Aggregation kinetics in a model colloidal suspension

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We present molecular dynamics simulations of aggregation kinetics in a colloidal suspension modeled as a highly asymmetric binary mixture. Starting from a configuration with largely uncorrelated colloidal particles the system relaxes by coagulation-fragmentation dynamics to a structured state of low-dimensionality clusters with an exponential size distribution. The results show that short-range repulsive interactions alone can give rise to so-called cluster phases. For the present model and probably other, more common colloids, the observed clusters appear to be equilibrium phase fluctuations induced by the entropic inter-colloidal attractions.

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Colloidal suspensions are common in biological and geological systems and their unusual properties are intensely exploited in many industrial applications, from pharmaceutics to food processing. Although some, e.g. noble metal colloids, have been produced and used for centuries, the systematic study of these systems is a more recent endeavor \[1]. One of the remarkable features of colloids is the possibility of tuning the colloidal interactions \[2]. This multiplies the already rich physical behavior exhibited by colloids \[3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14] and significantly increases their technological potential, but it also underscores the need for understanding the general principles and mechanisms that govern their equilibrium and kinetic behavior. The low volume fractions kinetic arrest leading to gelation in weakly attractive hard-sphere colloids has been found for example to share many of the features of the high volume fractions colloidal glass transition \[14]. The aggregation of colloids and the ensuing highly structured states such as gels are a direct consequence of the short-ranged attractions typically encountered in these systems, known as depletion or entropic interactions \[15]. The role of repulsions however is also important, and recently much attention has been devoted to elucidating the effect of long-range repulsions in the formation of cluster phases and the gelation transition \[7, 16, 17]. Understanding clustering effects and the mechanisms that control them is important conceptually as well as practically in systems ranging from biological to engineered ones, e.g. nanofluids. In the present paper we explore using molecular dynamics (MD) simulations the characteristics of aggregation kinetics in a model colloidal suspension with short-range repulsions that explicitly includes both colloidal and solvent particles. We find that the system evolution is well described as coagulation-fragmentation dynamics surprisingly similar to a simple polymerization-depolymerization process. The final state is a structured phase of low-dimensionality clusters with an exponential size distribution. These properties appear to be all due to the bond-like character of the induced short-ranged inter-colloidal attractions.

The simplest instance of a colloidal suspension is perhaps a binary hard-sphere mixture with significant size asymmetry. Dijkstra et al. \[18] have shown that the phase behavior of such a system can be accurately predicted by integrating out the effect of the small particles and introducing an effective depletion potential \[15] between the large (colloidal) particles. This approach provides significant insight into the structure of the phase diagram and simplifies considerably the computational study of the mixture. However, it may not be necessarily appropriate for the study of non-equilibrium processes in colloids. The motion of colloidal particles suspended in viscous liquids is generally coupled, even at large separations, by the induced suspending-liquid flows. These so-called hydrodynamic interactions are believed for example to play an important role in the aggregation kinetics \[19]. In the following we consider and fully model using MD simulations a system related to the binary hard-sphere mixture, consisting of two types of particles, 1 - colloid and 2 - solvent, with equal masses \(m\). The interactions between the particles are based on the inverse-12, 'soft-sphere' potential, whose properties have been well studied \[20\].

\[u(r) = \epsilon \left(\frac{\sigma}{r}\right)^{12}\] (1)

, and which we truncate and shift at \(r/\sigma = 2\). (We also define \(u(r) = \infty\) for \(r < 0\).) The interactions are:

\[u_{11}(r) = u(r - 2R_c)\] (2a)
\[u_{12}(r) = u(r - R_c)\] (2b)
\[u_{22}(r) = u(r)\] (2c)

Similar potentials, that take into account the 'size' of the colloidal particles by introducing a hard core radius \(R_c\), have been employed before to model suspensions \[21\]. For temperatures \(k_B T \approx \epsilon\) the effective diameters corresponding...
to the above interactions should be well approximated by $\sigma_2 = \sigma$, $\sigma_{12} = R_c + \sigma$, and $\sigma_1 \equiv \sigma_c = 2R_c + \sigma$, and satisfy additivity, $\sigma_{12} = (\sigma_1 + \sigma_2)/2$. The particular system that we focus on has a colloid-solvent ‘diameter’ ratio $\sigma_c/\sigma = 5$, colloid ‘volume fraction’ $\phi_c = \pi n_c \sigma_c^3/6 = 0.1$ and solvent ‘volume fraction’ $\phi = \pi n \sigma^3/6 = 0.37$; $n_c$ and $n$ are the corresponding number densities, $n_c = N_c/V$, $n = N/V$. We perform MD simulations of this system in the microcanonical (NVE) ensemble with $N_c = 200$, $N \approx 10^5$ and an average temperature set to $k_B T = \epsilon$. Such simulations of binary mixtures with large size ratios have been done before [22], but not, to our knowledge, for colloid volume fractions as small as the present one.

The simulations are initiated from a configuration of non-overlapping (as defined by the effective diameters), but otherwise randomly distributed colloid and solvent particles, and the system is equilibrated at $k_B T = \epsilon$ using a combination of velocity rescaling and Anderson thermostatiing. Thermal equilibrium is achieved over time scales much shorter than the scales over which the colloidal particles move over any significant distances, about 5000 MD steps, leading to a state corresponding to a fully equilibrated solvent and essentially uncorrelated colloidal particles. A largely similar configuration would be expected for example in the case of a hard-sphere colloid shortly after a rapid quench from a high temperature state. The system evolution is subsequently followed using conventional (constant energy) MD for over $5 \times 10^5$ time steps.

The evolving configuration of the large (colloidal) particles during the simulations is best characterized by using a standard aggregation criterion, that places two particles in the same cluster when their relative distance is smaller than a certain value $r_0$. For the present work we set $r_0 = 1.05 \sigma_c$, but our conclusions are independent of this value in a fairly wide range. The application of this criterion yields a time dependent cluster size distribution $N_k(t)$, or in scaled form $n_k(t) = N_k(t)/N_c$, where $N_c = \sum_k n_k N_k(t)$ is the (constant) total number of particles. We show in Fig. 1 the evolution of the (normalized) number of monomers $n_1(t)$, along with the inverse of the (normalized) total number of clusters, $s_1(t) = \sum_k n_k(t)/\sum_k N_k(t)$; $s_1$ can also be interpreted as a measure of the average cluster size. The number of monomers in the system steadily decreases, while the cluster size increases, with both quantities apparently reaching at late times well defined plateaus. Visual inspection of the clustering dynamics reveals tenuous structures reminiscent of branched polymers - see Fig. 2 - whose evolution is driven by aggregation and breakup processes. These features suggest that general concepts developed to describe coagulation-fragmentation kinetics [23, 24] may also be relevant to the present system. The starting point for this analysis is the generalized Smoluchowski equation for the cluster size distribution:

$$\frac{dn_k}{dt} = \frac{1}{2} \sum_{i+j=k} [K_{ij} n_i n_j - F_{ij} n_{i+j}] - \sum_{j=1}^{\infty} [K_{kj} n_k n_j - F_{kj} n_{k+j}]$$

(3)

with coagulation and breakup rate coefficients $\{K_{ij}\}$ and $\{F_{ij}\}$, $i, j = 1, \infty$. The cluster population $n_k(t)$ whose evolution is described by the above equations is generally assumed to satisfy a dynamical scaling relation [23, 24]

$$n_k(t) = s(t)^{-2} \phi(x)$$

(4)

where $x = k/s(t)$, $s(t)$ is a characteristic cluster size and $\phi(x)$ a scaling function dependent on the details of process, i.e. the coefficients $\{K_{ij}\}$ and $\{F_{ij}\}$. The applicability of Eq. (4) to systems that reach equilibrium, i.e. where $s(t)$ goes to a constant value when $t \to \infty$, appears to be somewhat more limited [24] than initially assumed. Nevertheless, under certain reasonable assumptions for the coagulation and breakup rates and in particular for a classical model of polymerization-depolymerization with constant coefficients [26] dynamical scaling holds exactly. For small deviations from equilibrium it can also be shown [24] that dynamical scaling implies that $s(t)$, and therefore $s_1(t)$, relaxes exponentially to its final, equilibrium value [24]. Furthermore, for the case of Ref. [24] the product $n_1(t)s_1^2(t)$ is time independent, $n_1(t)s_1^2(t) = 1$, while in general should increase only slowly in a narrow range.

For the asymmetric binary mixture considered here as an archetype of a colloidal suspension we find that $s_1(t)$ relaxes to a very good approximation in an exponential fashion, and moreover that the relation $n_1(t)s_1^2(t) = 1$ is very close to being satisfied - see Fig. 1. This makes the behavior of this simple model, containing only short-range repulsive potentials, surprisingly similar to that of common associating systems. As a test of dynamical scaling we consider the usual measure of the mean cluster size [23, 24], $s(t) = \sum_k k^2 n_k(t)$, which depends linearly on $s_1(t)$ for the model of Blatz and Tobolsky [24], and in general should be proportional with it when $s$ is large enough. This is also rather well satisfied, see Fig. 1(inset).

The morphology of the typical clusters - see Fig. 2 - closely resembles the open structures usually encountered in colloidal aggregation [19, 24]. Although the observed assemblies are not nearly as large as those obtained upon irreversible flocculation [27], their dimensionality can be well quantified by considering their radius of gyration $kR_g^2(k) = \langle \sum_{i=1}^{\infty} [(x_i - x_{CM})^2 + (y_i - y_{CM})^2 + (z_i - z_{CM})^2] \rangle$, where $i$ denotes the particles in a cluster of size $k$,
CM is the cluster center of mass, and the average is taken over the final population of clusters of size $k$. The resulting radii - see Fig. 3 - obey $R_g(k) \propto k^{1/D_f}$, with $D_f \approx 2$. The fractal dimension $D_f$ is slightly higher than the one observed for typical colloidal aggregates \cite{27}, but in agreement with that of randomly branched polymers \cite{28}.

The cluster size distribution in the final state - $n_n(k)$, see Fig. 4 - compares well with the distributions obtained in model coagulation-fragmentation dynamics \cite{23,26} and may be reasonably described at large cluster sizes as a decaying exponential. From the point of view of the kinetics this appears to be a consequence of an approximately cluster-independent break-up rate \cite{23,26}, which is consistent with the low cluster dimensionality and the equivalence for this system of the 'bonds' that hold together structures of different sizes. In fact, thermodynamic arguments applicable to equilibrium polymers systems \cite{24} yield a similar prediction, while general concepts pertaining to phase fluctuations (and nucleation) in solutions \cite{30} would also predict an exponentially decreasing abundance of large clusters for single-phase equilibrium close to a two-phase coexistence region. Nevertheless, since the size of our system in terms of the number of large particles modeled is fairly small, we further address this issue by considering an equivalent one-component model with effective interactions.

To this end we consider the structure of the mixture, in particular the colloid-colloid pair correlation function, $g_{11}(r)$ - Fig. 5. This function exhibits the sharp nearest-neighbor peak also encountered in the effective one-component modeling of asymmetric binary hard-sphere mixtures \cite{18}, which is due to the strong, entropically induced attractions. However, $g_{11}(r)$ is also considerably more structured than such one-component correlation functions due to the explicit presence of the solvent. For our own single component modeling we consider particles interacting through a potential $V_{eff}(r)$ defined by $\beta V_{eff}(r) = -\ln[g_{11}(r)]$, i.e. the potential of mean force, but which is truncated after its first maximum at a distance $r_m$ satisfying $V_{eff}(r_m) = 0$. We expect the resulting interaction to be an appropriate effective pair potential for the rather low colloid volume fraction system studied here. $V_{eff}$ has a fairly deep well followed by a barrier - see Fig. 5 (inset) - and is analogous with the effective potentials used to study the phase diagram of asymmetric binary hard-sphere mixtures \cite{18}. MD simulations with 864 particles interacting through this potential at a temperature $k_B T = \epsilon$ were carried out. The system requires very long equilibration times, of the order of $10^6$ steps, which we then follow by standard (NVE) MD to determine the fluid structure. The main features of the resulting pair correlation function - Fig. 5 - are not surprising. The sharp nearest neighbor peak of the binary mixture along with the next (colloid-separated) nearest neighbor maximum are very well reproduced, but the intermediate structure due to the small particles is missing. To further elucidate the one-component system structure and compare it with that of the binary mixture we perform the same clustering analysis as before, using the same distance $r_s$. This yields the cluster population shown in Fig. 4, which agrees well with the fully modeled mixture result and exhibits a rather clear exponential decay at large cluster sizes. Somewhat surprisingly, the dimensionality of the clusters so obtained is also in very good agreement with that of the clusters encountered in the two-component simulation of the mixture. This appears to cast some doubt on the role played by hydrodynamic interactions in the formation of open structures \cite{19}.

The present MD results and analysis suggest that in strongly asymmetric binary mixtures exclusion-like interactions alone can give rise to highly structured, so-called cluster phases. For the binary mixture studied here as a paradigm of a colloidal suspension the clustering kinetics is dominated by two-body coagulation and fragmentation processes. This may be expected to hold with increasing accuracy as the colloidal volume fraction decreases. The resulting colloidal clusters are tenuous, open structures that can be classified as randomly branched polymers, which is largely a consequence of the very short range of the induced inter-colloidal attractions and not hydrodynamic interactions. In the final state the clusters are exponentially distributed in size for both the fully modeled mixture and an equivalent effective one-component fluid. This is consistent with single-phase equilibrium of the mixture close to a two-phase region of its phase diagram. The observed cluster 'phase' is then simply a signature of equilibrium phase fluctuations, controllable for example by temperature and pressure changes. This may also be the case with other, more standard examples of colloids where clusters form through reversible aggregation and are quite small in size \cite{7}. When considered together with the observed properties of the aggregation kinetics these features lend support to the idea \cite{24} that the relaxation dynamics in such systems is well described as coagulation-fragmentation kinetics of phase fluctuations.

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\cite{1} For a recent review see, e.g., V.J. Anderson, H.N.W. Lekkerkerker, Nature 416, 811 (2002).

\cite{2} A. Yethiraj, A. van Blaaderen, Nature 421, 513 (2003).
[3] I.M. de Schepper, H.E. Smorenburg, E.G.D. Cohen, Phys. Rev. Lett. 70, 2178 (1993).
[4] D.T. Wasan, A.D. Nikolov, Nature 423, 156 (2003).
[5] Y. Han, D.G. Grier, Phys. Rev. Lett. 91, 038302 (2003).
[6] S. Auer, D. Frenkel, Phys. Rev. Lett. 91, 015703 (2003).
[7] A. Stradner, H. Sedgwick, F. Cardinaux, W.C.K. Poon, S.U. Egelhaaf, P. Schurtenberger, Nature 432, 492 (2004).
[8] J. Liu, E. Luijten, Phys. Rev. Lett. 93, 247802 (2004).
[9] M.D. Haw, Phys. Rev. Lett. 92, 185506 (2004).
[10] A.M. Puertas, M. Fuchs, M.E. Cates, J. Chem. Phys. 121, 2813 (2004).
[11] S. Manley et al., Phys. Rev. Lett. 93, 108302 (2004).
[12] M. Schmidt, M. Dijkstra, J.-P. Hansen, Phys. Rev. Lett. 93, 088303 (2004).
[13] J. Forsman, C.E. Woodward, Phys. Rev. Lett. 94, 118301 (2005).
[14] P.N. Segré, V. Prasad, A.B. Schofield, D.A. Weitz, Phys. Rev. Lett. 86, 6042 (2001).
[15] S. Asakura, F. Oosawa, J. Chem. Phys. 22, 1255 (1954).
[16] F. Sciortino, S. Mossa, E. Zaccarelli, P. Tartaglia, Phys. Rev. Lett. 93, 055701 (2004).
[17] A.I. Campbell, V.J. Anderson, J.S. van Duijneveldt, P. Bartlett, Phys. Rev. Lett. 94, 208301 (2005).
[18] M. Dijkstra, R. van Roij, R. Evans, Phys. Rev. E 59, 5744 (1999).
[19] H. Tanaka, T. Araki, Phys. Rev. Lett. 85, 1338 (2000).
[20] W.G. Hoover, M. Ross, K.W. Johnson, D. Henderson, J.A. Barker, B.C. Brown, J. Chem. Phys. 52, 4931 (1970).
[21] M.J. Nuevo, J.J. Morales, D.M. Heyes, Phys. Rev. E 58, 5845 (1998).
[22] G. Jackson, J.S. Rowlinson, F. van Sol, J. Phys. Chem. 91, 4907 (1987).
[23] F. Family, P. Meakin, J.M. Deutch, Phys. Rev. Lett. 57, 727 (1986).
[24] C.M. Sorensen, H.X. Zhang, T.W. Taylor, Phys. Rev. Lett. 59, 363 (1987).
[25] R.D. Vigil, R.M. Ziff, Phys. Rev. Lett. 61, 1431 (1988).
[26] P.J. Blatz, A.V. Tobolsky, J. Phys. Chem. 49, 77 (1945).
[27] D.A. Weitz, M. Oliveria, Phys. Rev. Lett. 52, 1433 (1984).
[28] G. Parisi, N. Sourlas, Phys. Rev. Lett. 46, 871 (1981).
[29] M.E. Cates, S.J. Candau, J. Phys.: Condens. Matter 2, 6869 (1990).
[30] See, e.g., L.D. Landau, E.M. Lifshitz, Statistical Physics, 3rd edition, (Butterworth-Heinemann, Oxford, 1980).
FIG. 1: Time evolution of the (normalized) number of monomers - $n_1$ (bottom dotted line), inverse of the (normalized) total number of clusters - $s_1$ (top dotted line), and $n_1 s_1^2$ (middle dotted line); solid line is an exponential relaxation fit; $t_0 = \sigma(m/\epsilon)^{1/2}$. Inset: characteristic cluster size $s$ (see text) versus $s_1$.

FIG. 2: A cluster of 33 colloidal particles.
FIG. 3: Radius of gyration $R_g$ (see text) as a function of cluster size $k$: from simulations (dashed line) and power law ($R_g \propto k^{1/D_f}$) fit (solid line); $D_f = 2$. 
FIG. 4: Cluster size distribution from MD simulations of the binary mixture (diamonds), simulations of the one-component fluid (dotted line) and exponential decay fit (solid line).
FIG. 5: Colloid-colloid pair correlation function of the binary mixture (solid line) and effective one-component system (dotted line). Inset: interaction potential $V_{\text{eff}}$ (see text) of the one-component system; $\beta = 1/k_B T$. 