\alpha,i}} d^3 \mathbf{x} z^2 \quad [3]$$

where z is the distance from the AB interface, where junction points are localized, from which chain trajectories are assumed to extend along the radial lines extending from the cell center \mathbf{x}_i to the outer wall of cell i , and $\int_{V_{\alpha,i}} d^3 \mathbf{x}$ is the integral over volume. For spherical domains, these integrals are evaluated by describing the cell shape as a function of the radial directions $\hat{\Omega}$ extending from the cell center at \mathbf{x}_i : $R_i(\hat{\Omega})$ and $R'_i(\hat{\Omega})$ are the respective distances to the interface and outer wall of the cell in direction $\hat{\Omega}$. Because the core chains occupy a fixed fraction f of each ‘‘wedge’’ in the PIL, we have $R'_i(\hat{\Omega}) = f^{1/2} R_i(\hat{\Omega})$, and the stretching contributions from each block are proportional to the same geometric stretching moment,

$$F_{st,i}^{(A)} = \frac{3\pi^2 \rho_0}{8N_A^2 a_A^2} \int d^2 \hat{\Omega} \int_0^{R'_i(\hat{\Omega})} dr r^2 [R'_i(\hat{\Omega}) - r]^2 = \frac{\pi^2 \rho_0}{80 f^{1/3} N^2 a_A^2} S_i, \quad [4]$$

*Note that Σ is varies with conformational asymmetry ϵ , and that it reduces to the standard result for interfacial tension between immiscible polymer melts in the symmetric limit $\Sigma(\epsilon \rightarrow 1) = \rho_0 a (\chi/6)^{1/2}$.

125 and

$$126 \quad F_{st,i}^{(B)} = \frac{3\pi^2\rho_0}{8N_B^2 a_B^2} \int d^2\hat{\Omega} \int_{R'_i(\hat{\Omega})}^{R_i(\hat{\Omega})} dr r^2 [R'_i(\hat{\Omega}) - r]^2 = \frac{\pi^2\rho_0(1-f^{1/3})^3(6+3f^{1/3}+f^{2/3})}{80(1-f)^2 N^2 a_B^2} S_i, \quad [5] \quad 187-189$$

128 where

$$129 \quad S_i \equiv \int d^2\hat{\Omega} R_i^5(\hat{\Omega}), \quad [6] \quad 190-191$$

130 and is propotional to the second-moment of cell volume defined in the main text,

$$131 \quad I_i = \int_{V_i} d^3\mathbf{x} |\mathbf{x} - \mathbf{x}_i|^2 = \int d^2\hat{\Omega} \int_0^{R_i(\hat{\Omega})} dr r^4 = S_i/5. \quad [7] \quad 192-193$$

134 Combining these together and summing over the cells in the periodic repeat we have the total stretching free energy

$$135 \quad F_{st} = \frac{\pi^2\rho_0}{16N^2 a^2} \left[\frac{f^{1/3}}{\epsilon} + \frac{\epsilon(1-f^{1/3})^3(6+3f^{1/3}+f^{2/3})}{(1-f)^2} \right] \sum_{i=1}^{n_X} I_i. \quad [8] \quad 194-195$$

136 Since melt assembly occurs at fixed total density, equilibrium states correspond to states of minimal free energy
137 per chain. Defining the mean volume of the cells in X as $V_0 = n_X^{-1} \sum_{i=1}^{n_X} V_i$, the total number of chains per periodic
138 repeat is $n_X V_0 \rho_0 / N$. The mean volume per cell also defines a measure of the mean cell dimension $R_0 = (3V_0/4\pi)^{1/3}$,
139 the radius of a sphere of equal volume to V_0 . Using this definition we can rewrite the area per volume as

$$140 \quad \frac{\sum_{i=1}^{n_X} A_i}{n_X V_0} = \frac{3\mathcal{A}(X)}{R_0} \quad [9] \quad 200-201$$

141 and the stretching per volume as

$$142 \quad \frac{\sum_{i=1}^{n_X} I_i}{n_X V_0} = \frac{3}{5} \mathcal{I}(X) R_0^2 \quad [10] \quad 202-203$$

143 where the dimensionless quantities $\mathcal{A}(X) = n_X^{-1} \sum_{i=1}^{n_X} A_i / (4\pi R_0^2)$ and $\mathcal{I}(X) = n_X^{-1} \sum_{i=1}^{n_X} I_i / (4\pi R_0^5/5)$ depend only on
144 cell shapes and are independent of R_0 , or mean domain size. Using these quantities and dividing $F_{int} + F_{st}$ by the
145 total chain number we arrive at eq. [1] in main text, where the coefficients are given by,

$$146 \quad \gamma = Na\sqrt{2\chi/3} \left(\frac{\epsilon^{3/2} - \epsilon^{-3/2}}{\epsilon - \epsilon^{-1}} \right) f^{2/3}, \quad [11] \quad 204-205$$

147 and

$$148 \quad \kappa = \frac{3\pi^2}{40Na^2} \left[\frac{1}{\epsilon f^{1/3}} + \frac{\epsilon(1-f^{1/3})^3(6+3f^{1/3}+f^{2/3})}{(1-f)^2} \right], \quad [12] \quad 206-207$$

149 which are independent of structure X and are fixed for a given set chain properties. Optimizing $F(X)$ with respect to
150 R_0 , we find a equilibrium mean domain size

$$151 \quad (R_0)_{eq} = R_s \left(\frac{\mathcal{A}(X)}{\mathcal{I}(X)} \right)^{1/3}, \quad [13] \quad 208-209$$

152 where $R_s = (\gamma/\kappa)^{1/3} \propto (\chi N)^{1/6} N^{1/2} a$ is the thermodynamically selected radius of domains if cells were equal volume
153 spheres (i.e. $\mathcal{A} = \mathcal{I} = 1$).

154 We note that this model relies on the so-called parabolic brush theory (6) in the expressions for Gaussian chain
155 entropy in eq. (3), which are known to fail for brush-like domains with convex curvature due to the presence
156 end-exclusions zones missing from the parabolic model. Notwithstanding, the failure to properly account for these
157 exclusion zones in the coronal blocks of this calculation (8), this approximation only modifies the coefficient κ and its
158 dependence of f^\dagger . The proportionality of the stretching free energy with $\int d^2\hat{\Omega} R^5(\hat{\Omega})$ follows on the general grounds
159 that each ‘‘wedge’’ of the domain includes a number of chains proportional to $d^2\hat{\Omega} R^3(\hat{\Omega})$, each of which is stretched a
160 distance proportional to $R(\hat{\Omega})$ and hence acquires a free energy penalty proportional to $R^2(\hat{\Omega})$.

161 [†]Specifically, it can be shown that the coronal brush free energy in a spherical geometry is proportional to h^2 , where h is the brush thickness, times a function of h/R_s , where R_s is the spherical radius.
162 In this geometry $h(\hat{\Omega}) = R(\hat{\Omega}) - R'(\hat{\Omega}) = (1 - f^{1/3})R(\hat{\Omega})$ and $R_s = R'(\hat{\Omega})$ so that $h/R_s = (f^{-1/3} - 1)$ for each wedge, independent of $\hat{\Omega}$

B. Unequal Domain Volumes in A15 Lattices: Weighted Voronoi Partitions. Here we illustrate the dependence of the DFM energy $\mathcal{F}(X) = [\mathcal{A}^2(X)\mathcal{I}(X)]^{1/3}$ on the volume difference between symmetry-distinct cells of FK lattices. For this purpose we consider the A15 lattice, which can be decomposed into two Z12 cells (at the center and corners of the primitive, cubic cell) and six Z14 cells (two positions decorating each face of the primitive, cubic cell). An analytical relation for $\mathcal{F}(\text{A15})$ can be obtained using expressions for cellular area, volume and second-moments of volume for the weighted Voronoi cells of A15. Standard Voronoi partitions derive from the polyhedra constructed by planes that bisect the center-to-center neighbor separation vectors normally. Here, we use the weighted partitions of A15 derived by Kusner and Sullivan (17), correspond to the (flat-face) polyhedra constructed from planes at variable separation between the Z12 and Z14 sites (i.e., non-bisecting).

Fixing the length of the primitive cubic cell to 2, and the mean volume per cell is fixed to $V_0 = 1$ (non-dimensional lengths), the total area of the cells can be expressed as a function of c , which parameterizes the size of the dodecahedral Z12 cells: the volumes of these cells are $V_{Z12} = c^3/2$; which implies $V_{Z14} = (4 - V_{Z12})/3$. The mean cell area (17) is

$$\frac{1}{8} \sum_{i=1}^8 A_i(c) = \frac{3}{2} + \frac{3}{2}\sqrt{6} + \frac{1}{8}(6\sqrt{5} - 4\sqrt{6} - 3)c^2 \quad [14]$$

while the total second moments per cubic repeat was calculated by Kashyap and Neuhoff (11) as

$$\frac{1}{8} \sum_{i=1}^8 I_i(c) = \frac{1}{32}(3c^4 - 5c^3 + 10). \quad [15]$$

Normalizing these by the area and second moment of spheres of $V_0 = 1$ gives the dimensionless free energy of A15 for flat faced cells,

$$\mathcal{F}_c(\text{A15}) = \left(\frac{5}{55296}\right)^{1/3} \left([12 + 12\sqrt{6} + (6\sqrt{5} - 4\sqrt{6} - 3)c^2]^2(3c^4 - 5c^3 + 10)\right)^{1/3} \quad [16]$$

The dimensionless area, stretching and free energy are plotted as function of the volume difference

$$\frac{\Delta V}{V_0} = \frac{2}{3}(c^3 - 2), \quad [17]$$

where $\Delta V = V_{Z12} - V_{Z14}$ between Z12 and Z14 cells in Fig. 1. This shows that the dimensionless area is minimal for vanishing Z12 volume ($c = 0$), while dimensionless stretching is in fact minimized by the standard (centroidal) Voronoi partition ($c = 5/4$), which has a very nearly equal volumes $\Delta V/V_0 = -0.03$. The competition between these drives for unequal cell volumes which results in optimal free energy ($\mathcal{F}_c(\text{A15}) = 1.070$) for flat faced cells with $c_{min} = 1.22$ and $\Delta V/V_0 = -0.12$.

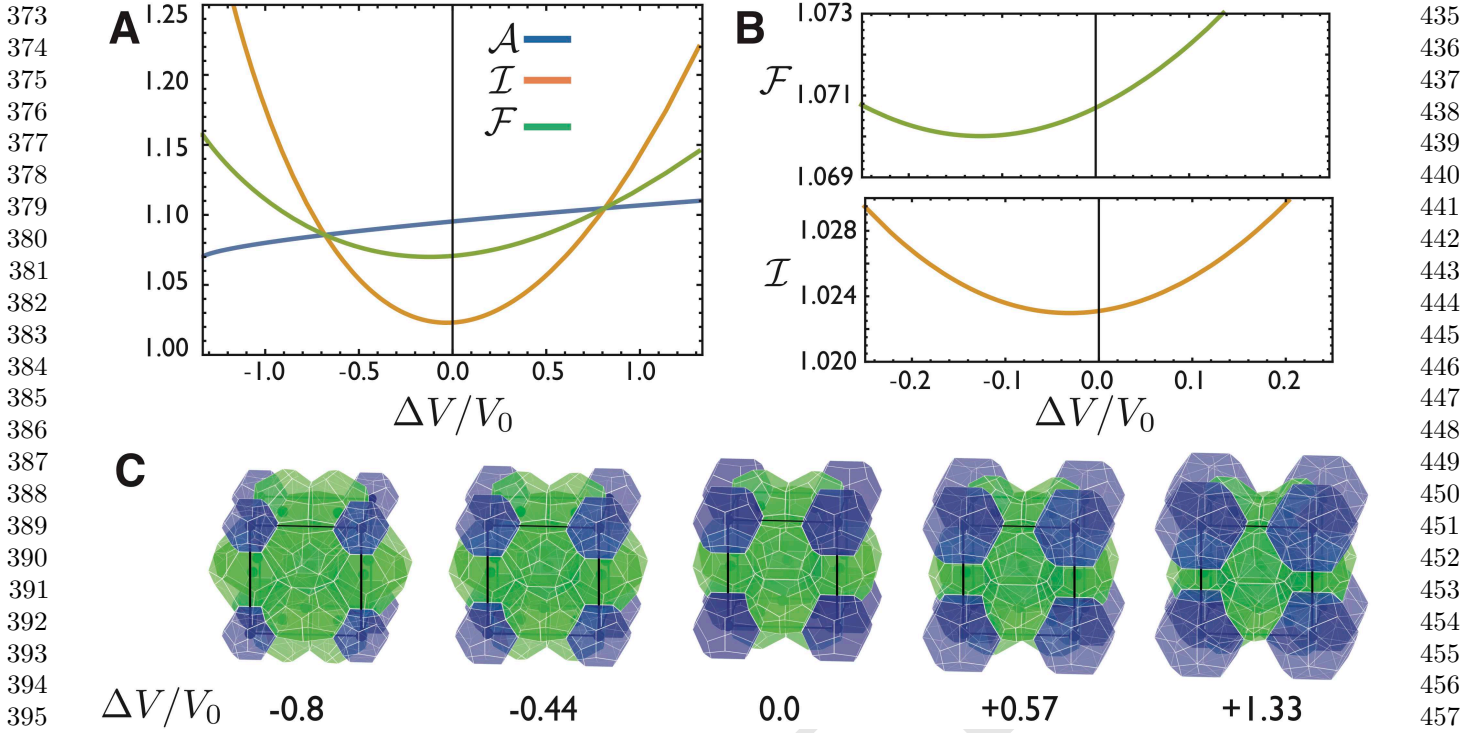


Fig. 1. (A) Plot of \mathcal{A} , \mathcal{I} and \mathcal{F} for A15 lattice as a function of varying volumes of Z12 and Z14 cells; in (B) zoom in of minimal for free energy and stretching. In (C), schematics of A15 primitive cell with unequal cell volumes: Z12 (blue) and Z14 (green) polyhedra.

2. Numerical optimization of cell geometry

Beyond the case of the flat-faced A15 cell results shown in Fig. 1 and discussed in the previous section, the relaxation of the cellular partitions from competing SD phases is performed using the *Surface Evolver* (SE) (9). While most commonly used for area optimization problems (e.g. dry foam models, minimal surfaces), SE generically optimizes a target function (e.g., energy, area) defined on the facets of triangulated surface mesh subject to various geometric constraints, for example periodic boundary conditions or volumes within bodies enclosed bodies (e.g., cells or bubbles).

For the DFM (and for Kelvin and Quantizer problem results below) we construct initial configurations that are input into SE by generating the Voronoi partitions from the point lattice positions of competitor structures within triply periodic, rectilinear box. The aspect ratio of the periodic cell dimensions and the initial coordinates of the cell centers are extracted from references listed for each FK lattice in Sec. C below. Voronoi cells are computed using *Voro++* (10) and then converted to SE input files via a custom python script. In addition to the initial topology of the “foam” structure, the SE input file also defines a discrete approximation of the dimensionless stretching, \mathcal{I} , area \mathcal{A} , and DFM free energy, \mathcal{F} .

Dimensionless area derives directly from computed total facet area and enclosed volumes of cells, while the stretching is computed as follows. For cell i in structure X , an initial center \mathbf{x}'_i is chosen as fixed reference point within the body. Since \mathbf{x}'_i is, in general, not the centroid of i , the stretching integral I_i splits into two contributions,

$$I_i = I'_i(\mathbf{x}'_i) - V_i |\mathbf{x}_i - \mathbf{x}'_i|^2 \quad [18]$$

where $I'_i(\mathbf{x}'_i) \equiv \int_{V_i} d^3\mathbf{x} |\mathbf{x} - \mathbf{x}'_i|^2$ is the second moment of V_i with respect to the reference point \mathbf{x}'_i . For a triangulated mesh composed of triangular facets with center \mathbf{X}_f , area ΔA_f , (outward) normal \mathbf{N}_f , these quantities can be approximated using the discrete sums,

$$I'_i(\mathbf{x}'_i) = \frac{1}{5} \sum_{f \in i} \Delta A_f \mathbf{N}_f \cdot (\mathbf{X}_f - \mathbf{x}'_i) |\mathbf{X}_f - \mathbf{x}'_i|^2, \quad [19]$$

and

$$\mathbf{x}_i - \mathbf{x}'_i = \frac{1}{4V_i} \sum_{f \in i} \Delta A_f [\mathbf{N}_f \cdot (\mathbf{X}_f - \mathbf{x}'_i)] (\mathbf{X}_f - \mathbf{x}'_i). \quad [20]$$

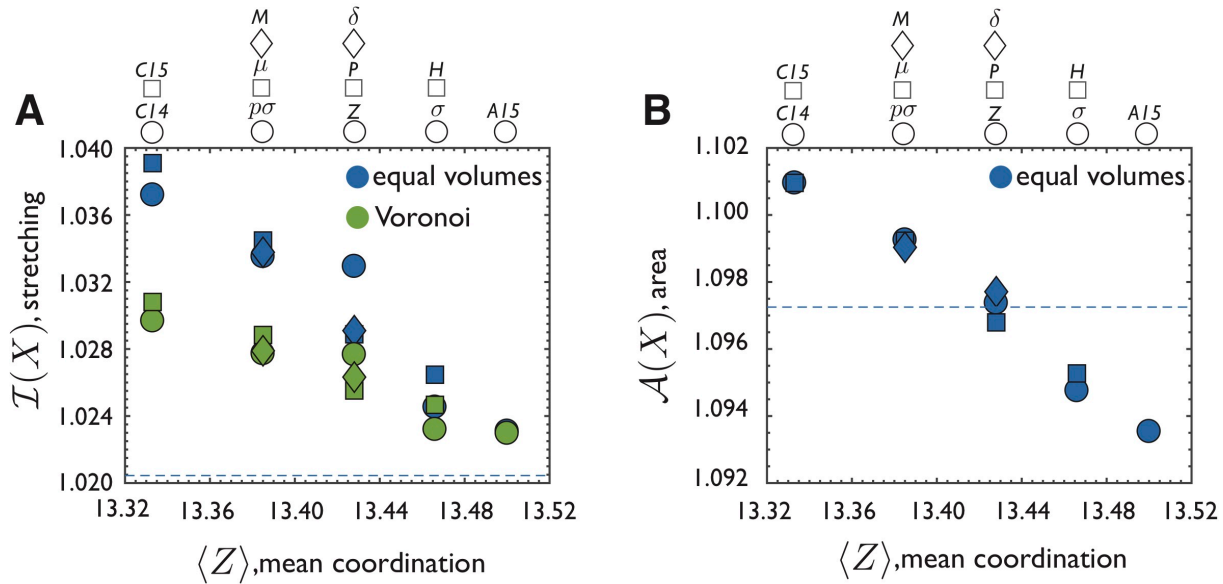


Fig. 2. (A) Computed minimal dimensionless stretching moments ($\mathcal{I}(X)$) computed from: equal volume constraint (blue) and Centroidal Voronoi Partitions (green). (B) Computed minimal dimensionless area ($\mathcal{A}(X)$) for equal volume cells (i.e. Kelvin problem results). Dashed lines show results for BCC.

These quantities are evaluated by use of `facet_general_integral` in SE. In the limit of $\Delta A_f \rightarrow 0$ these sums converge to the integral quantity with an error inversely proportional to the number of facets.

Cell shape optimizations are performed in order to minimize dimensionless free energy (\mathcal{F}), area (\mathcal{A}) or stretching (\mathcal{I}) for a given set of constraints, fixed periodicity, number of cells and with or without enforcing equal volume among distinct cells. Numerical optimization proceeds by successive interaction of vertex relaxation followed by mesh refinement steps. For each mesh refinement, vertices are relaxed until the optimized quantity (\mathcal{F} , \mathcal{A} or \mathcal{I}) changes by less than 10^{-6} . Mesh refinements proceed until the total change of post-relaxation value of the target quantity falls below 10^{-6} upon successive mesh refinements.

A. Minimal stretching cells: Quantizer problem. For comparison to DFM result, we compute the cell geometries that optimize \mathcal{I} for the competing FK lattices. This optimization is directly related to the *Quantizer problem* (11–13) that seeks the optimal decomposition of lattice into cells, at a given cell density, whose average square distance to the generating points \mathbf{x}'_i is minimal (i.e., minimal sum of the second moments $\int_{V_i} d^3\mathbf{x} |\mathbf{x} - \mathbf{x}'_i|^2$). For a fixed set of *generators*, \mathbf{x}'_i , the cells that minimize \mathcal{I} are given by the (unweighted) Voronoi partition (VP) derived from \mathbf{x}'_i . Additionally, for a fixed set of *cells*, V_i , the generators that minimize \mathcal{I} are given by the centroids $\mathbf{x}_i = V_i^{-1} \int d^3\mathbf{x} \mathbf{x}$ of V_i . Hence, VP whose generating points are cell centroids, so-called *centroidal Voronoi partitions* (CVP), are local minima of \mathcal{I} , for a given cell topology (14, 15). Therefore, in the context of the DFM, the CVP correspond to cell geometries that rigorously minimize the entropic cost of chain stretching.

We compute the CVP for competing lattices by minimizing $\mathcal{I}(X)$ within SE starting with generating points corresponding to reported lattice site positions of FK lattices (as summarized in Sec. 2C). CVP results from the minimization of \mathcal{I} at fixed mean volume of cells V_0 (fixed dimensions of the periodic repeat) but without constraints on the individual cell volumes V_i are shown in Fig. 2A. For comparison, we also compute the minimal stretching cells for *fixed, equal cell volumes*, $V_i = V_0$, a constraint which accounts for increased values of $\mathcal{I}(X)$ that decrease with mean coordination of cells $\langle Z \rangle$. This trend is consistent with main text Fig. 2 where the asymmetry in volume among cells for CVP is smallest for large- $\langle Z \rangle$ structures (A15) and largest at for small- $\langle Z \rangle$ structures (C14, C15). This trend implies that imposing the equal cell volume constraints requires a smaller distortion from the optimal stretching cell geometry for structures whose CVP are closest to equal volume (i.e. at large- $\langle Z \rangle$). Notably, no FK structure beats the minimal stretching value of BCC partition, $\mathcal{I}(\text{BCC}) = 1.0205$, proven to be the best quantizer among lattices in 3D (13).

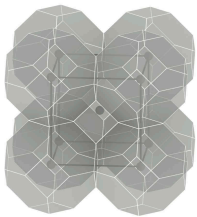
B. Minimal area cells: Kelvin problem. For comparison to DFM result, we also compute the cell geometries that optimize \mathcal{A} for the competing FK lattices. As shown earlier in Sec. 1B for flat faced cells (weighted VP) of A15, optimizing \mathcal{A} in the absence of the volume constraints is unstable due to shrinking of cells to zero volume, and hence is not well-defined with respect to comparison of partitions of different lattices X . A well-defined comparison is

621 possible with constraints on the relative volumes among cells, such as in the case of the *Kelvin problem*, which seeks 683
622 partitions into *equal volume cells* ($V_i = V_0$) that minimize mean cell area, or equivalently, surface area energy. The 684
623 results from the minimization of \mathcal{A} at fixed, equal volume of cells is shown in Fig. 2B. Notably, the minimal area 685
624 partition deriving from A15, the Weaire Phelan foam, is minimal (16), beating Kelvin’s conjectured optimal foam, 686
625 deriving from BCC. Partitions of FK structures σ and H also been previously reported (18, 19) to achieve lower areas 687
626 than the Kelvin foam[‡] (yet still larger than A15). Here, we report a third counterexample of Kelvin’s conjecture, 688
627 deriving from the FK lattice P ($\mathcal{A}(\text{BCC}) = 1.097251 > \mathcal{A}(\text{P}) = 1.096795 > \mathcal{A}(\text{A15}) = 1.093541$) as well as minimal 689
628 area (equal volume) results for four other previously unreported FK lattices, $\rho\sigma$, δ and M, all with dimensionless area 690
629 exceeding the Kelvin foam. 691

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640 **C. Competing lattices: Results.** Here, we summarize the SE results for competing structures in terms of dimensionless 702
641 free energy \mathcal{F} , dimensionless surface area energy \mathcal{A} , dimensionless stretching moment \mathcal{I} for various optimizations 703
642 (optimal area, stretching or DFM free energy) and under different constraints (with and without equal cell volume 704
643 constraints). For comparison, we also include dimensionless area, stretching and DFM free energy for Voronoi 705
644 partitions corresponding to the initial generating points (not necessarily centroidal). Additionally, we compare volume 706
645 histograms for both centroidal Voronoi partitions (upper histogram) and unconstrained DFM cells (lower histogram), 707
646 plotted in terms of the volume fraction ϕ_i occupied by cell type i , in each structure. For each structure, cell types 708
647 are classified and color coded in terms of number of faces: Z12 (blue); Z14 (green); Z15 (orange); and Z16 (red). 709
648 Additionally, in the histograms, cell types are annotated according to the Wyckoff positions corresponding to their 710
649 generating points. 711

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	Minimal Area	Minimal Stretching	Minimal \mathcal{F}		
			\mathcal{A}	\mathcal{I}	\mathcal{F}
Equal	1.097251	1.020450	1.097288	1.020622	1.071111
CVP	1.099000	1.020481	1.099000	1.020481	1.072177
Unconstrained	--	1.022976	1.097288	1.020622	1.071111
Initial Voronoi	1.099000	1.020481	1.099000	1.020481	1.072177

665 **Fig. 3.** BCC; space group: $Im\bar{3}m$; periodic cell: (cubic) a:b:c=1:1:1; $n_X = 2$, $\langle Z \rangle = 14$ 727

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676 [‡]We note that reported values for dimensionless area by Phelan, which from Fig. 2 of ref. (18) gives $\mathcal{A}(\sigma) = 1.094759 > \mathcal{A}(\text{H}) = 1.095271$, are slightly discrepant (and smaller than) with our 738
677 findings, reported in Sec. 2C 739

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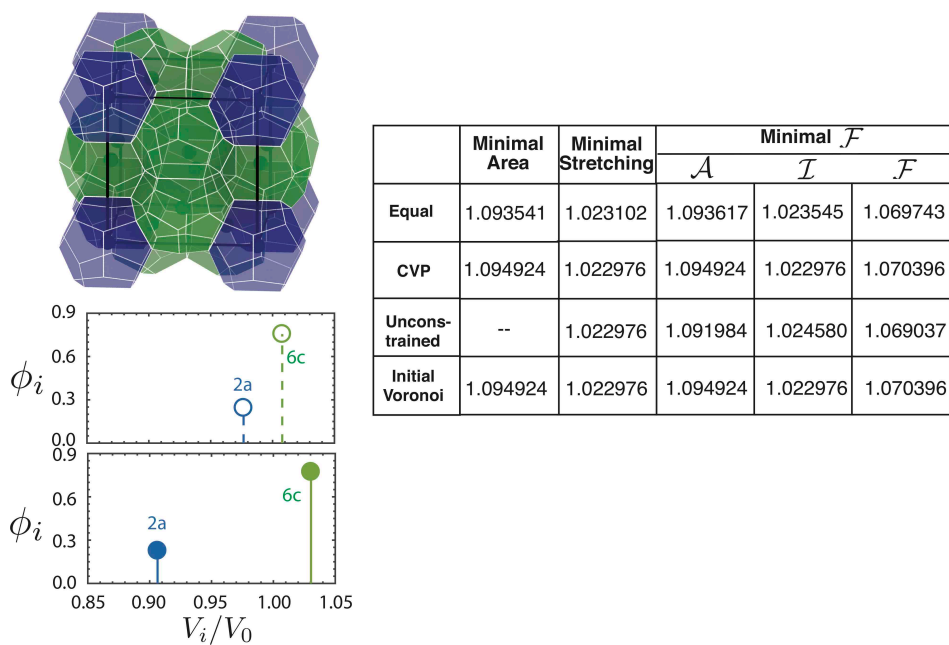


Fig. 4. A_{15} ; space group: $Pm\bar{3}n$; periodic cell: (cubic) a:b:c=1:1:1; $n_X = 8$; $\langle Z \rangle = 13.5$, init. coords.: Cr_3Si from ref. (20)

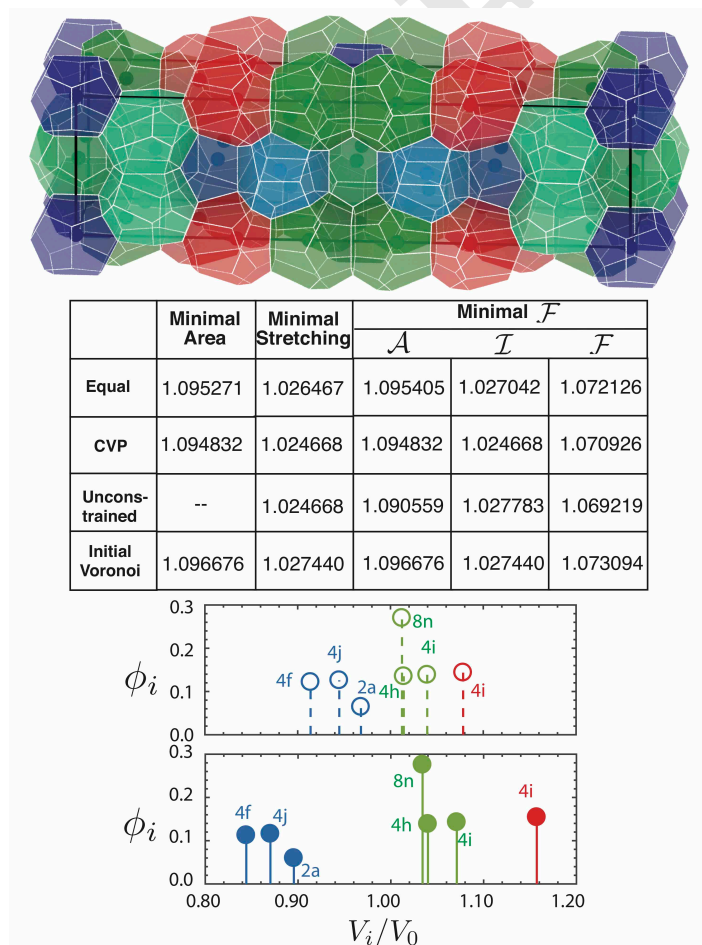


Fig. 5. H ; space group: $Cmmm$; periodic cell: (orthorhombic) a:b:c=1:3.88:1; $n_X = 30$; $\langle Z \rangle = 13.466$; init. coords.: ref. (22) (note that reference indicates Z12 spheres are situated at Wyckoff positions 2(a) and 4(e) whereas we have used positions 2(a) and 4(f))

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	Minimal Area	Minimal Stretching	Minimal \mathcal{F}		
			\mathcal{A}	\mathcal{I}	\mathcal{F}
Equal	1.094759	1.024536	1.094935	1.025203	1.071179
CVP	1.094909	1.023207	1.094909	1.023207	1.070467
Unconstrained	--	1.024536	1.090531	1.026001	1.068583
Initial Voronoi	1.094267	1.023697	1.094267	1.023697	1.070219

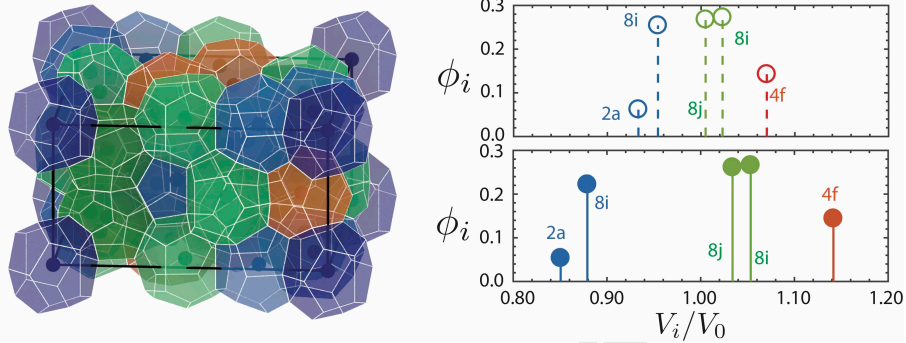


Fig. 6. σ ; space group: $P4_2/mnm$; periodic cell: (tetragonal) a:b:c=1.9:1.9:1; $n_X = 30$; $\langle Z \rangle = 13.466$; init. coords.: ref. (21)

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	Minimal Area	Minimal Stretching	Minimal \mathcal{F}		
			\mathcal{A}	\mathcal{I}	\mathcal{F}
Equal	1.097376	1.032937	1.097490	1.033350	1.075679
CVP	1.093898	1.027670	1.093898	1.027670	1.071360
Unconstrained	--	1.027670	1.088499	1.031766	1.069250
Initial Voronoi	1.093039	1.028173	1.093039	1.028173	1.070975

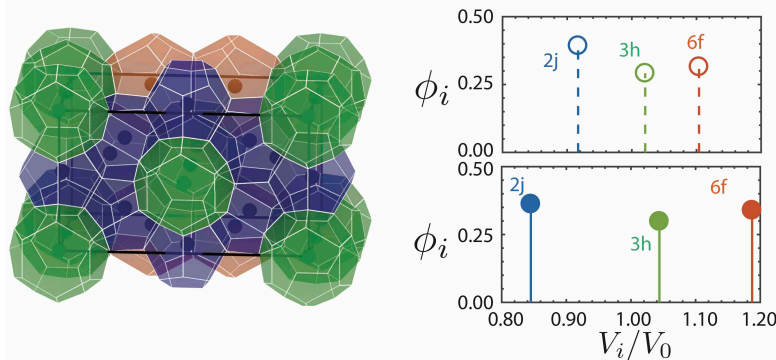
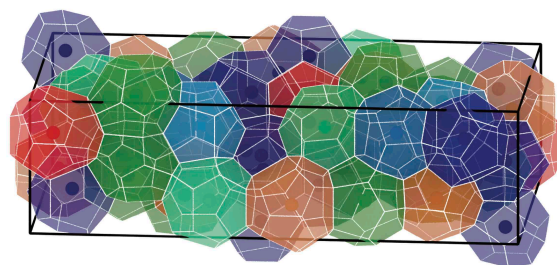


Fig. 7. Z ; space group: $P6/mmm$; periodic cell: (hexagonal*) a:b:c=1:1:0.993; $n_X = 7$; $\langle Z \rangle = 13.428$; init. coords.: ref. (23) (*For SE calculations, we have used an equivalent orthorhombic unit cell with twice the number of cells than that of hexagonal unit cell)

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	Minimal Area	Minimal Stretching	Minimal \mathcal{F}		
			\mathcal{A}	\mathcal{I}	\mathcal{F}
Equal	1.096795	1.028879	1.096934	1.029429	1.073954
CVP	1.094704	1.025525	1.094704	1.025525	1.071141
Unconstrained	--	1.025525	1.089421	1.029037	1.068909
Initial Voronoi	1.093971	1.026022	1.093971	1.026022	1.070836

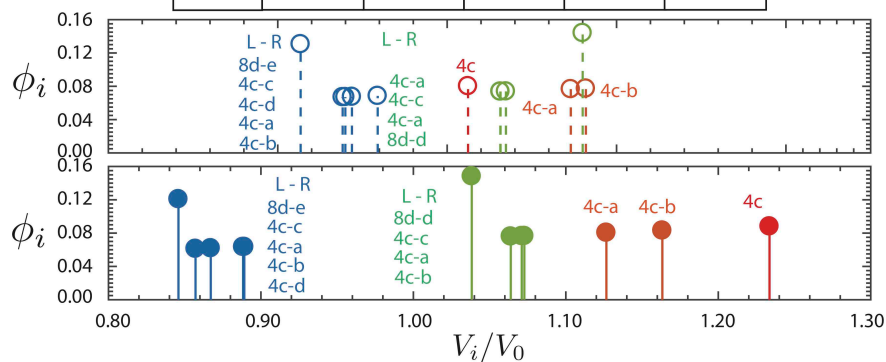


Fig. 8. P ; space group: $Pbnm$, periodic cell: (orthorhombic) $a:b:c=1.91:3.57:1$; $n_X = 56$; $\langle Z \rangle = 13.428$; init. coords.: ref. (24)

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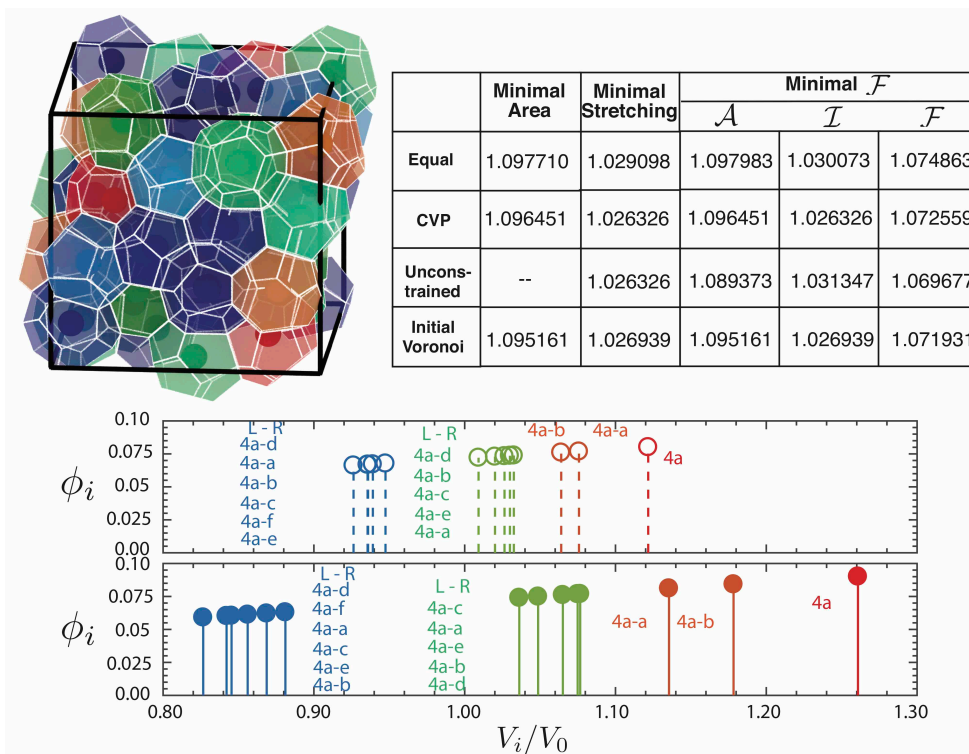


Fig. 9. δ ; space group: $P2_12_12_1$; periodic cell: (orthorhombic) $a:b:c=1.03:1.03:1$; $n_X = 56$; $\langle Z \rangle = 13.428$; init. coords.: ref. (25)

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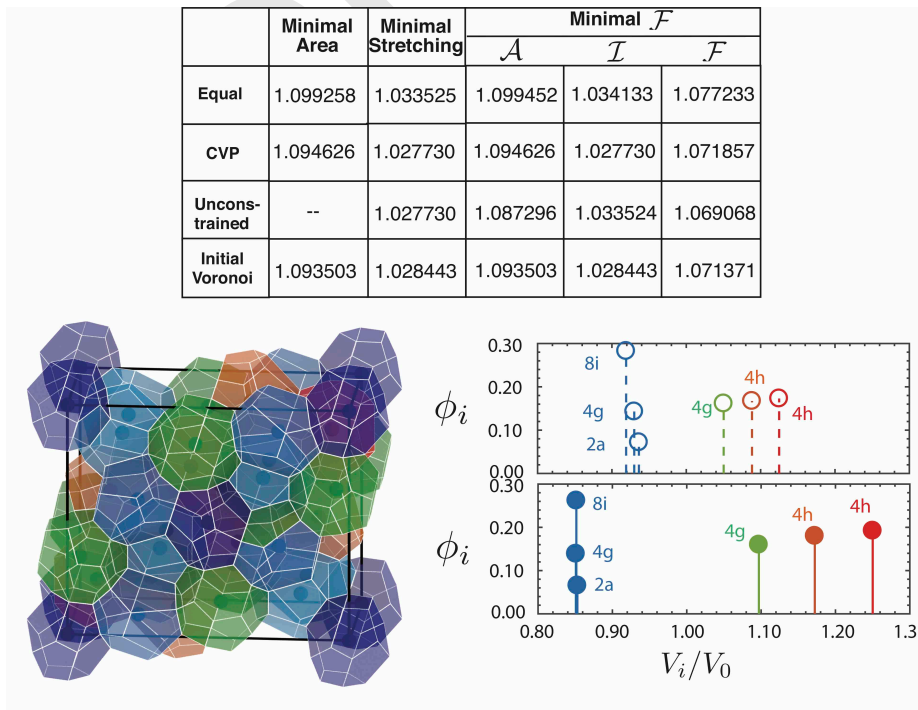


Fig. 10. $p\sigma$; space group: $Pbam$; periodic cell: (orthorhombic) $a:b:c=1.95:1.64:1$; $n_X = 26$; $\langle Z \rangle = 13.385$; init. coords.: ref. (26)

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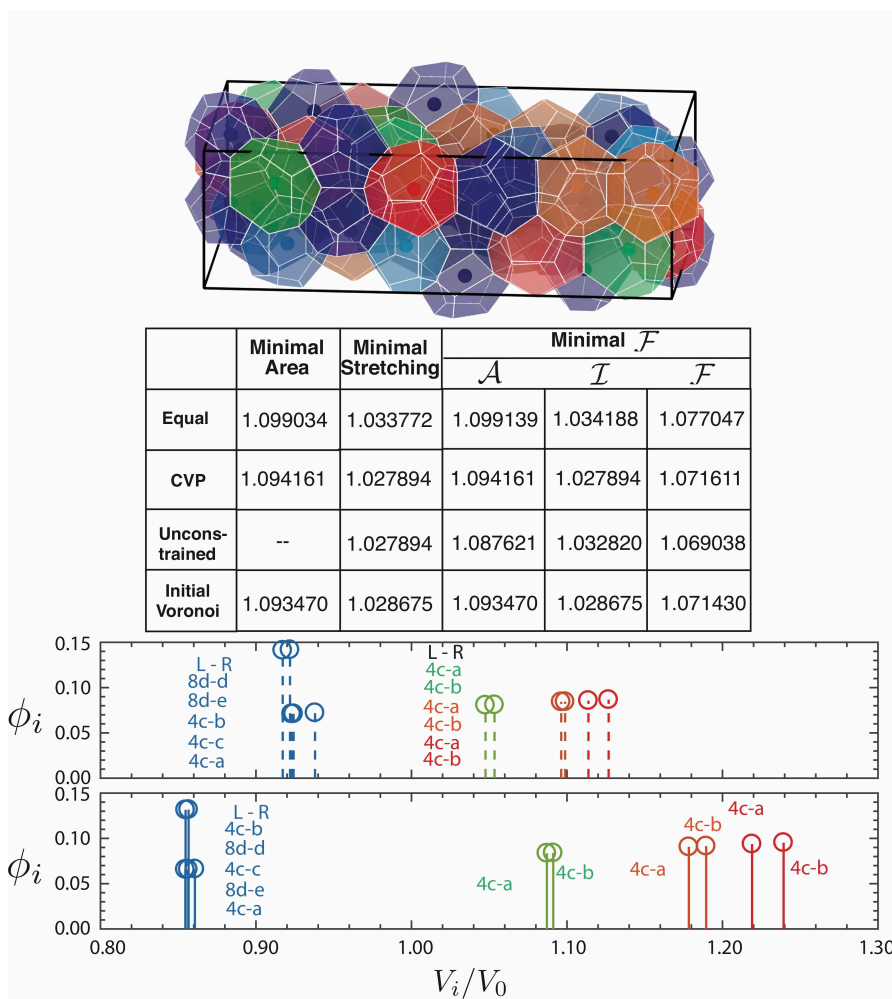


Fig. 11. M ; space group: $Pnam$; periodic cell: (orthorhombic) $a:b:c=1.89:3.3:1$; $n_X = 52$; $\langle Z \rangle = 13.385$; init. coords.: ref. (27)

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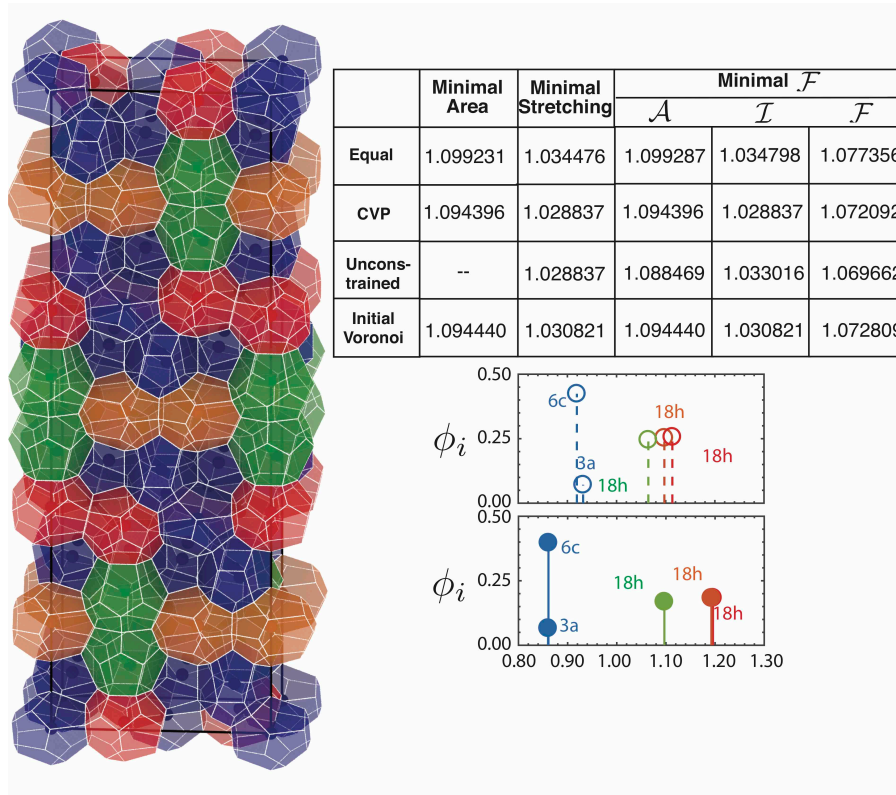


Fig. 12. μ ; space group: $R\bar{3}m$; periodic cell: (hexagonal*) a:b:c=1:1:5.2; $n_X = 39$; $\langle Z \rangle = 13.385$; init. coords.: W_6Fe_7 from ref. (20) (*For SE calculations, we have used an equivalent orthorhombic unit cell with twice the number of cells than that of hexagonal unit cell)

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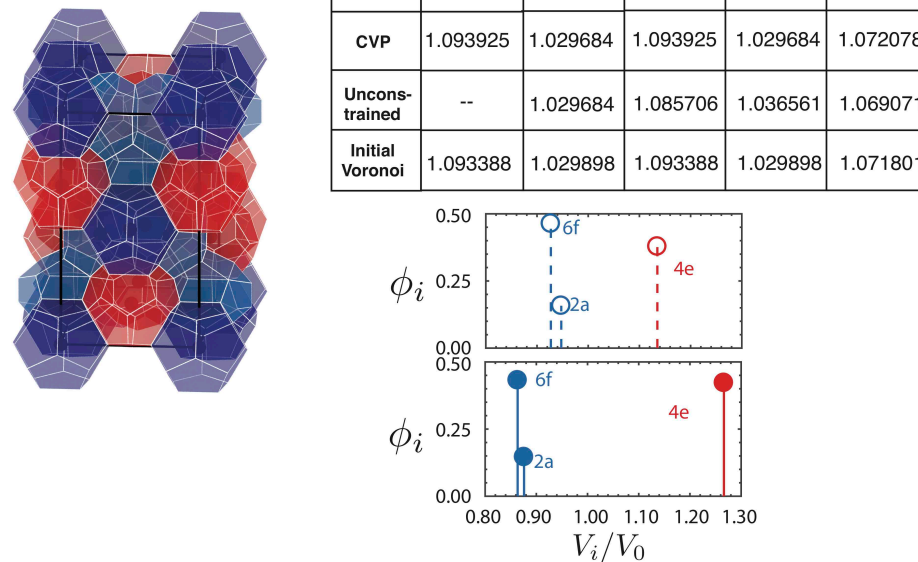


Fig. 13. $C14$; space group: $P6_3/mmc$; periodic cell: (hexagonal*) a:b:c=1:1:1.63; $n_X = 12$; $\langle Z \rangle = 13.333$; init. coords.: $MgZn_2$ from ref. (20) (*We have used an equivalent orthorhombic unit cell with twice the number of cells than that of hexagonal unit cell)

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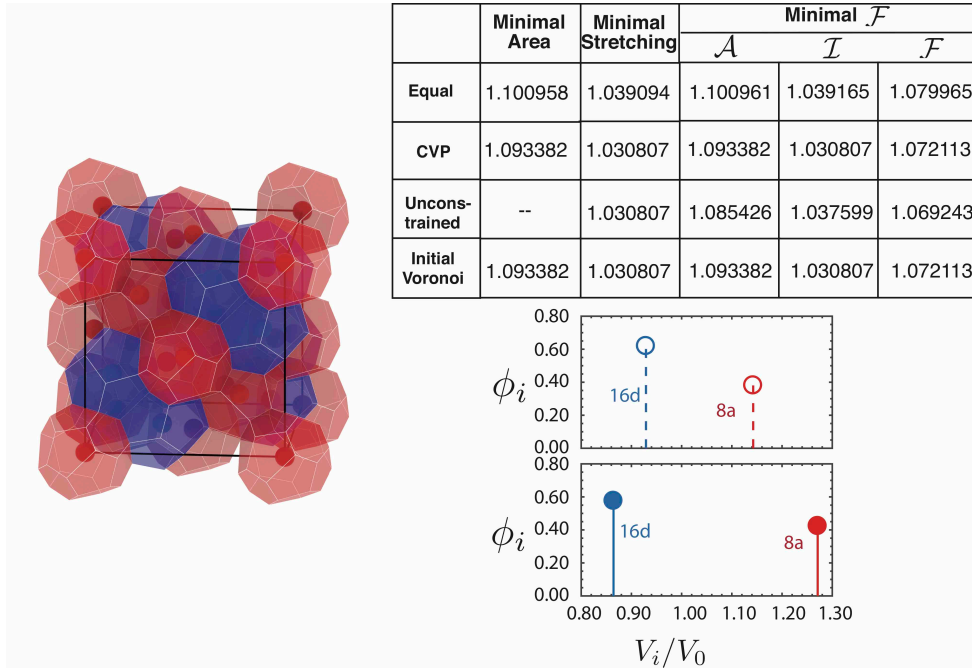


Fig. 14. C15; space group: $Fd\bar{3}m$, periodic cell: (cubic) a:b:c=1:1:1; $n_x = 24$; $\langle Z \rangle = 13.333$; init. coords.: Cu_2Mg from ref. (20)

3. Self-consistent field theory of conformationally asymmetric diblocks

We use self-consistent field theory (SCFT) of a Gaussian chain model of diblock copolymer melts (7) to predict structure and thermodynamics of a multi-chain qSD formation. In particular, we consider a model where chains possess $N_A = fN$ and $N_B = (1 - f)N$ segments of A and B type monomers each having statistical segment lengths as a_A and $a_B = \epsilon^{-1}a_A$ respectively but having the same segment volume ρ_0^{-1} , with the Flory-Huggins interaction parameter χ describing the enthalpic repulsion between A and B blocks. In SCFT, the key statistical quantities are the chain distribution functions $q(\mathbf{x}, n)$ and $q^\dagger(\mathbf{x}, n)$ which capture the statistical weights (constrained partial partition functions) of chains “diffusing” from their respective A and B ends to the n th segment located at position \mathbf{x} . Following methods described in ref. (30) and elsewhere, these are determined self-consistently according to inter-segment interactions deriving from the mean compositions profiles $\phi_{A,B}(\mathbf{x}) = \frac{V}{NQ} \int_{A,B} dn q(\mathbf{x}, n) q^\dagger(\mathbf{x}, n)$, where $Q = \int d\mathbf{x} q(\mathbf{x}, n) q^\dagger(\mathbf{x}, n)$ is the single chain partition function and $\int_{A,B} dn$ corresponds to the integration over the A or B block segments.

A. Thermodynamics of SD phases. Here, we summarize results for thermodynamics of qSD phases, in comparison to DFM predictions, for conformational asymmetries $\epsilon > 1$. For modest conformational asymmetry, i.e $\epsilon=1.5$ and 2, Xie *et al.* (31) and Kim *et al.* (32) have shown that σ is the equilibrium for AB diblock copolymers in melt over a range of compositions between a stable BCC (low- f) and hexagonally ordered cylinders (high- f). Kim *et al.* have additionally reported results for FK candidates, σ , A15, Z, C14 and C15, for $\chi N = 40$, which we analyze in more detail. In Fig. 15 we also report two new metastable structures H and $p\sigma$ for AB diblocks at $\chi N = 25$. Although metastable, these phases all beat BCC over a range of f , and H is shown to be competitive with σ and A15 over the entire range of metastable compositions studied, $0.23 \leq f \leq 0.33$.

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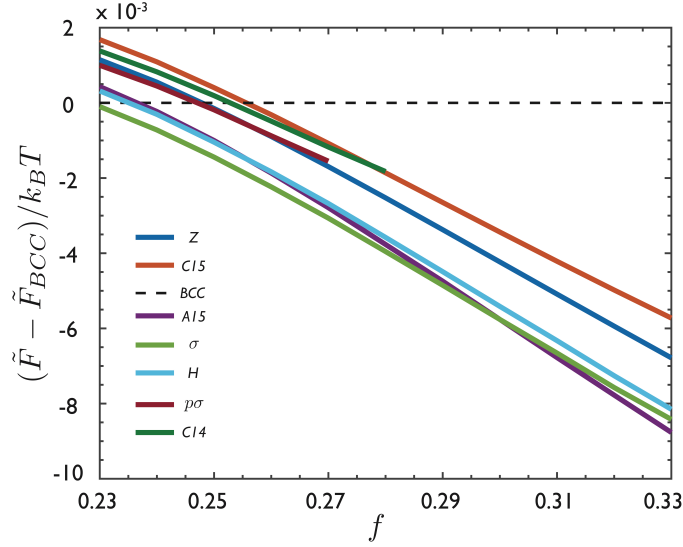


Fig. 15. Relative Free energy per chain of FK lattices w.r.t BCC as a function of volume fraction for $\chi N=25$

To compare the DFM predictions to SCFT results, for σ , A15, Z, C14 and C15, at the highest segregation strength computed ($\chi N = 40$) we normalize the free energy per chain by the value of A15, as plotted in Fig. 16A, since according to DFM, the free energy per chain for each structure is proportional to the same quantities (a function of χN , ϵ and f) that vary with cell geometry. While DFM models are strictly constant with f and SCFT results show at least slight variation of relative free energy with f , we note that the relatively close free energies of SCFT are remarkably consistent with scale of separation predicted by DFM predictions.

We also compare the relative ranking of σ , A15, Z, C14 and C15 in terms of the enthalpic and entropic contributions to the free energy per chain, $\tilde{F}'_{enthalpy} = V^{-1} \int d^3\mathbf{x} \chi N \phi_A(\mathbf{x}) \phi_B(\mathbf{x})$, and $\tilde{F}'_{entropy} = \tilde{F}'_{tot} - \tilde{F}'_{enthalpy}$, which are computed from SCFT solutions as described in ref. (7) and elsewhere (here, primed quantities refer to values derived from SCF and unprimed quantities refer to their values from DFM). To extract strictly the geometric dependence of these thermodynamic quantities, we note from the DFM model (predicated on the strong-segregation and the polyhedral interface limits) that

$$\tilde{F}'_{enthalpy} = \frac{\gamma \mathcal{A}}{R_0}; \quad \tilde{F}'_{entropy} = \frac{\kappa}{2} \mathcal{I} R_0^2 \quad (\text{DFM}) \quad [21]$$

which motivates the definition of scaled-enthalpy \mathcal{A}' and scaled-entropy \mathcal{I}' computed from SCF results for $\tilde{F}'_{enthalpy}$ and $\tilde{F}'_{entropy}$, appropriately scaled by the mean sphere radius R'_0 (the radius of a sphere of equal mean volume to equilibrium SD for a given structure) according to

$$\mathcal{A}' \equiv \gamma^{-1} \tilde{F}'_{enthalpy} R'_0; \quad \mathcal{I}' \equiv \frac{2 \tilde{F}'_{entropy}}{\kappa R_0'^2} \quad (\text{SCFT}). \quad [22]$$

This definition scales out the variation of enthalpic and entropic contributions due to the difference in mean domain sizes from structure to structure. Fig. 16 B and C plots the respective SCFT results for scaled enthalpy ($\mathcal{A}'(X)$) and scaled entropy ($\mathcal{I}'(X)$) normalized by the value for A15, and compared to DFM predictions. Additionally, Fig. 16 D plots the mean domain sizes $R'_0(X)$ (relative to A15) computed from SCFT, which largely confirms that generic prediction of DFM, in eq. (13) that structures corresponding to relatively small stretching costs favor relatively larger domain sizes (aggregation numbers per sphere).

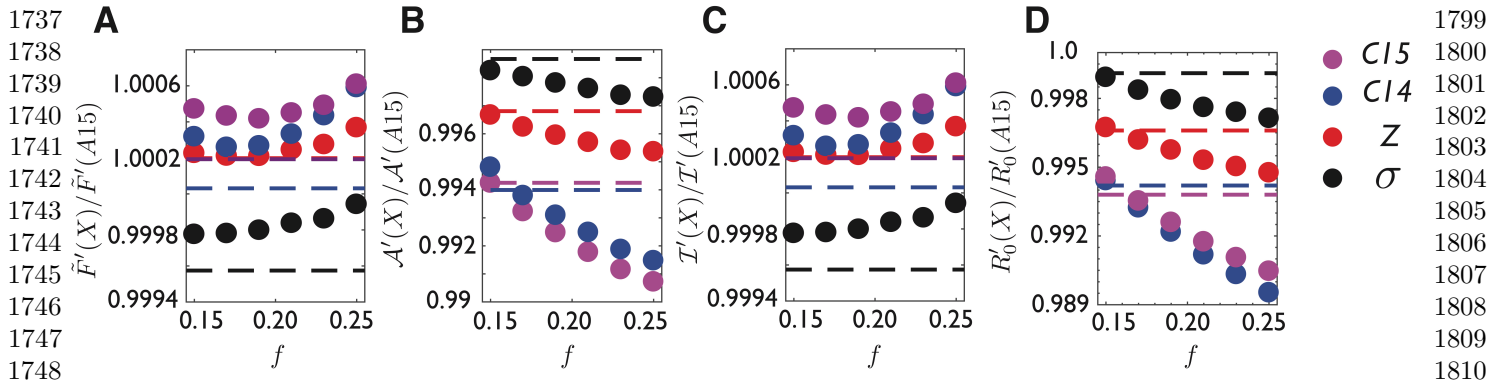


Fig. 16. (A) Relative free energy (B) Scaled enthalpy (C) Scaled entropy of FK lattices relative to A15 from SCF (circles) and DFM (dashed lines) (D) Relative mean domain sizes of FK lattices (relative to A15). SCF results are for $\chi N = 40$ and $\epsilon = 2$.

B. Geometric analysis of spherical domains.

The geometry of SD core shapes predicted by SCFT are analyzed in terms of the AB interface, which can be extracted from the equilibrium composition profiles, specifically the isosurfaces where A and B have equal volume fractions, $\phi_A(\mathbf{x}) = \phi_B(\mathbf{x}) = 0.5$. From the isosurfaces, numerically extracted using MatLab, the total areas and enclosed volumes within each SD in the predicted SCFT structure can be directly computed. Because the core blocks constitute a fixed fraction f of the entire chain, the core volume accounts for the same fixed fraction of the entire qSD. As shown in the main text for A15 and BCC at $f = 0.29$, the areal distortion parameter, α_i , of the core interface varies with conformational asymmetry. In Fig. 17 we also show results at higher core composition, $f = 0.34$. These indicate that degree of polyhedral warping of interface increases with f , due to the enhanced proximity of the interface to the outer boundary (or cell “wall”) between neighbor domains. Fig. 17 also shows the variation of α_i with f for C15, highlighting the presence of two populations of SD in the structure: relatively spherical Z16 domains (low α_i), and less spherical Z12 domains (higher α_i).

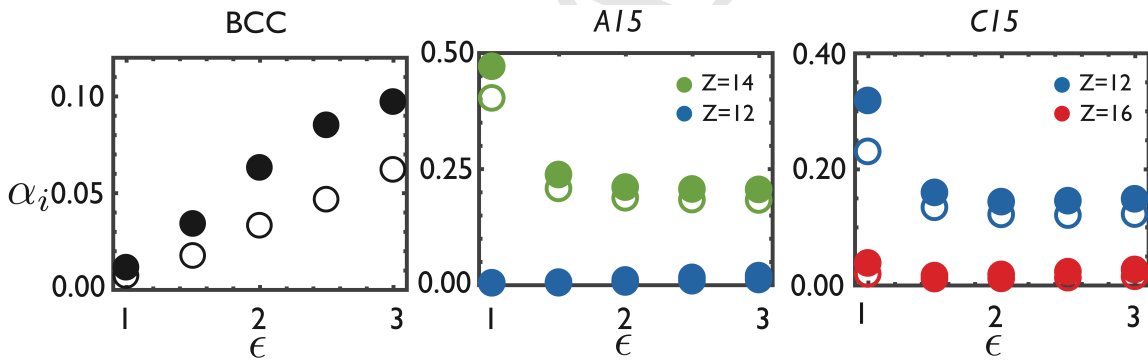


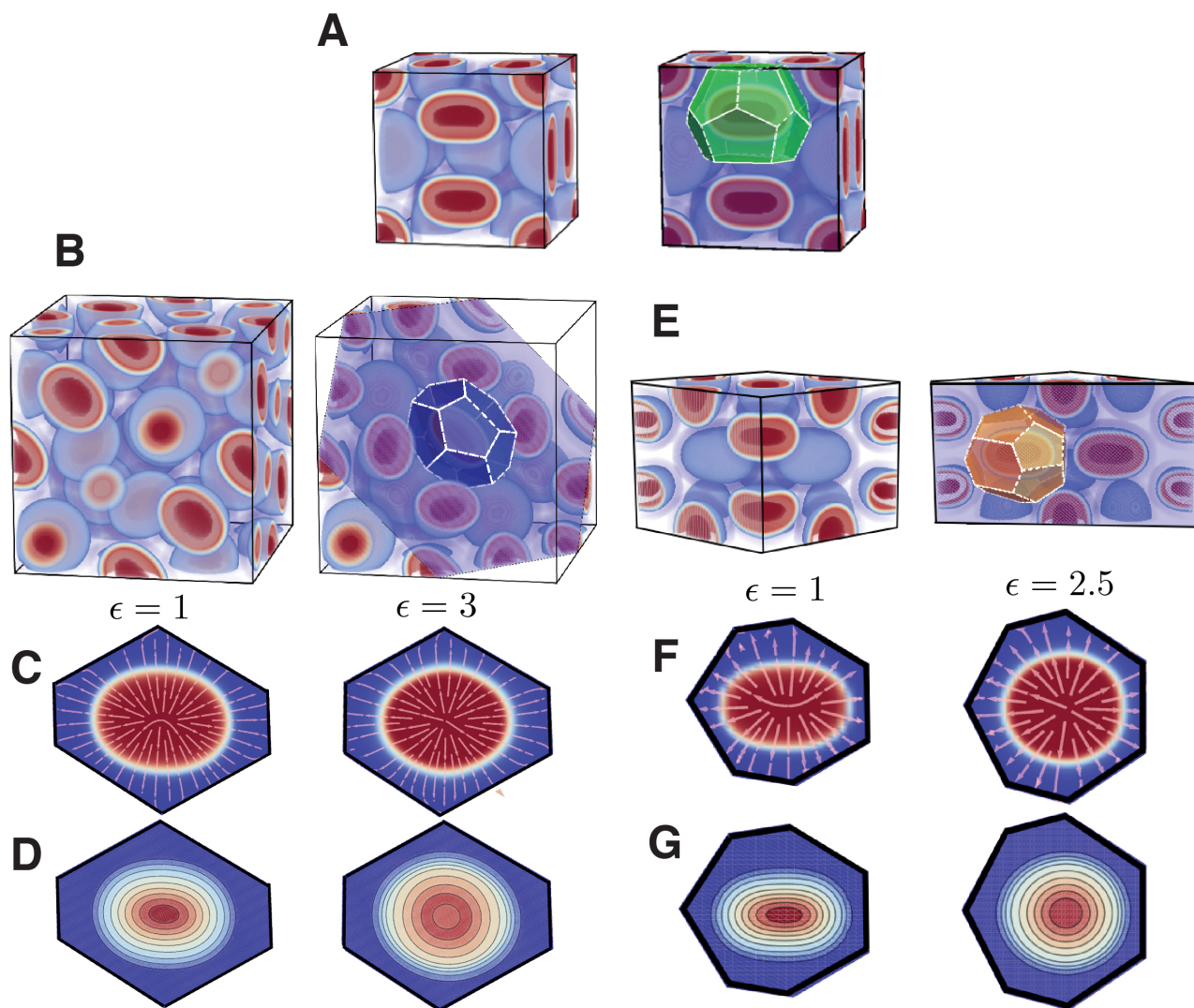
Fig. 17. Measure of areal distortion of AB interface of distinct domains for competing FK phases computed from SCF predictions at $\chi N = 40$, $f = 0.29$ (open circles) and $f = 0.34$ (filled circles).

Like the case for the Z14 cells of A15 (shown in main text Fig. 3), these higher- α_i cells of C15 and Z also undergo a discoidal-to-radial transition as ϵ is varied from 1 (conformationally symmetric) to ≈ 2 (conformationally asymmetric). To analyze the intra-domain structure of chain packing in more detail, we compute the *polar* orientational order parameter, $\mathbf{t}_A(\mathbf{x})$, of A block segments using methods described in ref. (29),

$$\mathbf{t}_A(\mathbf{x}) = \frac{V}{6NQ} \int_A dn [q \nabla q^\dagger - q^\dagger \nabla q], \quad [23]$$

where the vector orientation of the segments is defined to point from the free A end towards the junction point along the chains. These orientational profiles are shown in 2D sections through spherical domains of BCC and A15 in Fig. 3 of the main text. In Fig. 18 we also show 2D cuts through Z12 domains and Z15 domains of C15 (B,C) and Z (E,F), respectively. Note that only streamlines of $\mathbf{t}_A(\mathbf{x})$ are shown in these figure, and thus only the local orientation of A segments, but not the degree of alignment, is visible. Both of these domains show a trend consistent with the sub-domain morphology of the Z14 cell in A15. For conformationally symmetric chains ($\epsilon = 1$), the interface shape is more oblate than the polyhedral cell enclosing the domain (which itself can be observed from the flow lines

1861 about the orientational order parameter), and the core regions are composed of a quasi-lamellar “puck” encircled 1923
 1862 by quasi-toroidal rim. In contrast, when conformational asymmetry imposes a sufficiently larger cost on coronal 1924
 1863 stretching ($\epsilon \gtrsim 2$), the core interface shape is more consistent with an affinely shrunk (and somewhat rounded) copy 1925
 1864 of the polyhedral cell, and segment orientation becomes consistent with uniformly radial extension of the chains from 1926
 1865 the domain center. The sub-domain distinctions between radial and discoidal domain shapes is further highlighted by 1927
 1866 comparing the respectively focussed vs. spatially spread distributions of A-block ends, computed from $q^\dagger(\mathbf{x}, n = 0)$ 1928
 1867 and shown in Fig. 18 D and G. 1929



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Fig. 18. (A) 3D density plot of core block forming A15 from SCFT data at $f=0.29$, $\epsilon=1$, $\chi N = 40$ in primitive cell (on left) with the Z14 cell surrounding a corresponding qSD on the $[100]$ face shown in green (on right) corresponding to the sections shown in main text Fig. 3. (B) shows the same but for the C15 structure, and a 2D section through a $\langle 111 \rangle$ plane through the center of a Z12 cell shown in blue (on right). The composition and segment orientation for the Z12 domain of C15 are shown in (C), with the end distribution of the core A-block shown in (D) for conformationally symmetric and asymmetric cases. (E) shows a hexagonal cell of Z phase from SCFT results at the same conditions at (A) and (B) (on left), with a cut through the center of the Z15 cell shown in orange (on right). The composition and segment orientation for the Z15 domain of Z are shown in (F), with the end distribution of the core A-block shown in (G) for conformationally symmetric and asymmetric cases.

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