

# Exact methods for Campi plots

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## Abstract

We introduce for canonical fragmentation models an exact method for computing expectation values which exclude the largest cluster. This method allows for the computation of the reduced multiplicity and other quantities of interest introduced by Campi, and a comparison shows that the percolation model and a recent canonical model differ mostly only in small respects in these ensemble averages.

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Campi and Krivine [1,2] introduced a method for distinguishing fragmentation models from one another. By comparing various expectation values in which the largest cluster is excluded but the particle number and fragment multiplicity are held fixed, they showed that the percolation model has a distinctly different behavior than many competing nuclear fragmentation models.

In this paper, we analyze another statistical weight we have been using recently and show that it shares many of the same properties as percolation theory, a point already apparent from a consideration of critical exponents [3]. The method used in this paper is unusual in that it is an exact computational method. Monte Carlo sampling is avoided by exploiting some properties of the partition functions, and as such there are no statistical errors, an immense improvement over earlier methods.

We begin by assuming that each fragmentation outcome happens with a probability proportional to the Gibbs weight [4–7]

$$W(\vec{n}) = \prod_{k \geq 1} \frac{x_k^{n_k}}{n_k!}, \quad (1)$$

where  $n_k$  is the number of fragments of size (or charge)  $k$ , and  $x_k$  is a parameter associated with  $k$  sized fragments. We then define the microcanonical partition function as

$$Z_A^{(m)}(\vec{x}) = \sum_{\pi_m(A)} W(\vec{n}), \quad (2)$$

where  $\pi_m(A)$  is the set of partitions of  $A$  nucleons into  $m$  fragments, i.e.  $\sum_k k n_k = A$ ,  $\sum_k n_k = m$ . These partition functions satisfy the identity

$$\frac{\partial Z_A^{(m)}}{\partial x_k} = Z_{A-k}^{(m-1)}(\vec{x}), \quad (3)$$

which allows for the computation of the partition functions recursively from  $\sum_k \langle n_k \rangle = m$ , since

$$\langle n_k \rangle = x_k \frac{\partial}{\partial x_k} \ln Z_A^{(m)} = x_k \frac{Z_{A-k}^{(m-1)}}{Z_A^{(m)}}. \quad (4)$$

Campi defined reduced moments as moments in which the largest cluster is excluded from the measure, i.e.

$$M_s(\vec{n}) \equiv \sum_k k^s n_k - k_{\max}^s \quad (5)$$

where  $k_{\max}$  is the size of the largest cluster. This suggested the definition of the reduced variance  $\gamma_2$  [1,2] for a fragmentation event should be

$$\gamma_2(\vec{n}) = \frac{M_2 M_0}{M_1^2} = (m-1) \frac{\sum_k k^2 n_k - k_{\max}^2}{(A - k_{\max})^2}, \quad (6)$$

Its expectation value can be computed by breaking events into classes specified by  $k_{\max}$ , and summing the expectation values over those classes with the appropriate weight, i.e.

$$\langle \gamma_2 \rangle = (m-1) \sum_{k_{\max}} \Pr(k_{\max}) \frac{\sum_k k^2 \langle n_k \rangle(k_{\max}) - k_{\max}^2}{(A - k_{\max})^2}, \quad (7)$$

where  $\Pr(k_{\max})$  is the probability of  $k_{\max}$  being the largest cluster size and  $\langle n_k \rangle(k_{\max})$  is the expectation value of  $n_k$  when  $k_{\max}$  is fixed.

To compute expectation values in which the largest cluster size is fixed we need to compute the partition function for such ensembles. Clearly this partition function is given by all the terms in the microcanonical partition function which have  $x_{k_{\max}}$  as the highest  $x_k$  in the term. Consider

$$\begin{aligned} \Delta Z_A^{(m)}(k_{\max}) &\equiv Z_A^{(m)}(x_1, \dots, x_{k_{\max}}, 0, \dots, 0) \\ &\quad - Z_A^{(m)}(x_1, \dots, x_{k_{\max}-1}, 0, \dots, 0). \end{aligned} \quad (8)$$

We see that  $\Delta Z_A^{(m)}(k_{\max})$  is the partition function for ensembles with fixed maximum cluster size  $k_{\max}$ , since the first term collects all terms with  $x_{k_{\max}}$  or lower, and the second term eliminates those terms which don't have an  $x_{k_{\max}}$ . From this result we can determine  $\Pr(k_{\max})$  and  $\langle n_k \rangle(k_{\max})$ , which are given by

$$\Pr(k_{\max}) = \frac{\Delta Z_A^{(m)}(k_{\max})}{Z_A^{(m)}(x_1, \dots, x_A)} \quad (9)$$

$$\langle n_k \rangle(k_{\max}) = \begin{cases} 0 & k > k_{\max} \\ \frac{Z_{A-k}^{(m-1)}(x_1, \dots, x_{k_{\max}}, 0, \dots, 0)}{\Delta Z_A^{(m)}(k_{\max})} & k = k_{\max} \\ \frac{\Delta Z_{A-k}^{(m-1)}(k_{\max})}{\Delta Z_A^{(m)}(k_{\max})} & k < k_{\max} \end{cases} \quad (10)$$

This method is quite general and can be applied to other models. For example, equipartitioning models, which have weights given by

$$W(\vec{n}) = \prod_{k \geq 1} x_k^{n_k} \quad (11)$$

can also be analyzed by this method with some minor modifications. For example,  $x_k = 1$  is the model used by Sobotka and Moretto [8].

With these identities there is sufficient information to compute  $\langle \gamma_2 \rangle$  and other reduced moments for any  $x_k$ . We use  $x_k = x/k^\tau$  for a variety of reasons discussed elsewhere [9]. Campi and Krivine [2] following Mekjian [4] considered this model with  $\tau = 1.0$  and showed that its reduced variance and other related expectation values had a distinctly different behavior than the percolation model. Plotting the expected reduced variance vs.  $(m-1)/A$ ,  $\langle \gamma_2 \rangle(m)$  has a single peak. The location, height and width of this peak for the two models (and other models they considered) are completely different, suggesting the usefulness of this plot in distinguishing fragmentation models. The choice  $\tau = 0$  was considered by Gross, et. al. [10–12], and a different model was analyzed by Pan and Das Gupta [13].

Since that time our interest has turned to the choice  $\tau = 2.5$  because of similarities with percolation theory and Bose condensation. Namely, the sudden appearance of an infinite cluster in the infinite  $A$  limit and the presence of condensation phenomena. As such, we have recomputed the Campi plots for this model and have discovered that they duplicate the percolation model results in many respects. Figure 1(a) shows the results. The height and location of the peak of the reduced variance  $\langle \gamma_2 \rangle$  are the same in both models. The only significant difference is the width of the peak which is larger in the Gibbs model than in the percolation model. Plots of  $\langle k_{\max} \rangle(m)$  vs.  $(m-1)/A$  given in Fig. 1(b) are also very similar for both models, and the scaling behavior of the position, width and height with changing  $A$  also agree.

Another plot suggested by Campi [14] is to divide the event space by the maximum cluster size of the event  $k_{\max}$  and the reduced second moment  $M_2$  and plot the probability of the canonical model being at any particular point on the graph. This can be also be done

exactly for Gibbs models in a way completely analogous to the way given above. Define the partition function  $Z_A(m_2; \vec{x})$  as the sum of the Gibbs weight Eq. (1) over all partition vectors  $\vec{n}$  which satisfy  $\sum_k kn_k = A$ ,  $\sum_k k^2n_k = m_2$ . This can be computed by the following recursion,

$$Z_A(m_2; \vec{x}) = \frac{1}{A} \sum_k kx_k Z_{A-k}(m_2 - k^2; \vec{x}) \quad (12)$$

with  $Z_0(m_2; \vec{x}) = \delta_{m_2,0}$ . If we define  $\Delta Z$  as in Eq. (8), we find again the partition function conditioned on  $k_{\max}$  being fixed, which is proportional to the probability of having an event with both  $m_2$  and  $k_{\max}$ . Figure 2 plots this probability profile as a contour plot, which reveals the events are centered on a particular region in this phase space. The slopes of the edges of this region are related to the critical exponents according to Campi [14].

Clearly there are differences between percolation theory and a Gibbs model, but the differences are not as large as originally suggested by early computations. Indeed the reduced variance might not reliably distinguish percolation from a simple Gibbs model. A different method is needed to distinguish these models. However, the idea of excluding the largest cluster from the ensemble averages is a standard procedure in percolation theory [15], and this new technique for doing that analytically in the Gibbs models shows a particular advantage of these models over percolation models, which we hope will encourage further interest in them.

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## FIGURES

FIG. 1. Expected reduced multiplicity  $\langle\gamma_2\rangle(m)$  (a) and largest cluster size  $k_{\max}/A$  (b) vs.  $(m-1)/A$  for  $\tau = 1.0, 2.5$  and the percolation model at  $A = 125$ .

FIG. 2. Campi probability contour plot for  $Z = 79$ ,  $\tau = 2.5$  at the critical point  $x = x_c$ . The axes are logarithmic, with the largest cluster size on the  $y$ -axis, and the ratio of the reduced moments  $M_2/M_1$  on the  $x$ -axis. The central rings are higher in probability than the outer rings.