



UNITED NATIONS EDUCATIONAL, SCIENTIFIC AND CULTURAL ORGANIZATION
INTERNATIONAL ATOMIC ENERGY AGENCY
INTERNATIONAL CENTRE FOR THEORETICAL PHYSICS
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H4.SMR/994-28

**SPRING COLLEGES IN
COMPUTATIONAL PHYSICS**

19 May - 27 June 1997

**ADVANCED CLASSICAL
MONTE CARLO SIMULATIONS**

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1. Cluster Monte Carlo

2. Histograms

3. Bayesian analysis

Transition matrix MC

4. Simulation of
biological molecules

Dynamically Optimized MC

Emancipation of
computer simulations
from experimental paradigm

Non-physical dynamics
for greater efficiency
- MC , cluster methods

Measurement of quantities
not available to experiment

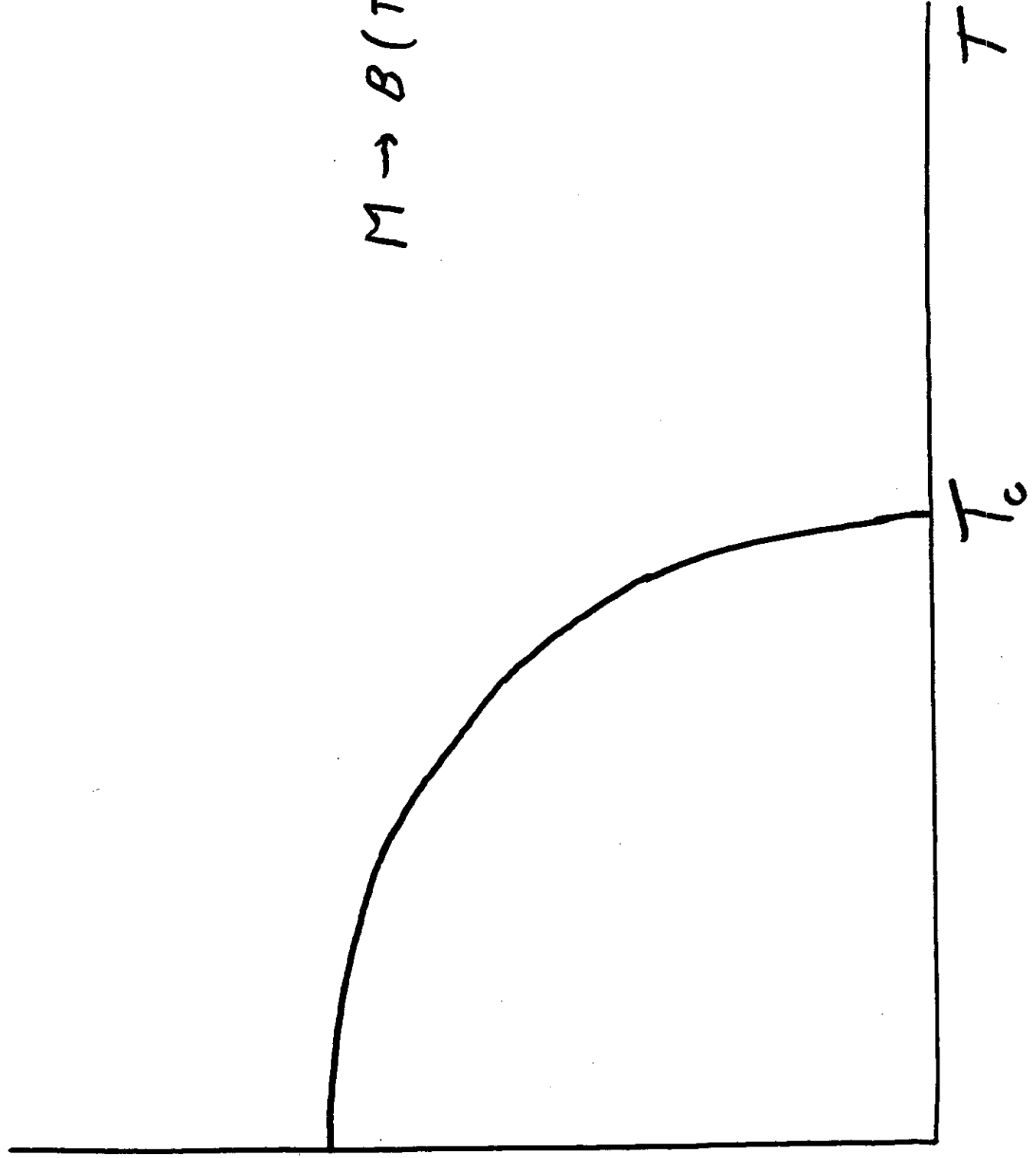
MLRG

Histograms

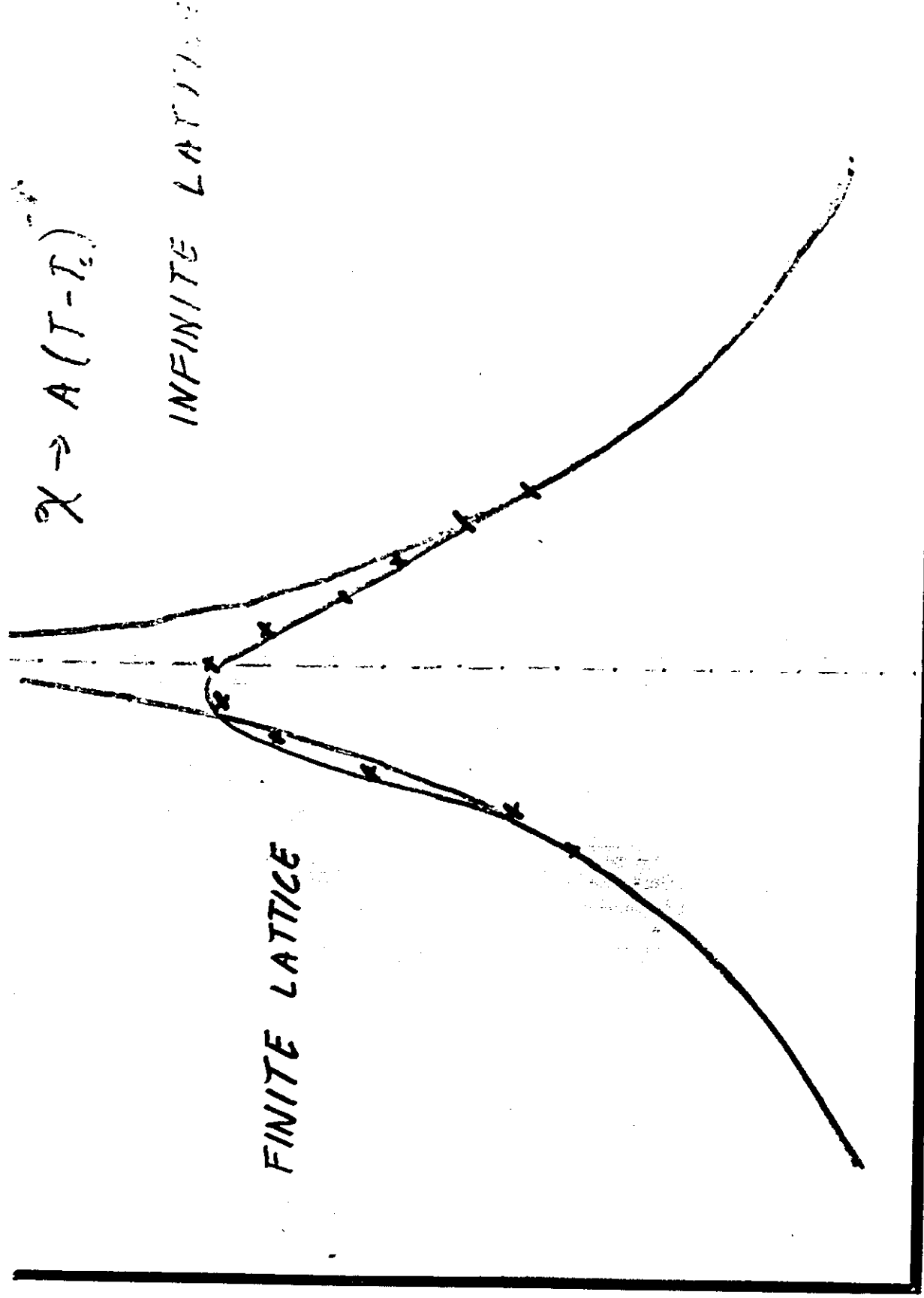
Improved estimators

M

$$M \rightarrow B(T_c - T)^p$$



χ



T_c

T

Monte Carlo Strategy

1) Test your program!

2) Think small

small lattices
short runs
fast scan of parameters
find "interesting" region
vary the initial conditions

3) Concentrate on "interesting" region

small lattices
longer runs (but $\ll$ budget)
calculate errors
estimate errors vs. run time
for larger lattices

4) Increase lattice size

compare
look for size effects
extrapolate
calculate errors

Equilibrium Probability Density

$$P_e(\sigma) = \frac{1}{Z} \exp[-E(\sigma)/k_B T]$$

$$Z = \text{Tr}_{\sigma} \exp[-E(\sigma)/k_B T]$$

$$\langle E \rangle = \text{Tr}_{\sigma} E(\sigma) P_e(\sigma)$$

Markov Process

New state depends on old state

but not

on previous states.

Master Equation:

$$P(\sigma, t+\Delta t) - P(\sigma, t) =$$

$$\sum_{\sigma'} [W(\sigma' \rightarrow \sigma) P(\sigma', t) - W(\sigma \rightarrow \sigma') P(\sigma, t)]$$

$$P(\sigma, t + \Delta t) - P(\sigma, t) =$$

$$\sum_{\sigma'} [W(\sigma' \rightarrow \sigma) P(\sigma', t) - W(\sigma \rightarrow \sigma') P(\sigma, t)]$$

Necessary condition for $P(\sigma, t) \rightarrow P_{eq}(\sigma)$

$$\sum_{\sigma'} [W(\sigma' \rightarrow \sigma) P_{eq}(\sigma') - W(\sigma \rightarrow \sigma') P_{eq}(\sigma)] = 0$$

Stronger condition

Detailed Balance

$$W(\sigma' \rightarrow \sigma) P_{eq}(\sigma') = W(\sigma \rightarrow \sigma') P_{eq}(\sigma)$$

Detailed Balance

$$W(\sigma' \rightarrow \sigma) P_{eq}(\sigma') = W(\sigma \rightarrow \sigma') P_{eq}(\sigma)$$

$$P_{eq}(\sigma) = \frac{1}{Z} \exp[-E(\sigma)/k_B T]$$

$$\frac{W(\sigma' \rightarrow \sigma)}{W(\sigma \rightarrow \sigma')} = \exp[(E(\sigma') - E(\sigma))/k_B T]$$

Theorem for Markov Processes

IF:

- 1) For any two states, σ_A and σ_B , there exists a sequence of states, $\sigma_1 \dots \sigma_n$, such that

$$W(\sigma_A \rightarrow \sigma_1) W(\sigma_1 \rightarrow \sigma_2) \dots W(\sigma_n \rightarrow \sigma_B) \neq 0$$

and

- 2) Detailed Balance

THEN:

$$P(\sigma, t) \xrightarrow[t \rightarrow \infty]{} P_{eq}(\sigma)$$

ISING MODEL

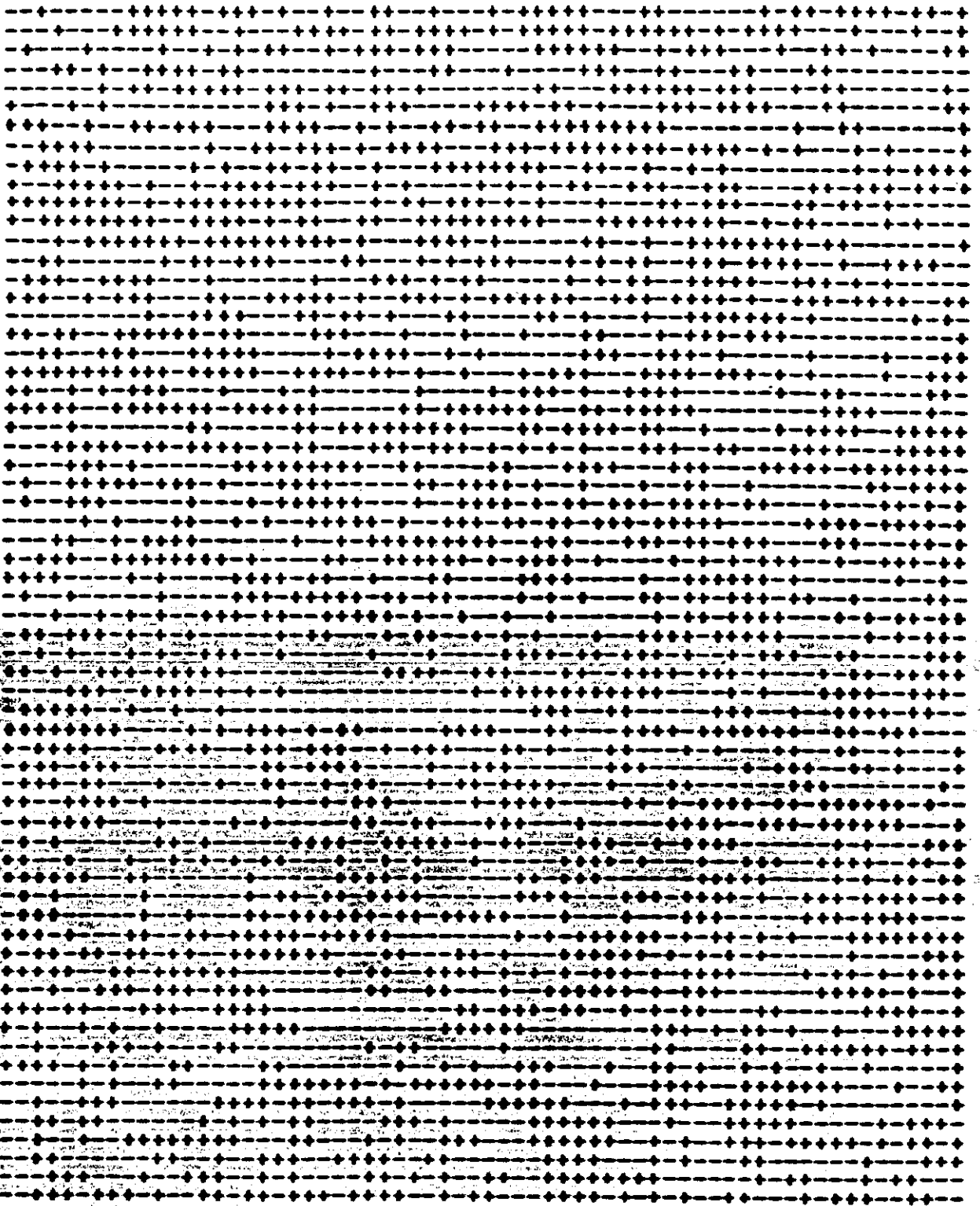
"Spins" σ_i on lattice sites

$$\sigma_i = +1 \quad \text{or} \quad -1$$

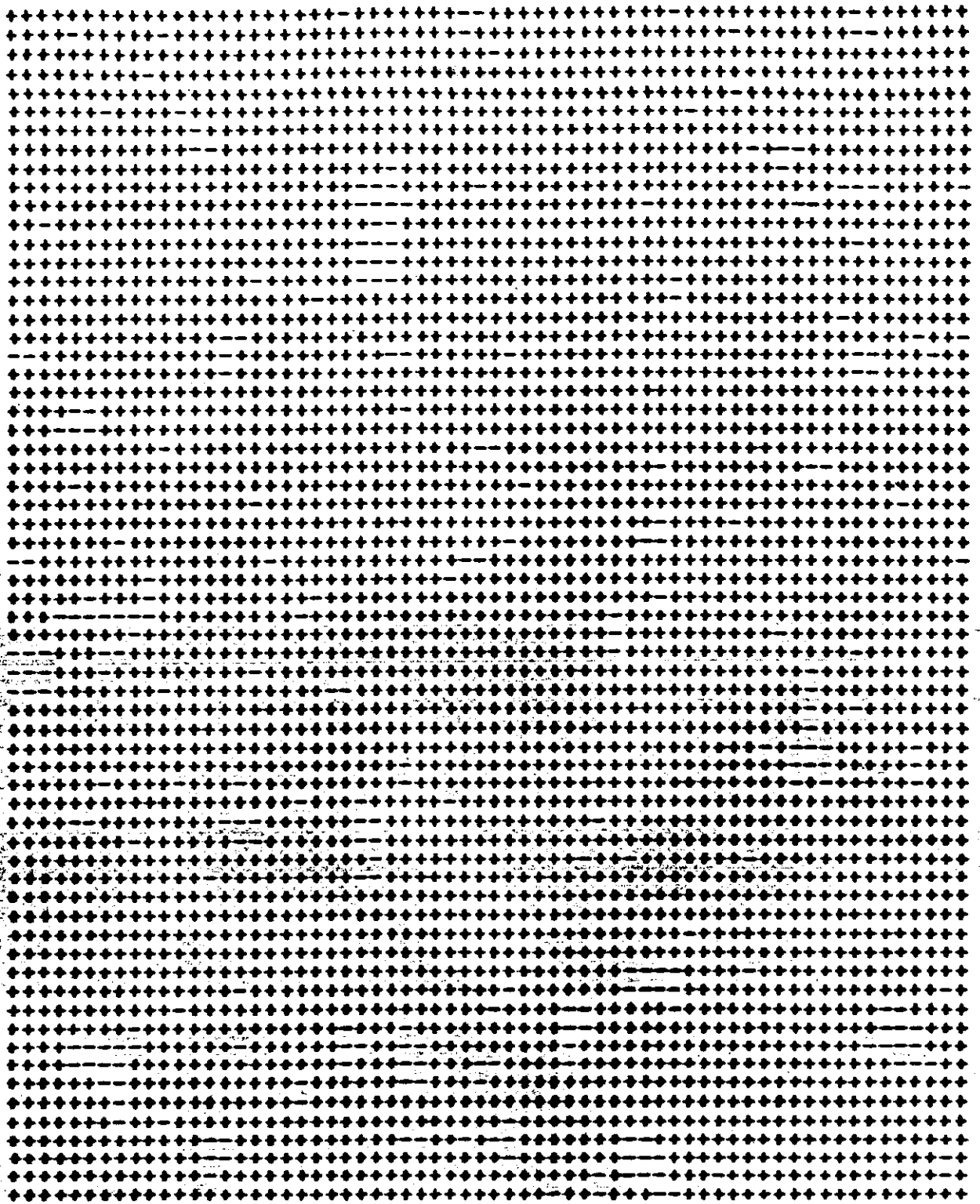
$$H = -J \sum_{\langle ij \rangle} \sigma_i \sigma_j$$

sum on
nearest neighbors

Neighbors		Energy
or	$\begin{array}{cc} + & + \\ - & - \end{array} \}$	$-J$
or	$\begin{array}{cc} + & - \\ - & + \end{array} \}$	$+J$









Critical Slowing Down

$$\tau \sim \xi^z$$

$z \equiv$ dynamical critical exponent

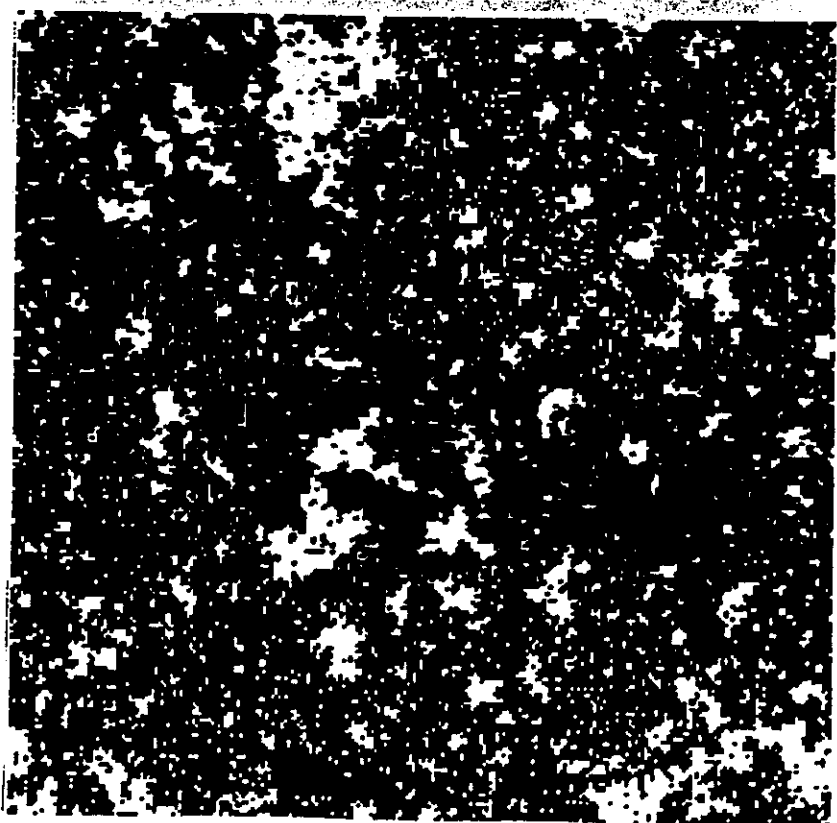
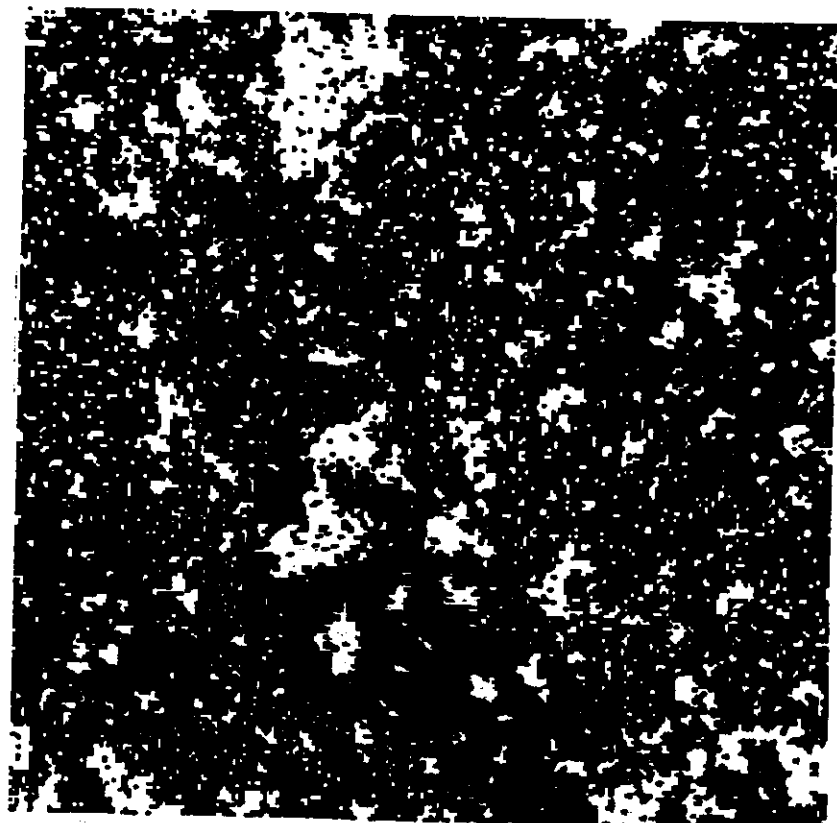
$$\xi \sim |T - T_c|^{-\nu}$$

$$\tau \sim |T - T_c|^{-z\nu}$$

For finite systems $[L \times L]$

Maximum $\xi \sim L$

Maximum $\tau \sim L^z$

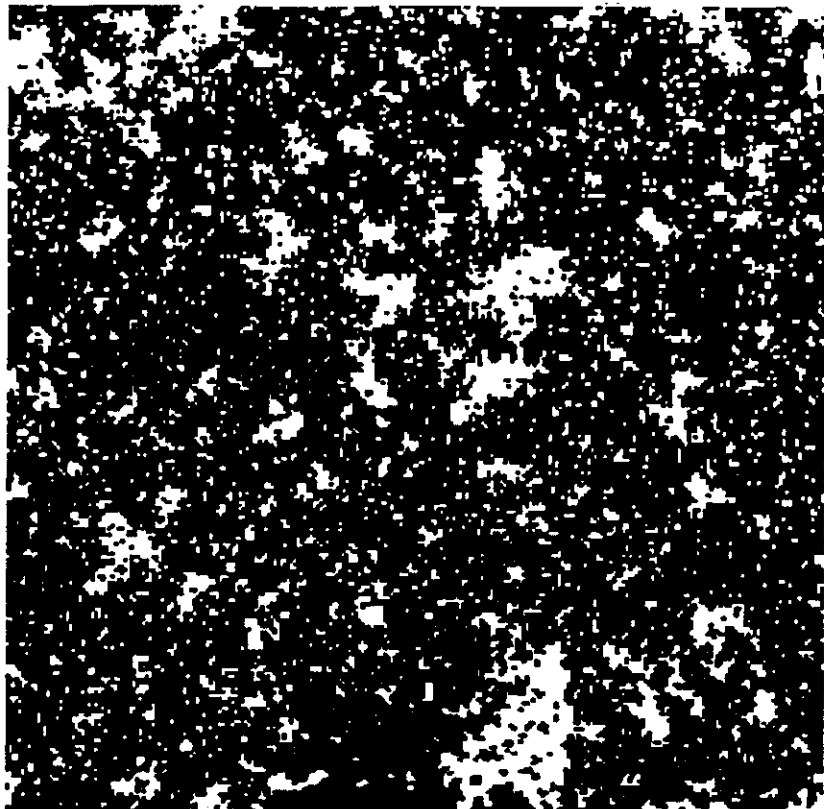


251^h

$T = T_c$

Standard Monte Carlo

3

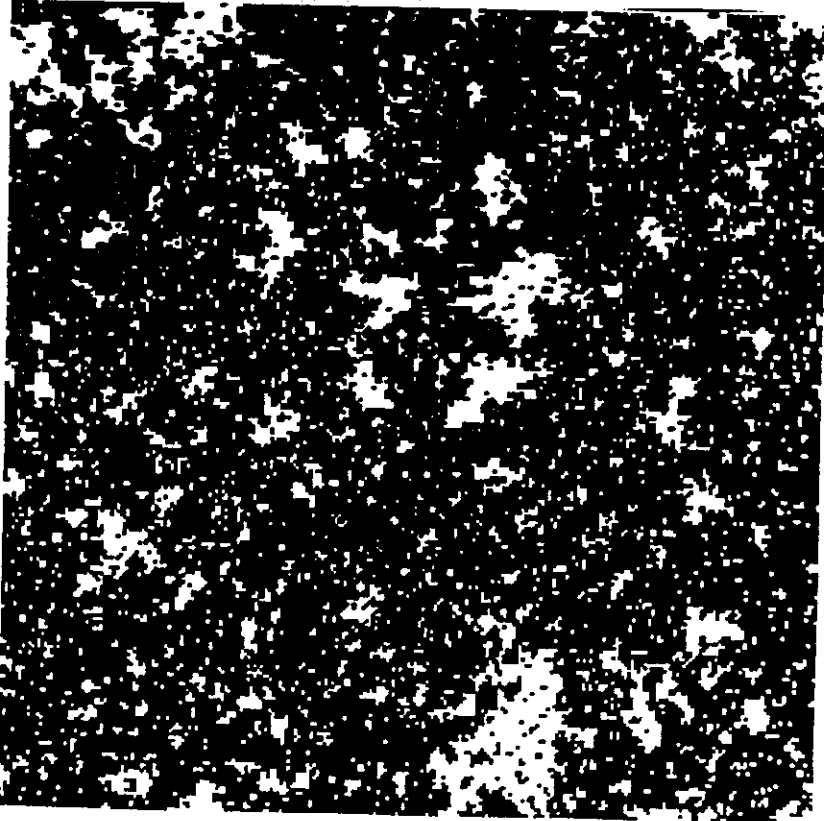


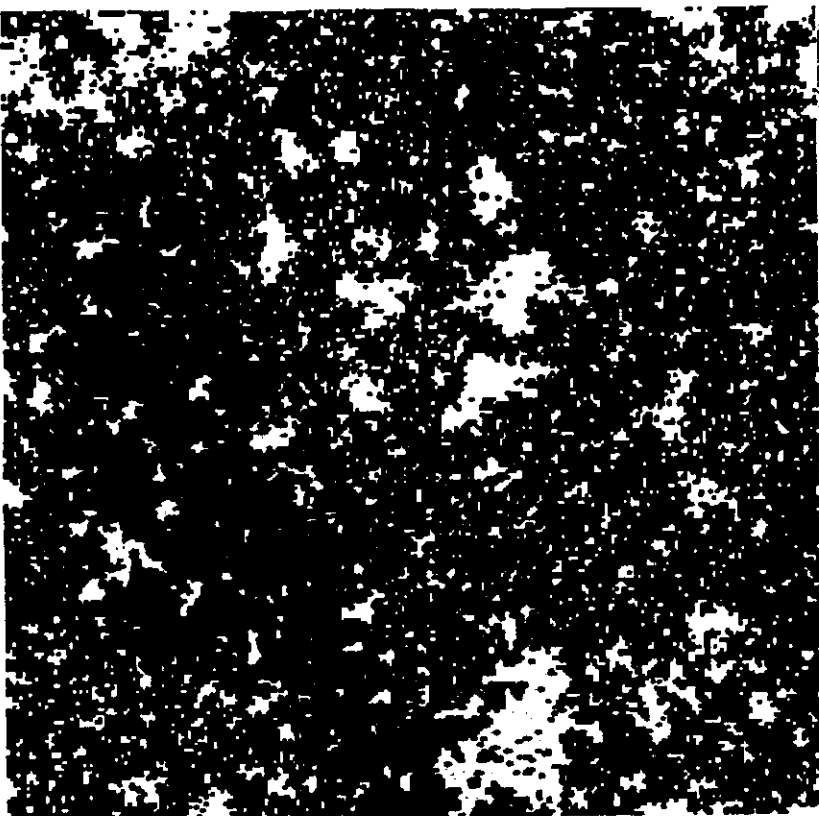
116

T-1

256

4



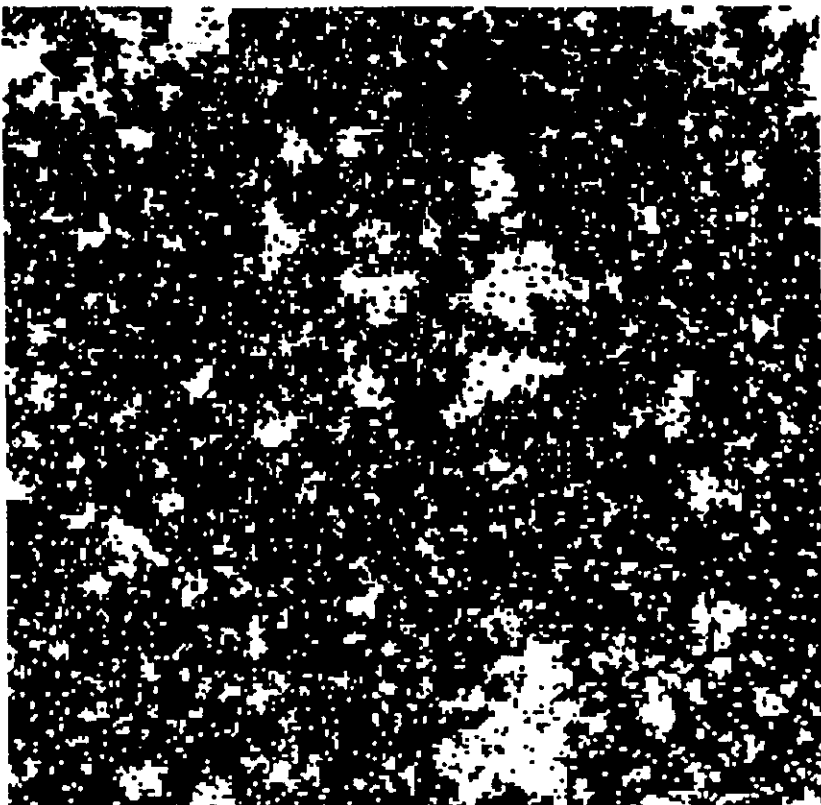


5

176

7-7

251

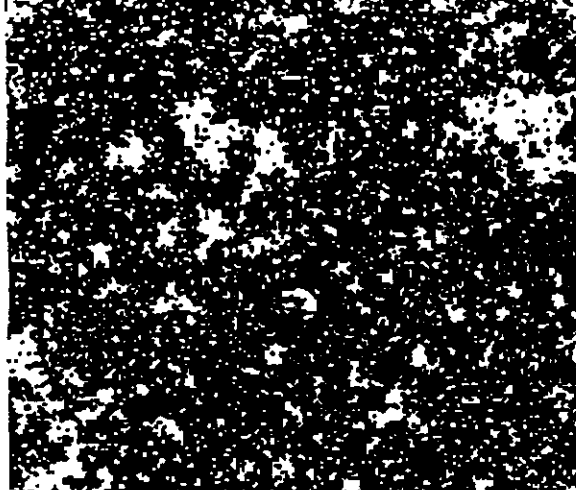


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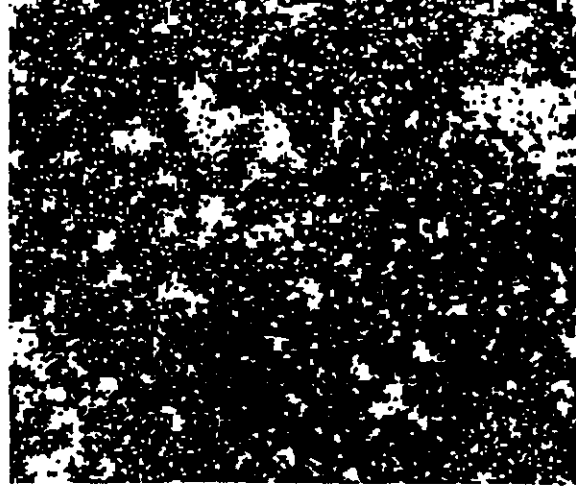
176 5 6

$T = T_c$

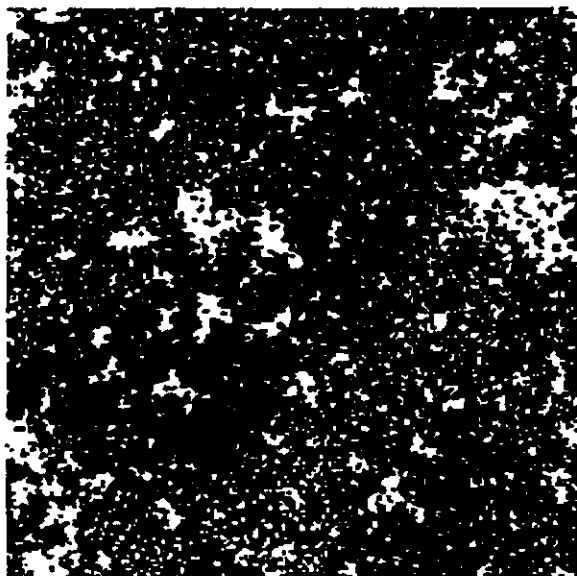
56 x 256



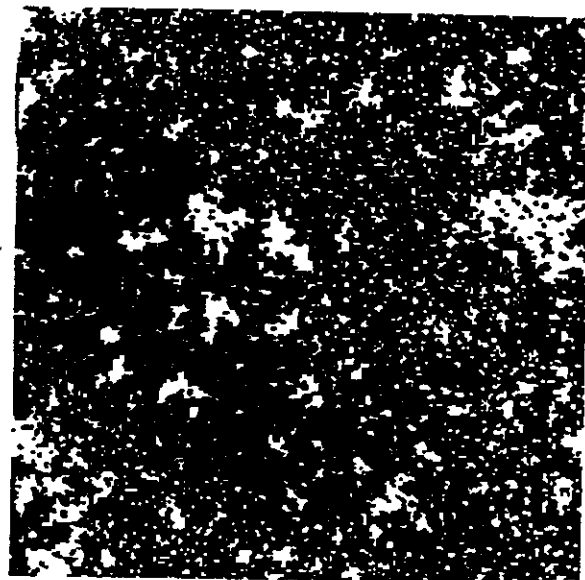
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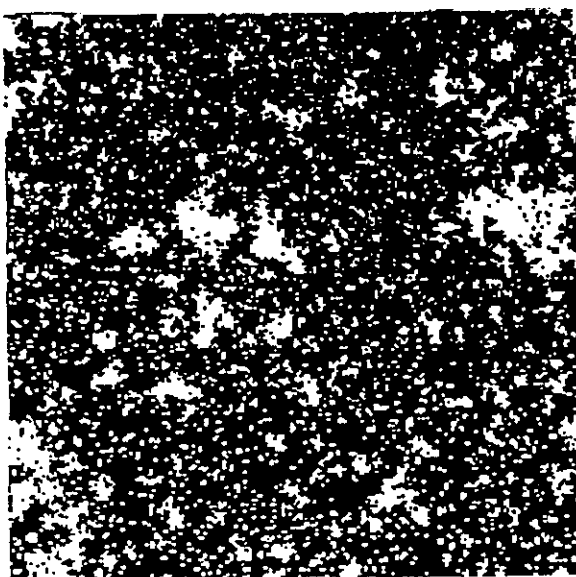
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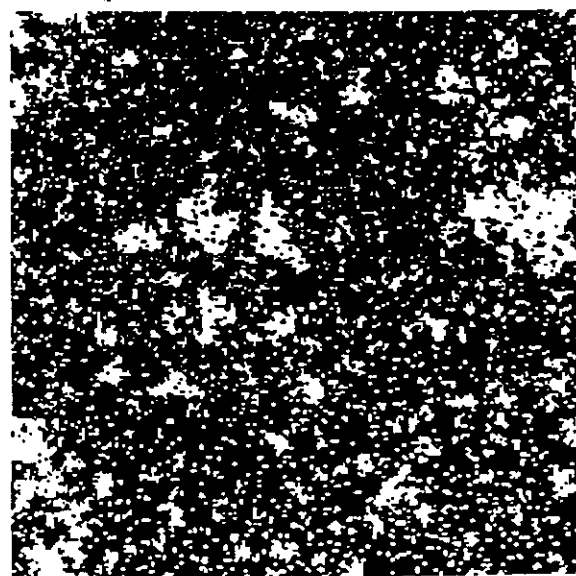
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6

Monte Carlo

Simulation of

Bond Percolation

$X(J,K)$: bond in x-direction
at site (J,K)

$Y(J,K)$: bond in y-direction

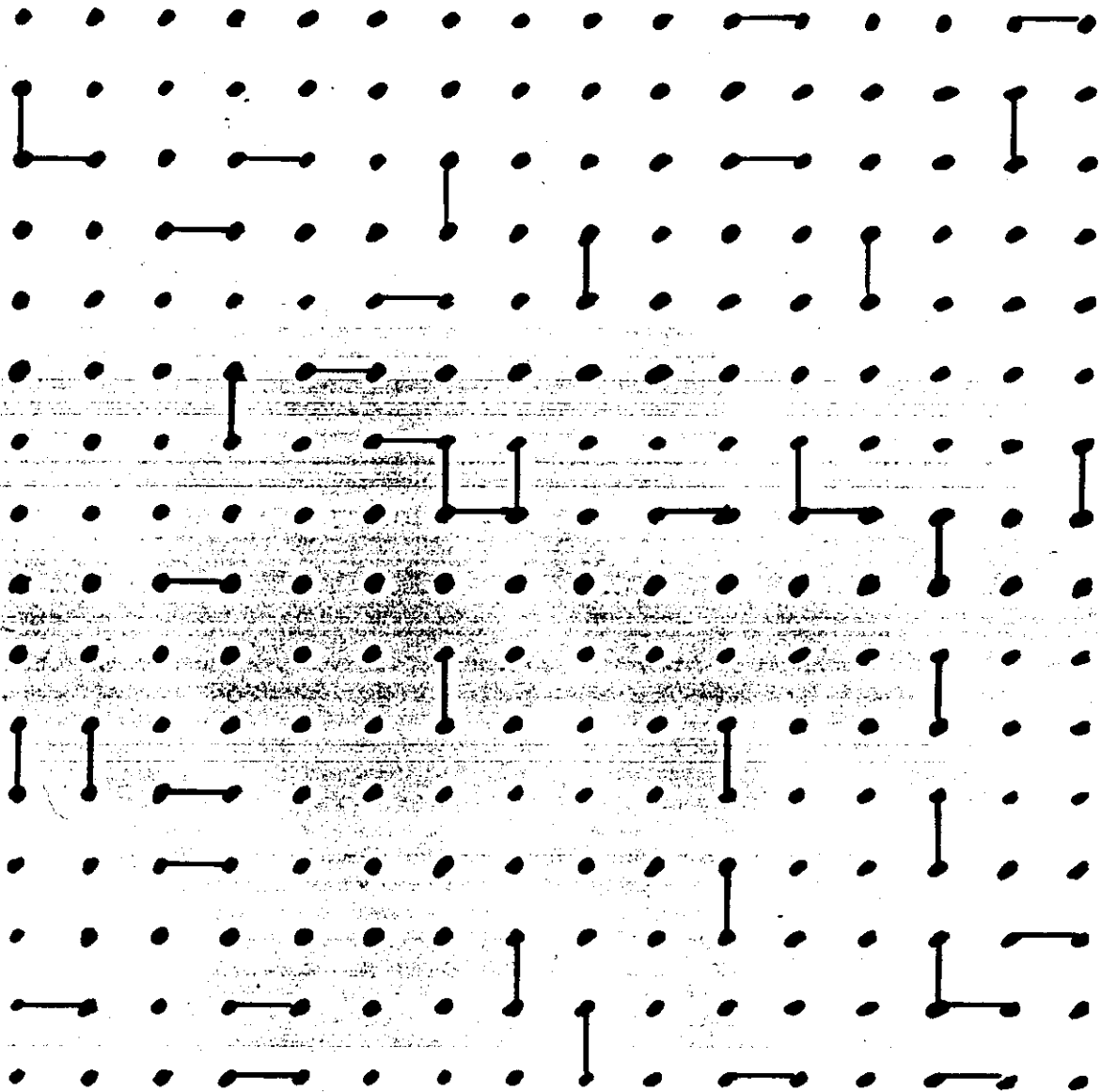
$X = Y = 0 \text{ or } 1$

Essential part of program:

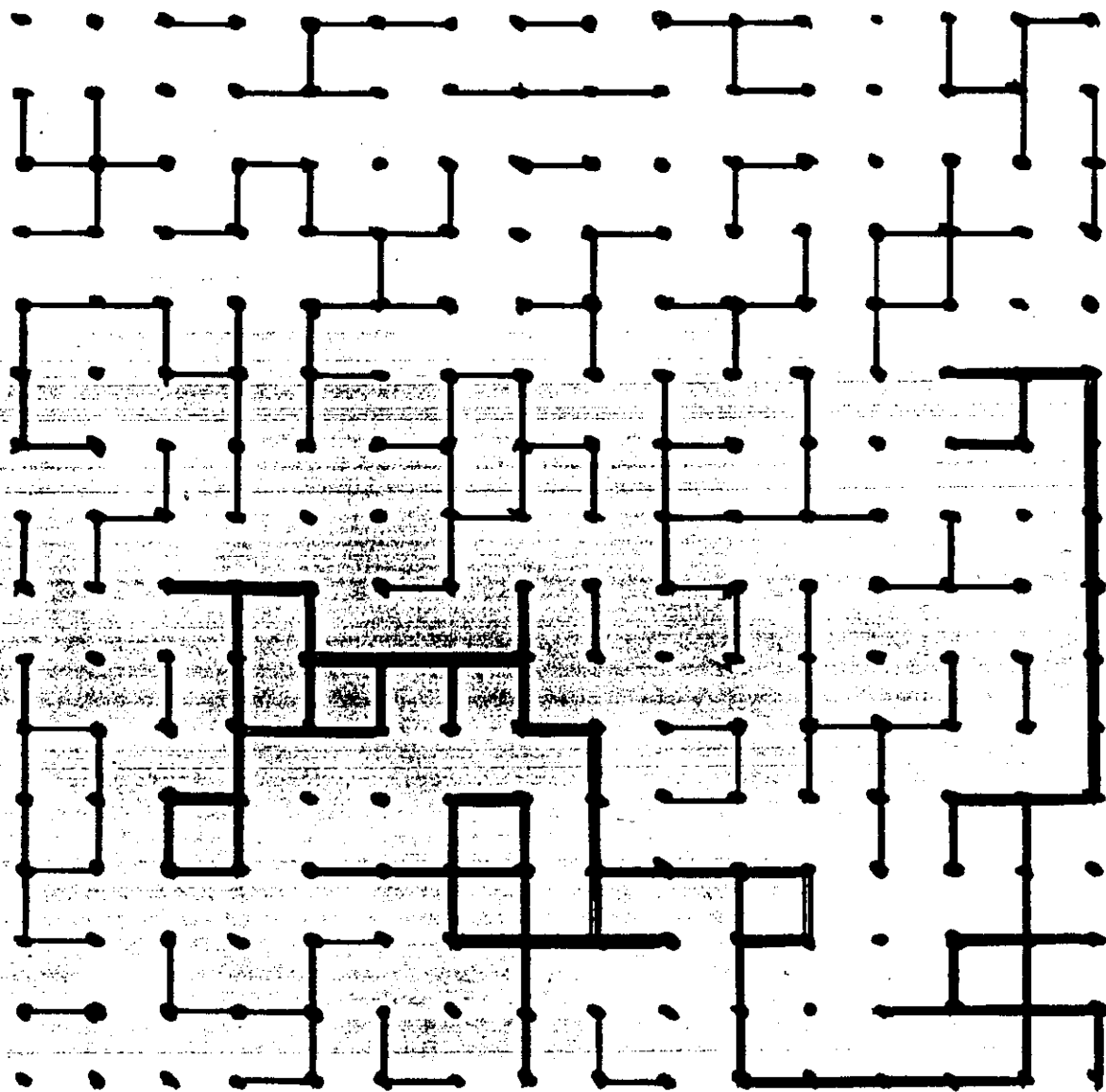
$X(J,K) = 0$

IF(RANX(J,K).LT.P) X(J,K)=1

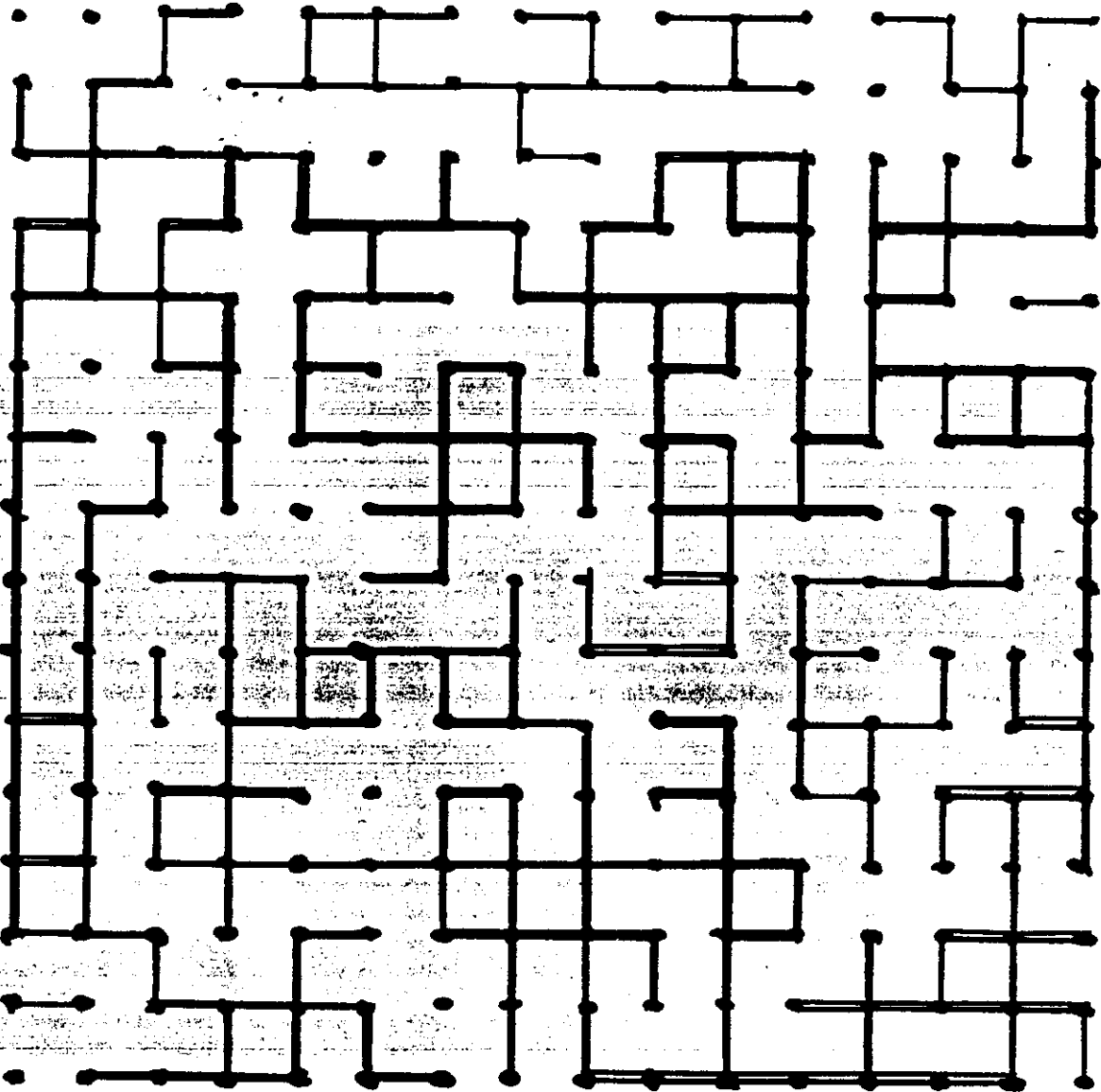
$$p = 0.10$$



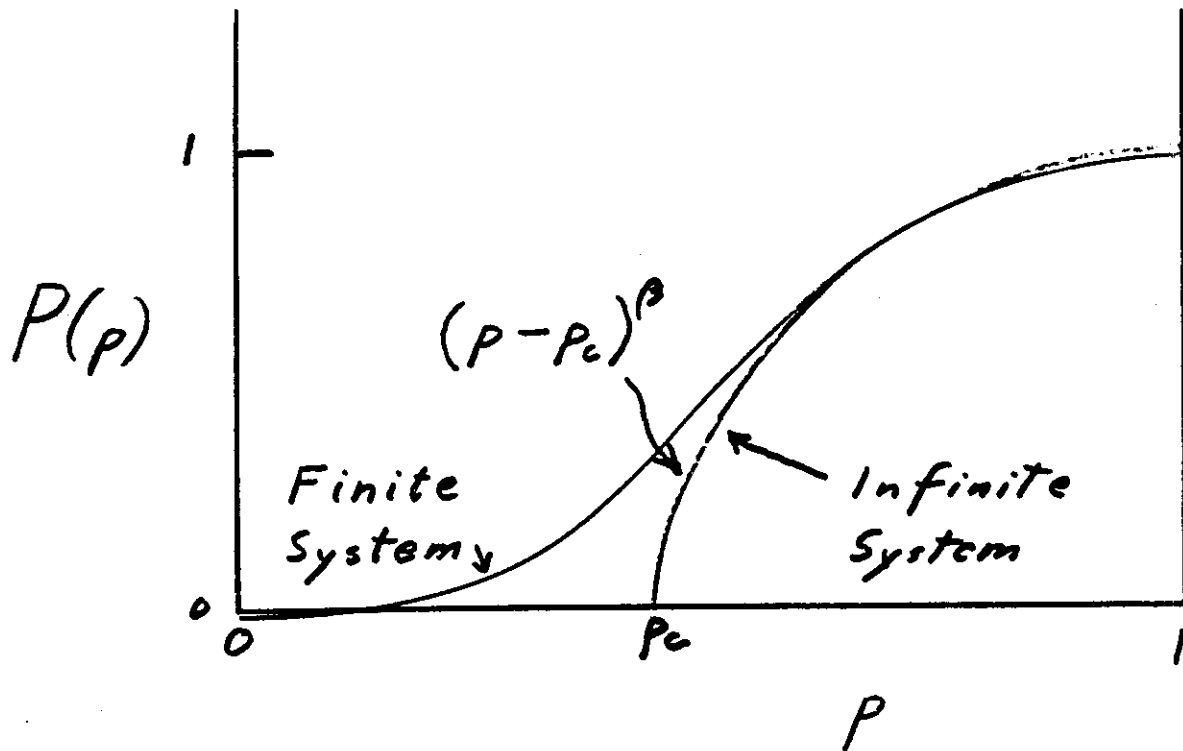
$$p = 0.40$$



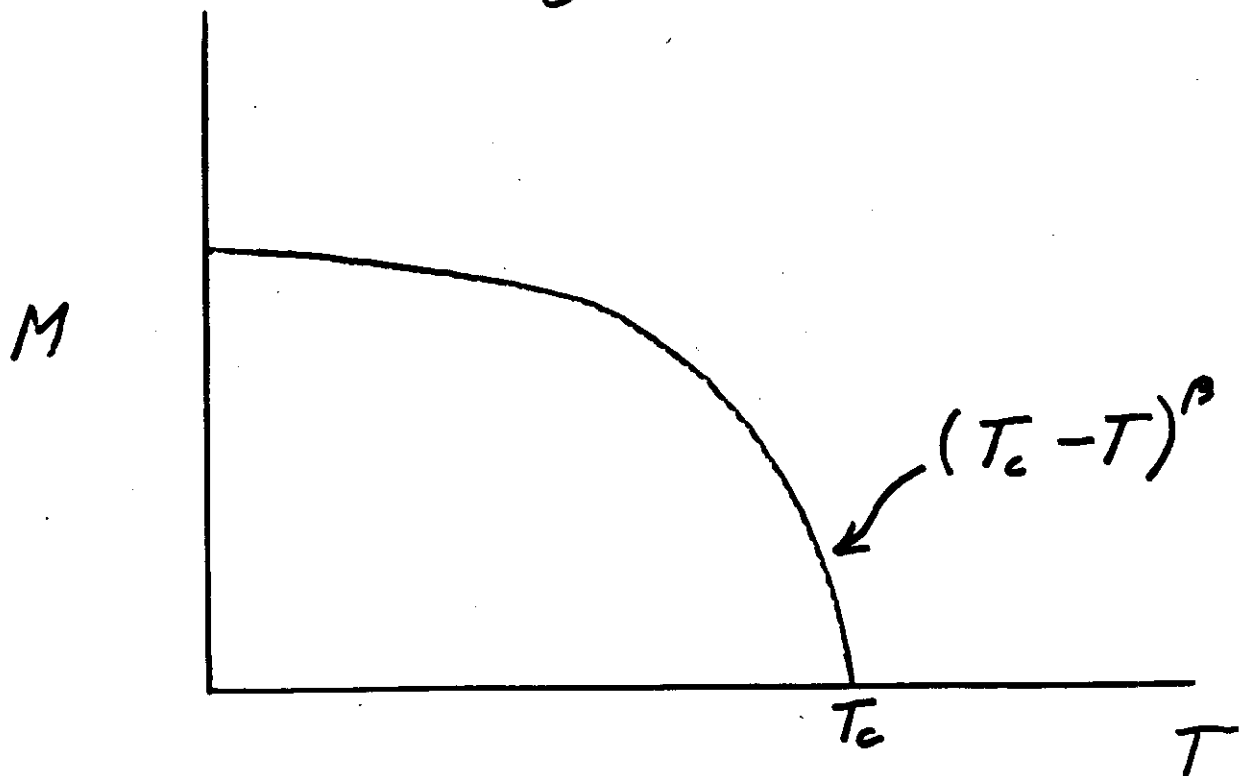
$$p = 0.55$$



Percolation



Magnetization



Ising Model

$$H = K \sum_{\langle i,j \rangle} \delta_{\sigma_i, \sigma_j}$$

$$\sigma_i = +1, -1, \dots$$

or

$$\sigma_i = 1, 2, \dots$$

Potts Model

$$\sigma_i = 1, 2, 3, \dots, q$$

Potts $\rightarrow$ Percolation

$$H = K \sum_{\langle ij \rangle} (\delta_{\sigma_i, \sigma_j} - 1)$$

$$Z = \text{Tr}_{\sigma} e^H$$

$$H_{\langle lm \rangle} = K \sum_{\substack{\langle ij \rangle \\ \neq \langle lm \rangle}} (\delta_{\sigma_i, \sigma_j} - 1)$$

$$Z_{\langle lm \rangle}^{\text{same}} \equiv \text{Tr}_{\sigma} e^{H_{\langle lm \rangle}} \delta_{\sigma_l, \sigma_m}$$

$$Z_{\langle lm \rangle}^{\text{diff}} \equiv \text{Tr}_{\sigma} e^{H_{\langle lm \rangle}} (1 - \delta_{\sigma_l, \sigma_m})$$

$$Z = Z_{\langle lm \rangle}^{\text{same}} + e^{-K} Z_{\langle lm \rangle}^{\text{diff}}$$

Potts $\rightarrow$ Percolation

$$Z = Z_{\langle lm \rangle}^{\text{same}} + e^{-K} Z_{\langle lm \rangle}^{\text{diff}}$$

$$Z_{\langle lm \rangle}^{\text{indep}} \equiv \text{Tr}_{\sigma} e^{H_{\langle lm \rangle}}$$

$$Z_{\langle lm \rangle}^{\text{indep}} = Z_{\langle lm \rangle}^{\text{same}} + Z_{\langle lm \rangle}^{\text{diff}}$$

$$Z = (1 - e^{-K}) Z_{\langle lm \rangle}^{\text{same}} + e^{-K} Z_{\langle lm \rangle}^{\text{indep}}$$

$\uparrow$
 p

$\uparrow$
 $(1-p)$

Potts $\rightarrow$ Percolation

$$Z = p Z_{\langle em \rangle}^{\text{same}} + (1-p) Z_{\langle em \rangle}^{\text{indep}}$$

$$p = 1 - e^{-K}$$

$$Z = \text{Tr}_{\{\text{links}\}} p^b (1-p)^b s^{N_c}$$

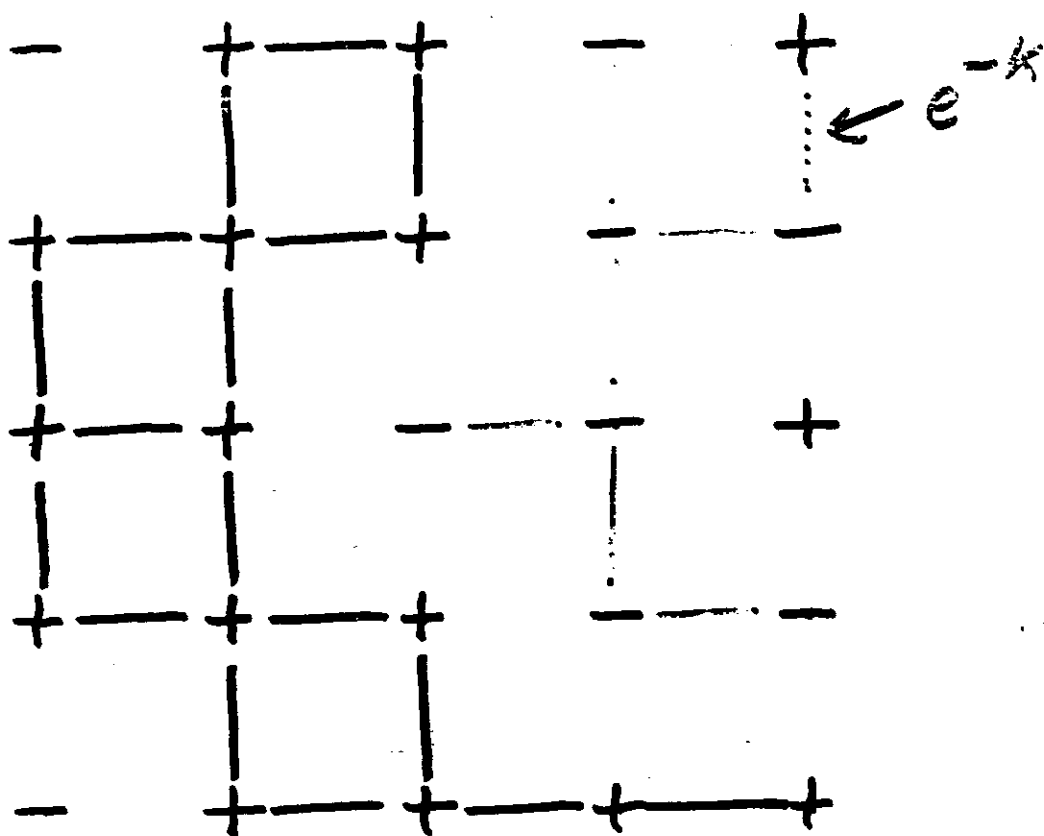
$$Z = \langle s^{N_c} \rangle_{\{\text{links}\}}$$

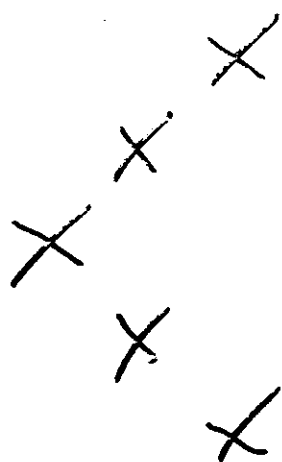
$$\langle N_c \rangle = \lim_{s \rightarrow 1} \left[\frac{\partial}{\partial s} \ln Z \right]$$

Kasteleyn + Fortuin, 1969

-	+	+	-	+
+	+	+	-	-
+	+	-	-	+
+	+	+	-	-
-	+	+	+	+

Potts Clusters





x

x

x

x

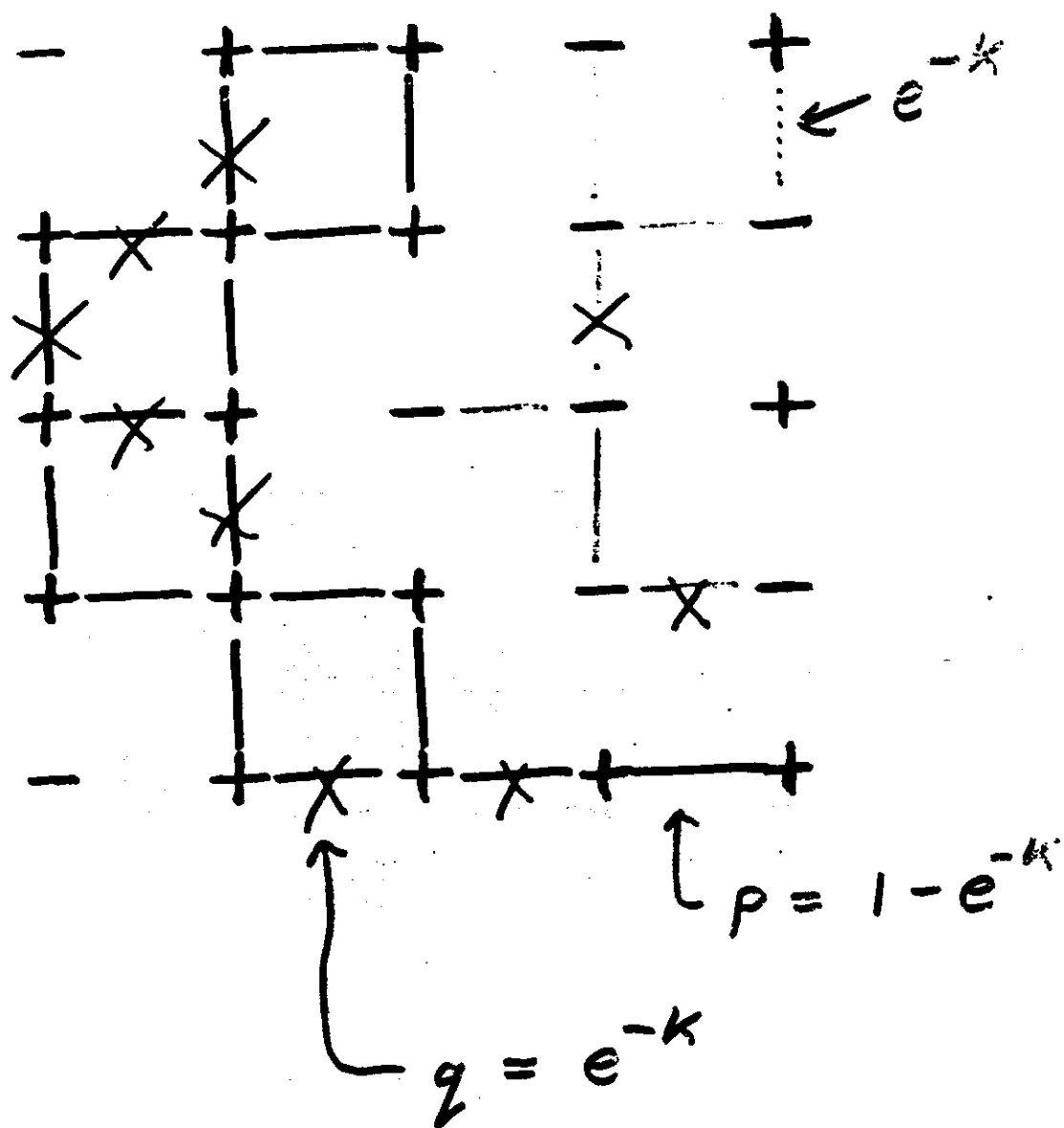
↑

$$p = 1 - e^{-k}$$



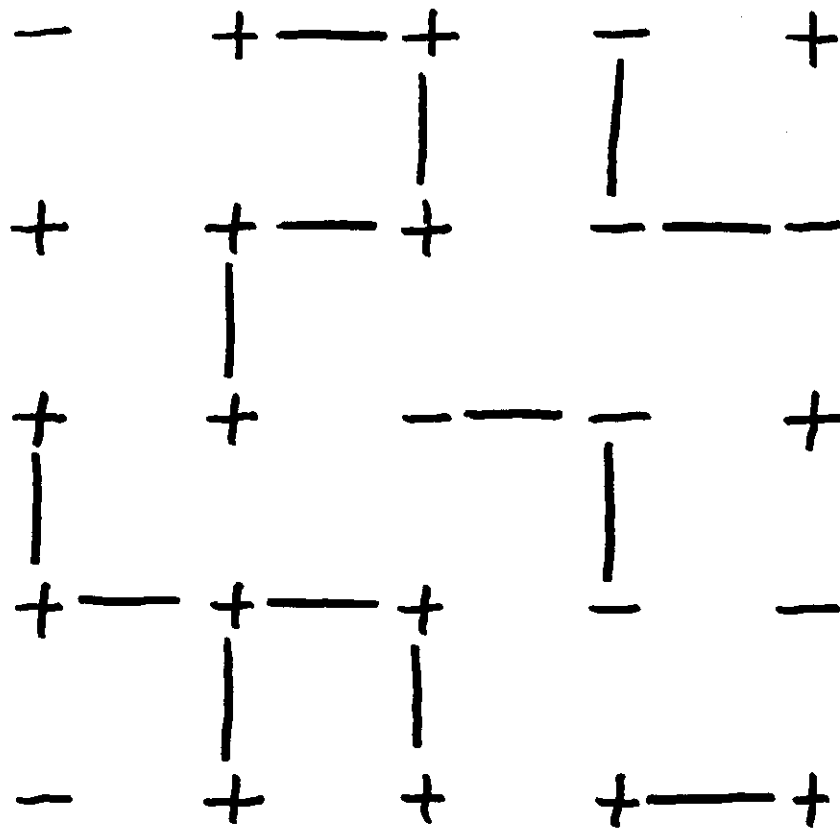
$$z = e^{-k}$$

Potts Clusters



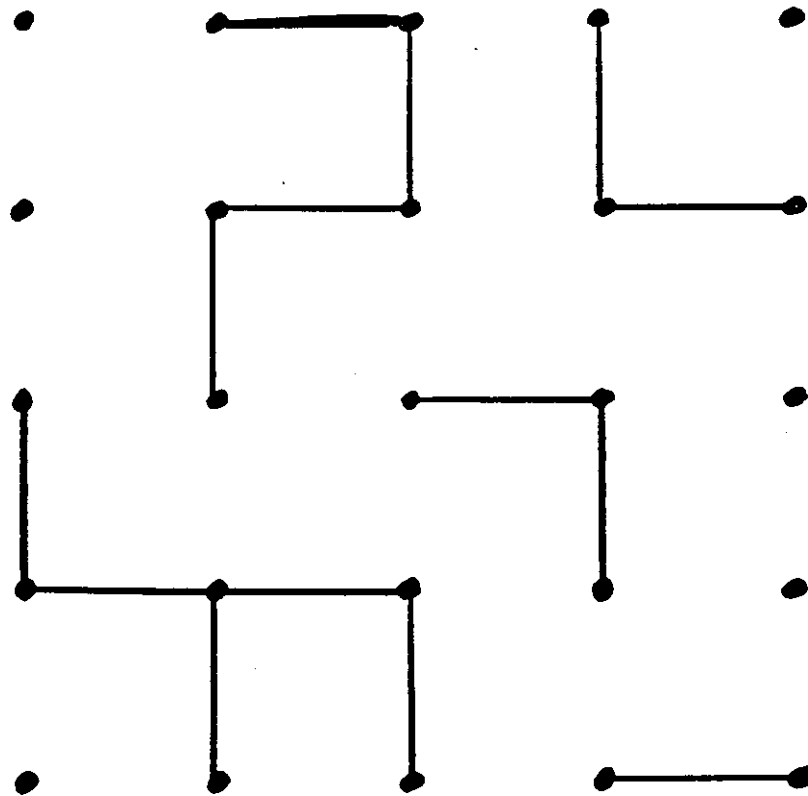
Site - Bond

Correlated Percolation



8 5-0

Percolation Clusters



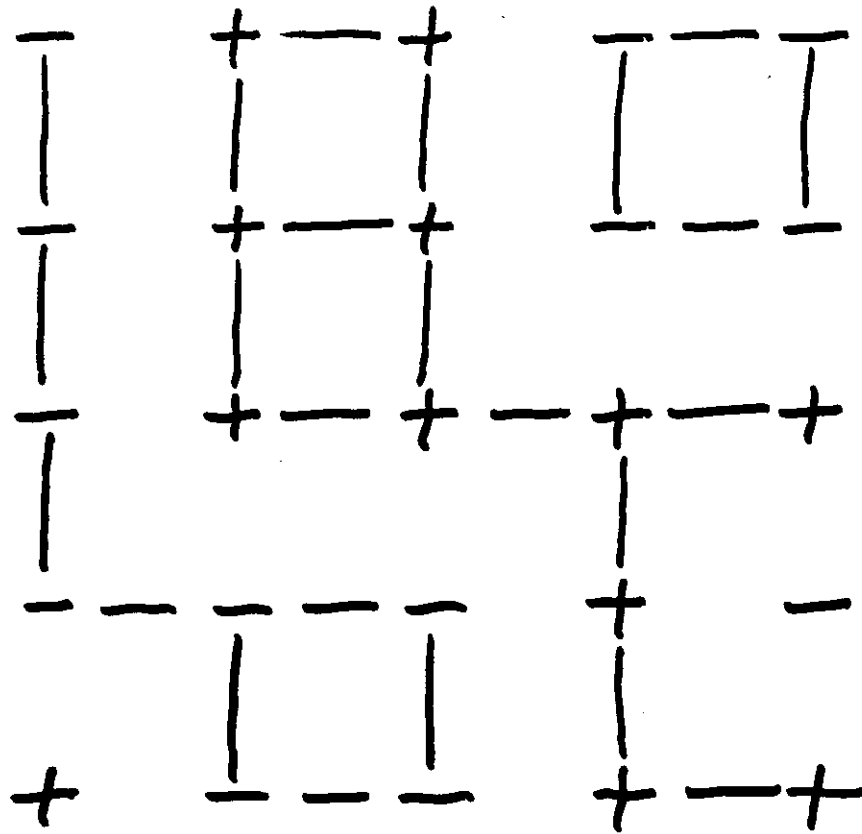
p $q = 1 - p$ s

$s = \# \text{ of Potts states}$

$$q = e^{-K} = 1 - p$$

-	+	+	-	-
-	+	+	-	-
-	+	+	+	+
-	-	-	+	-
+	-	-	+	+

New Potts Clusters



Order Parameter

Ising Ferromagnet

$$M = \lim_{H \rightarrow 0^+} \lim_{L \rightarrow \infty} L^{-d} \left\langle \sum_i \tau_i \right\rangle$$

$$= \lim_{H \rightarrow 0^+} \lim_{L \rightarrow \infty} L^{-d} \left\langle \sum_n C_n \sigma_n \right\rangle$$

$\uparrow$ sum on clusters

$$M = \lim_{H \rightarrow 0^+} \lim_{L \rightarrow \infty} L^{-d} \langle C_{\text{largest}} \rangle$$

Magnetic Susceptibility

$$\chi = L^{-d} \left\langle \sum_{i,j} \sigma_i \sigma_j \right\rangle$$

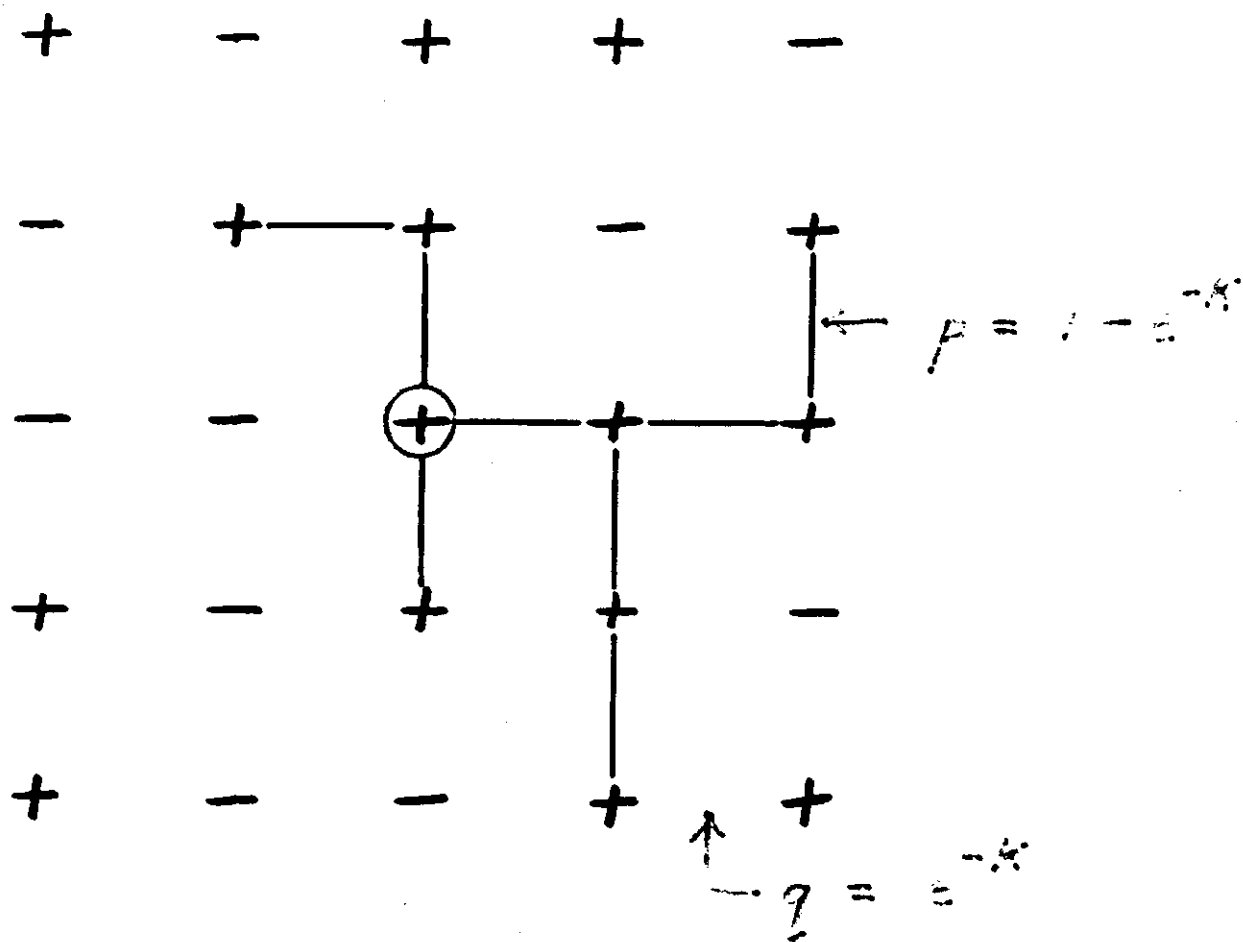
$$\chi = L^{-d} \left\langle \sum_{n,m} C_n C_m \sigma_n \sigma_m \right\rangle$$

$$\langle \sigma_n \sigma_m \rangle = \delta_{n,m}$$

$$\chi = L^{-d} \left\langle \sum_n C_n^2 \right\rangle$$

Single-Cluster Algorithm

U. Wolff

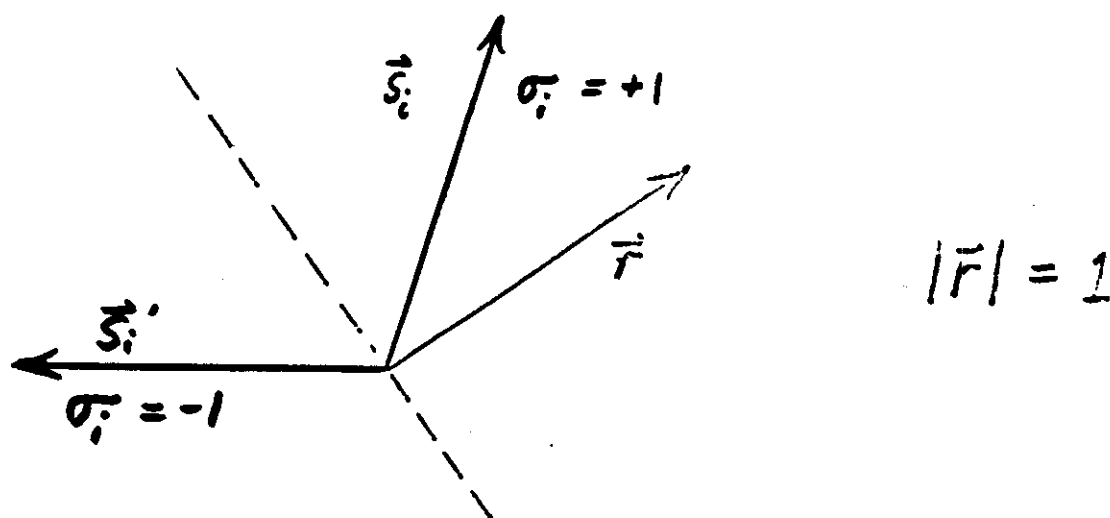


Embedded Ising Variables

U. Wolff

For $O(n)$ models

Example: $\vec{s}_i = (s_{ix}, s_{iy})$, $|\vec{s}_i| = 1$



$$\vec{s}_i' = \vec{s}_i + (\sigma_i - 1) (\vec{F} \cdot \vec{s}_i) \vec{F}$$

Application to

Anti ferromagnet

$$K < 0$$

PROBLEM:

$$p = 1 - e^{-K} < 0$$

$$q = e^{-K} > 1$$

Ising Antiferromagnet

$$K < 0$$

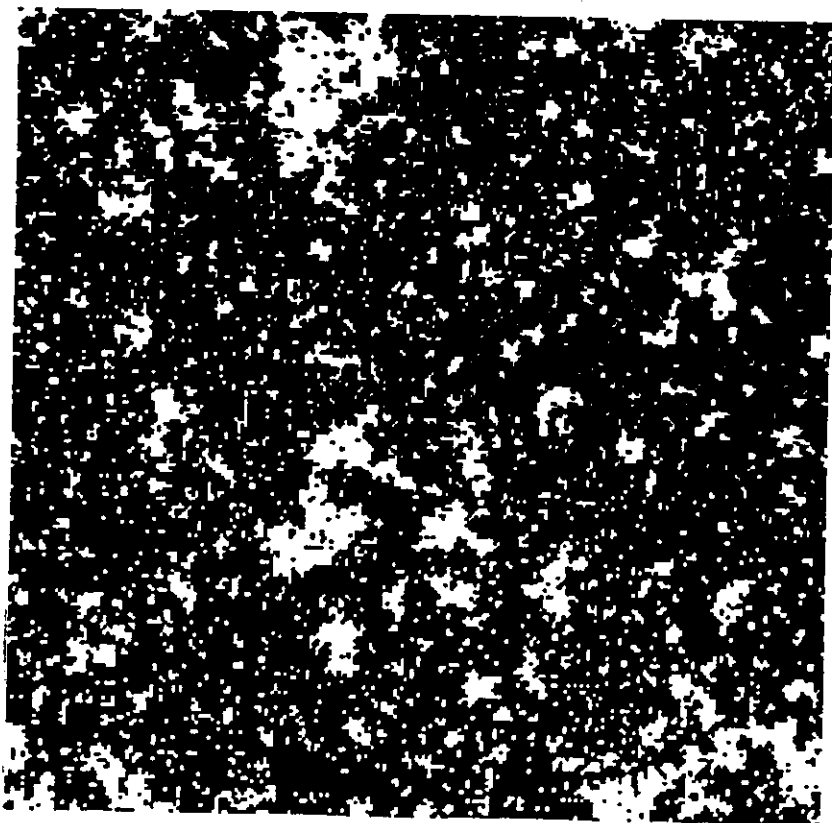
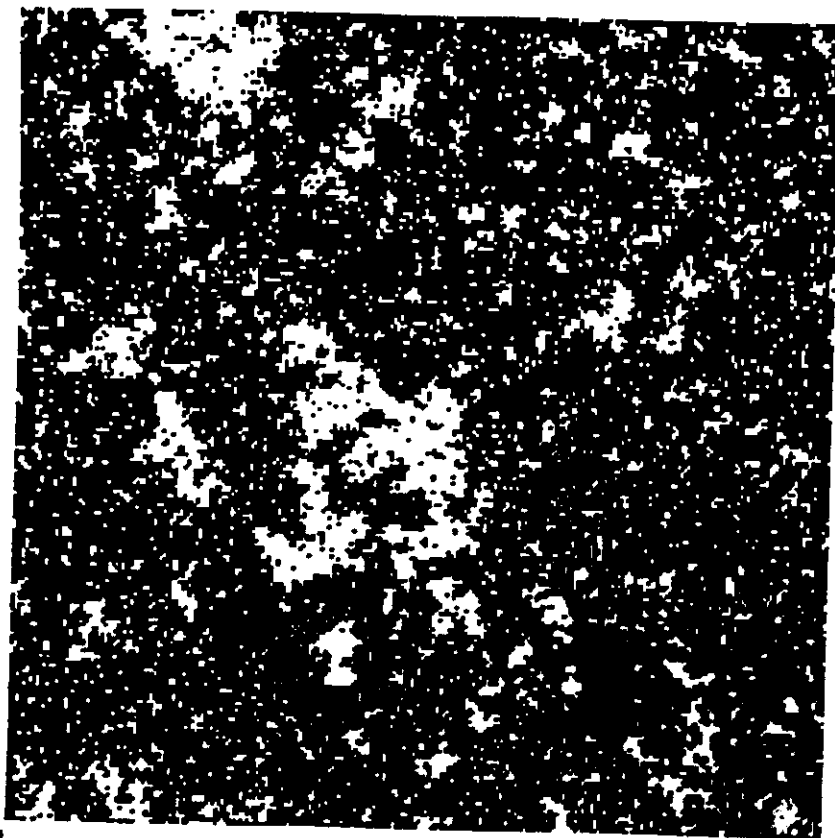
Eliminate $Z_{\langle lm \rangle}^{\text{same}}$

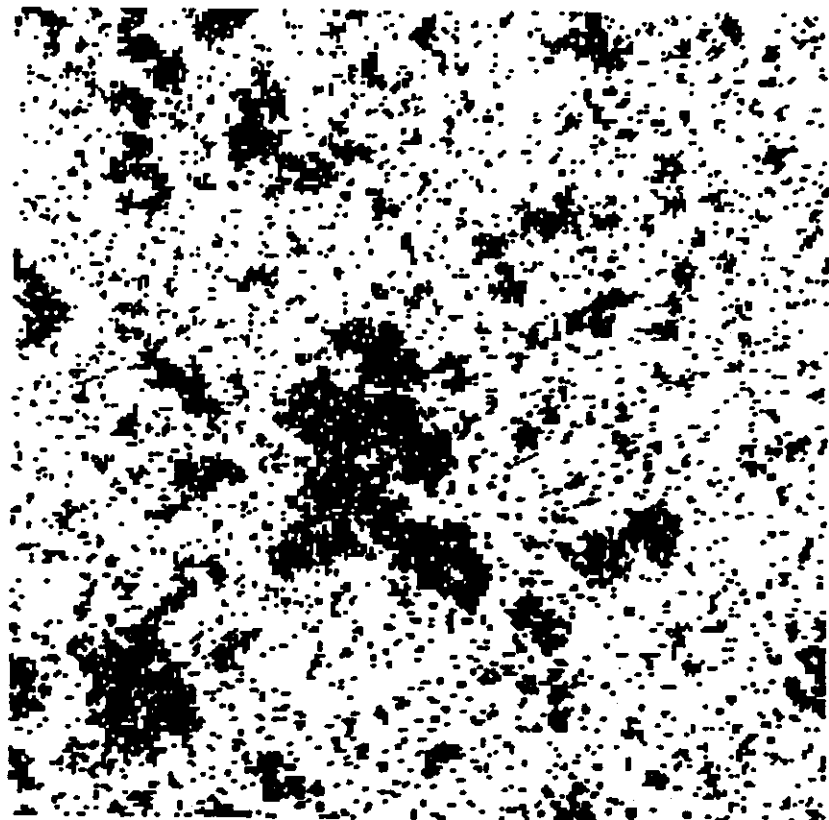
$$e^K Z = (1 - e^K) Z_{\langle lm \rangle}^{\text{diff}} + e^K Z_{\langle lm \rangle}^{\text{IND}}$$

$\uparrow$ $\uparrow$
 p $1-p$

$1 - e^K = \text{Probability of antibond}$

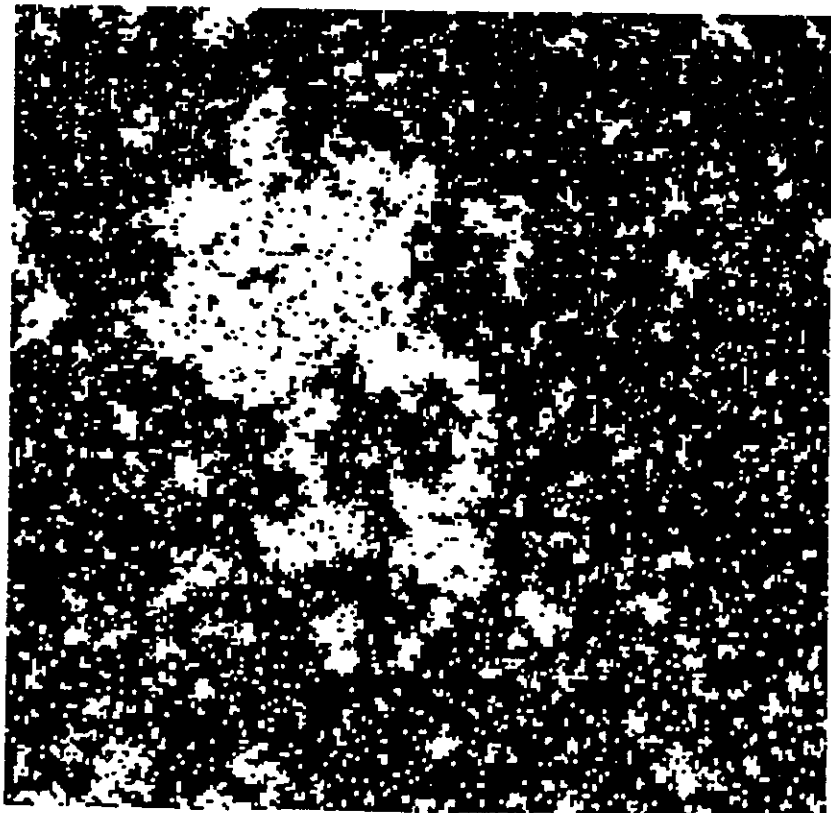
$\Rightarrow$ Anticlusters





4

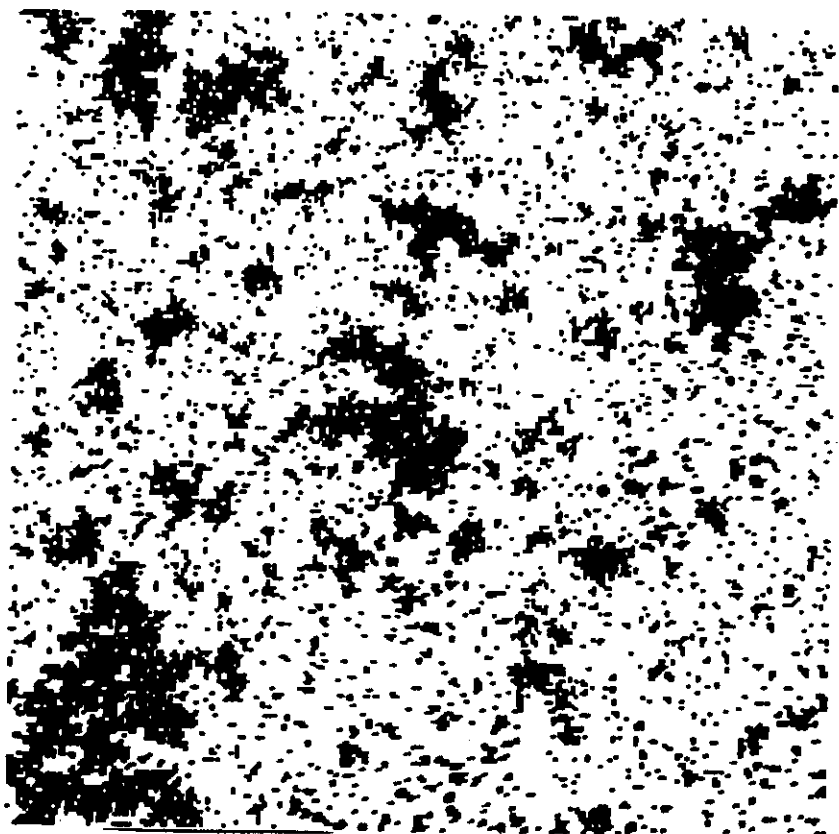
250



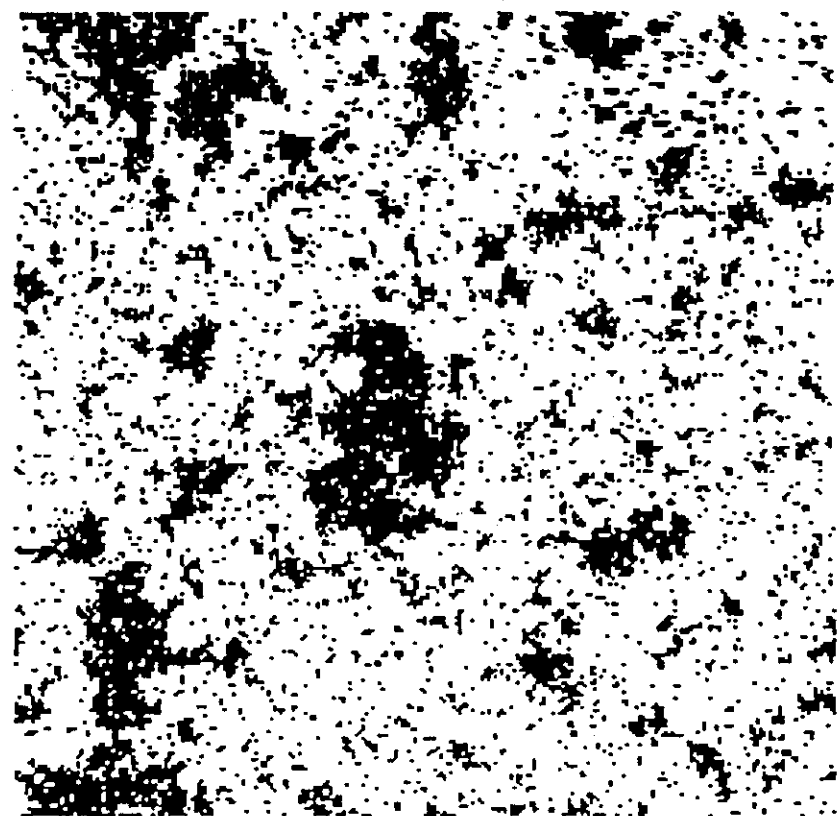
3

17-76

Field - 1000 ft. high



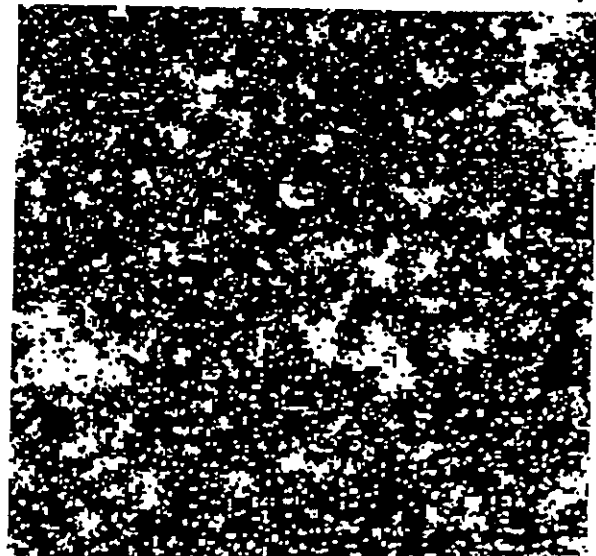
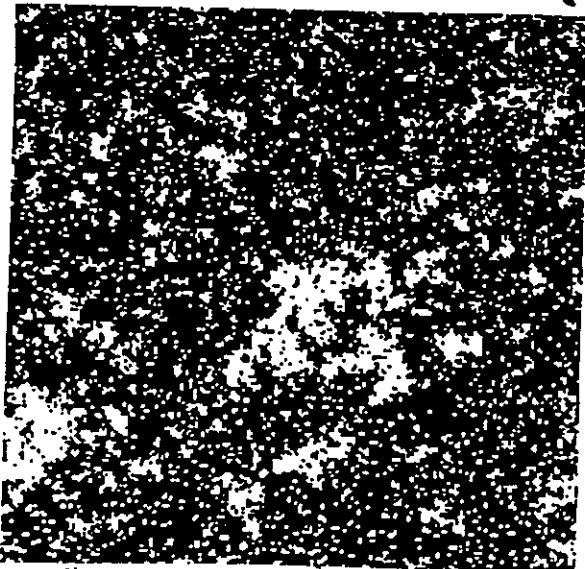
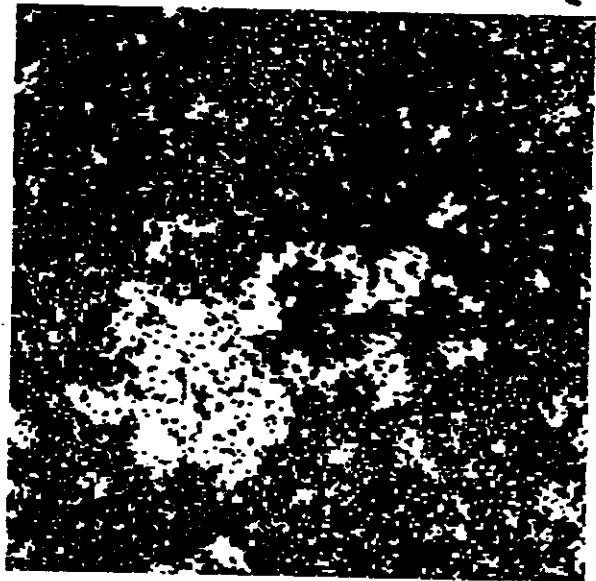
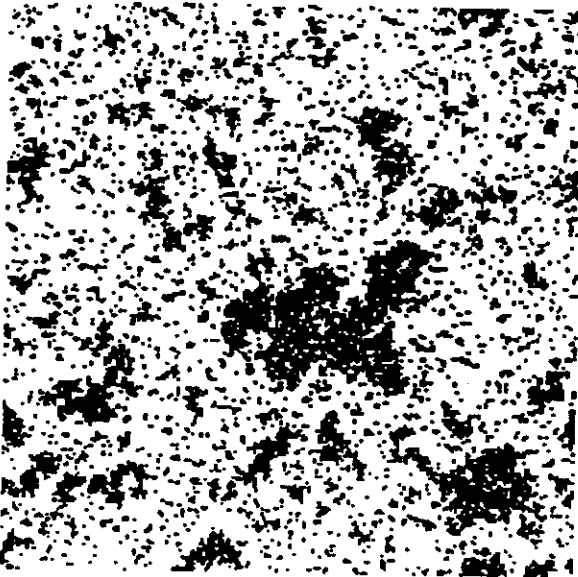
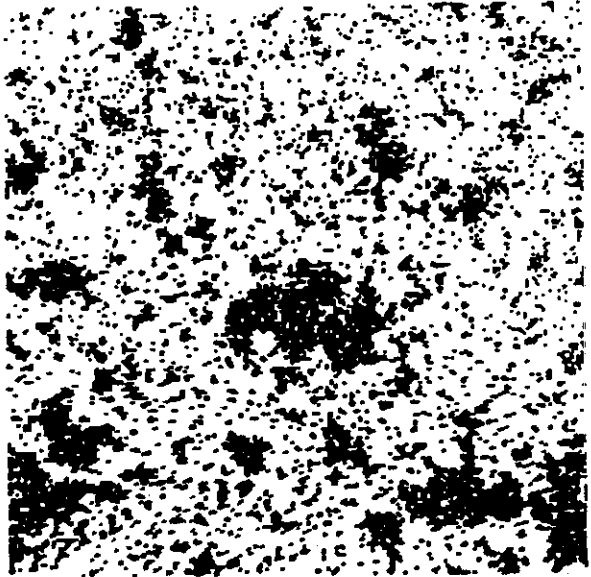
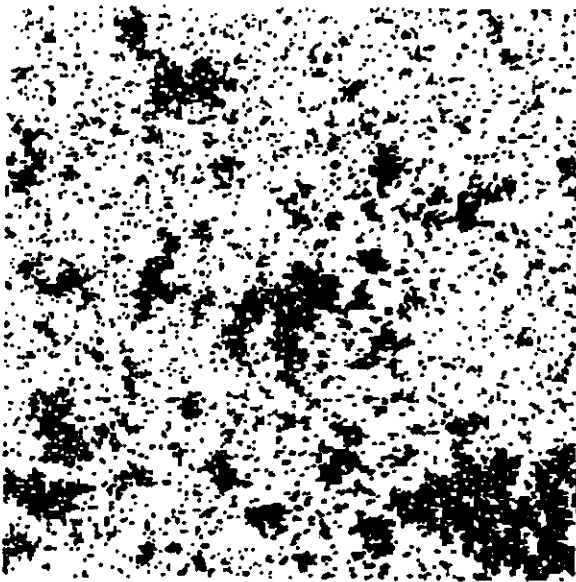
6



Path. microsc. lat. 100

T. 10

254



1
954x.
not 96/00.
-544

"Density of States"

$W(E)$ is the number of states with energy E

$$Z = \sum_{\{\sigma\}} \exp[-\beta E(\sigma)]$$

$$= \sum_E W(E) e^{-\beta E}$$

$$P(E) = \frac{1}{Z} W(E) e^{-\beta E}$$

Partition Function

$$\begin{aligned} Z &= e^{-\beta F} = \sum_{\{\sigma\}} \exp[-\beta \mathcal{H}(\sigma)] \\ &= \sum_E W(E) \exp[-\beta E] \end{aligned}$$

$$\beta = 1/k_B T$$

Monte Carlo simulation at β_i

$\Rightarrow$ histogram $N_i(E)$

$$\sum_E N_i(E) = n_i$$

$$\overline{N_i(E)} = n_i W(E) e^{-\beta_i E} e^{f_i}$$

$$f_i \equiv \beta_i F(\beta_i)$$

Approximation for $W(E)$

$$W(E) \approx n_i^{-1} N_i(E) e^{\beta_i E} e^{-f_i}$$

Then for other values of β

$$\langle A(E) \rangle = \sum_E A(E) W(E) e^{-\beta E} / Z(\beta)$$

$$Z(\beta) = \sum_E W(E) e^{-\beta E}$$

Single - Histogram Equation

$$\langle A(E) \rangle_{\beta} = \frac{\sum_E A(E) N_i(E) \exp[-(\beta - \beta_i) E]}{\sum_E N_i(E) \exp[-(\beta - \beta_i) E]}$$

Ising Model

$d=2$ 16×16

Fig. 1

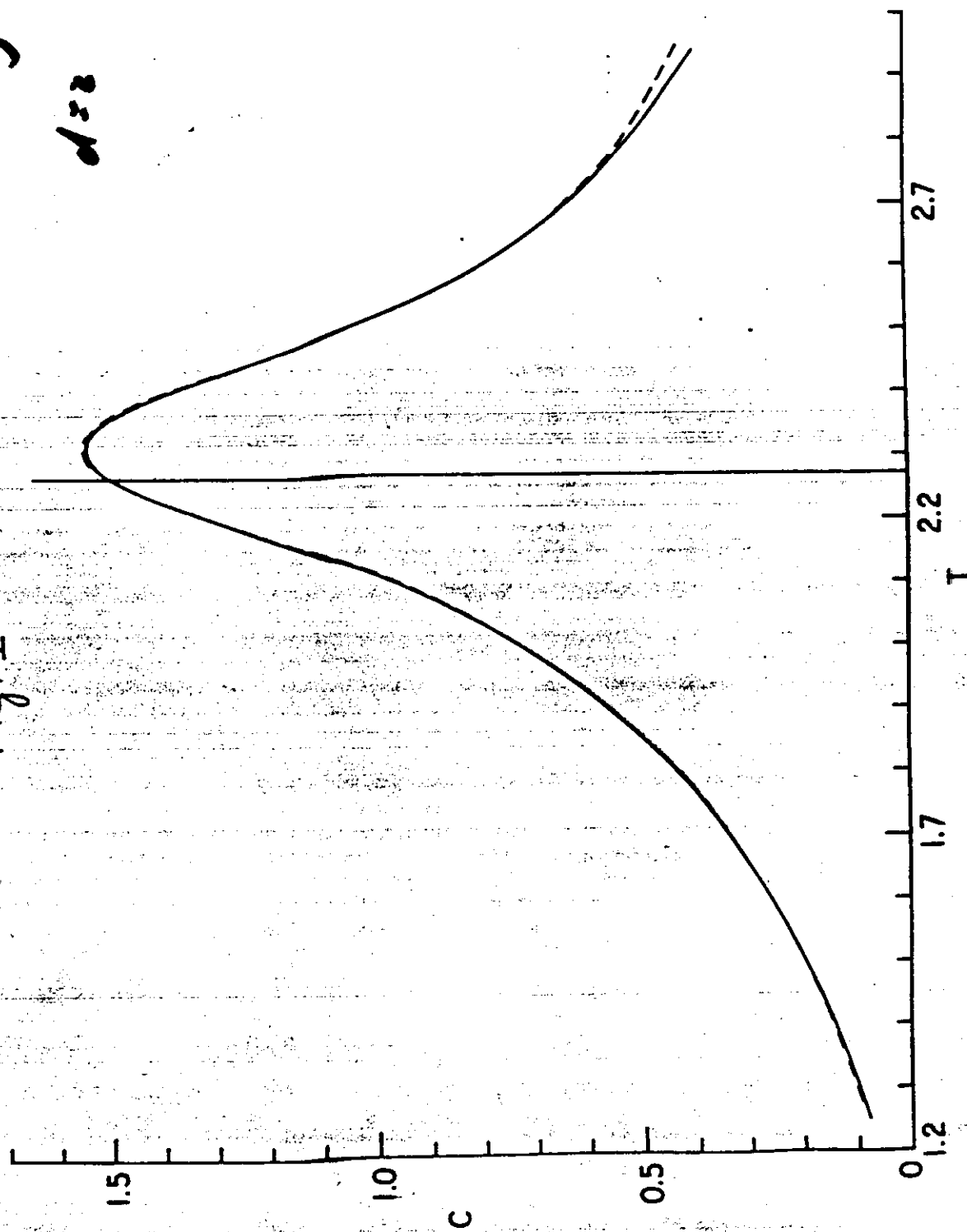
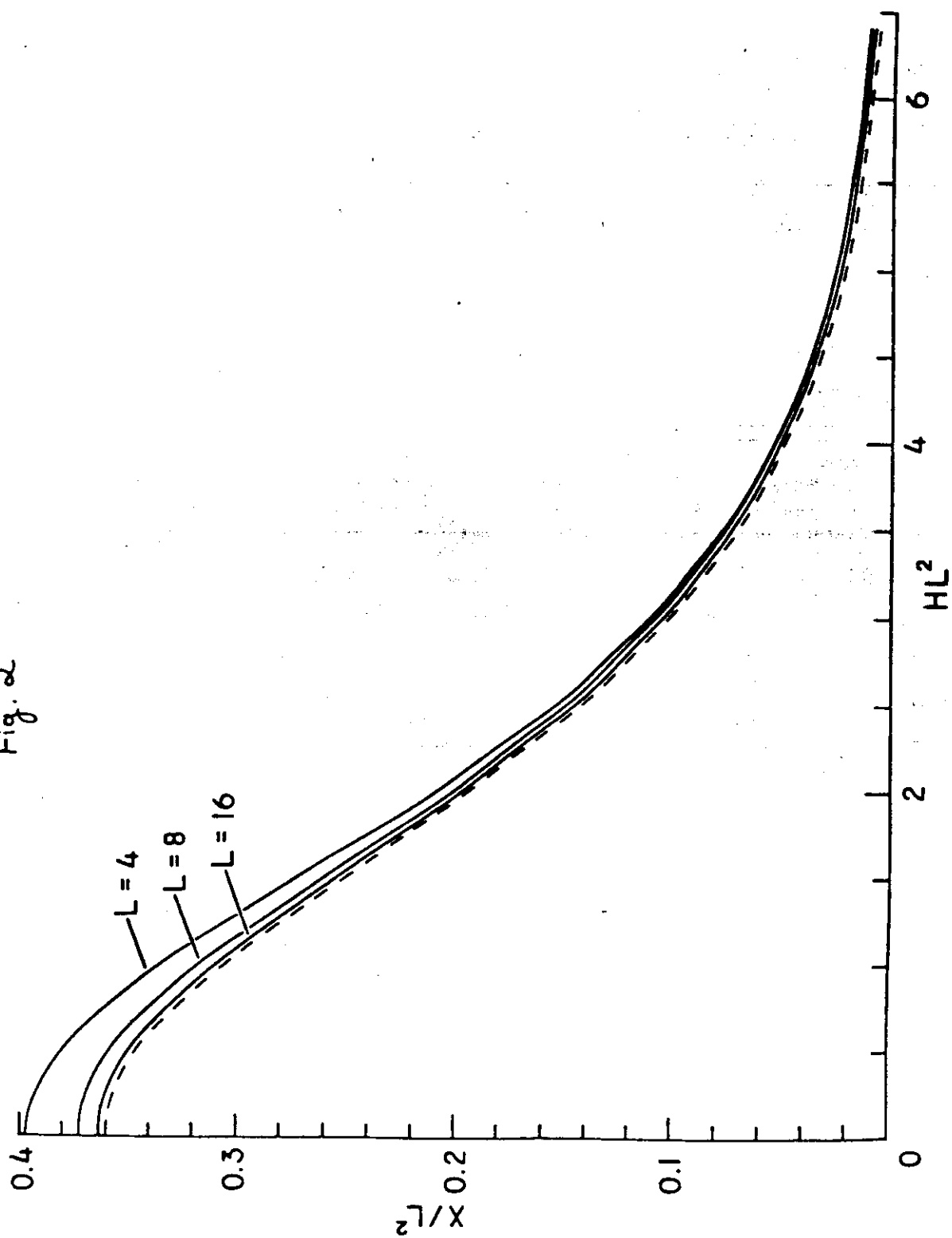


Fig. 2



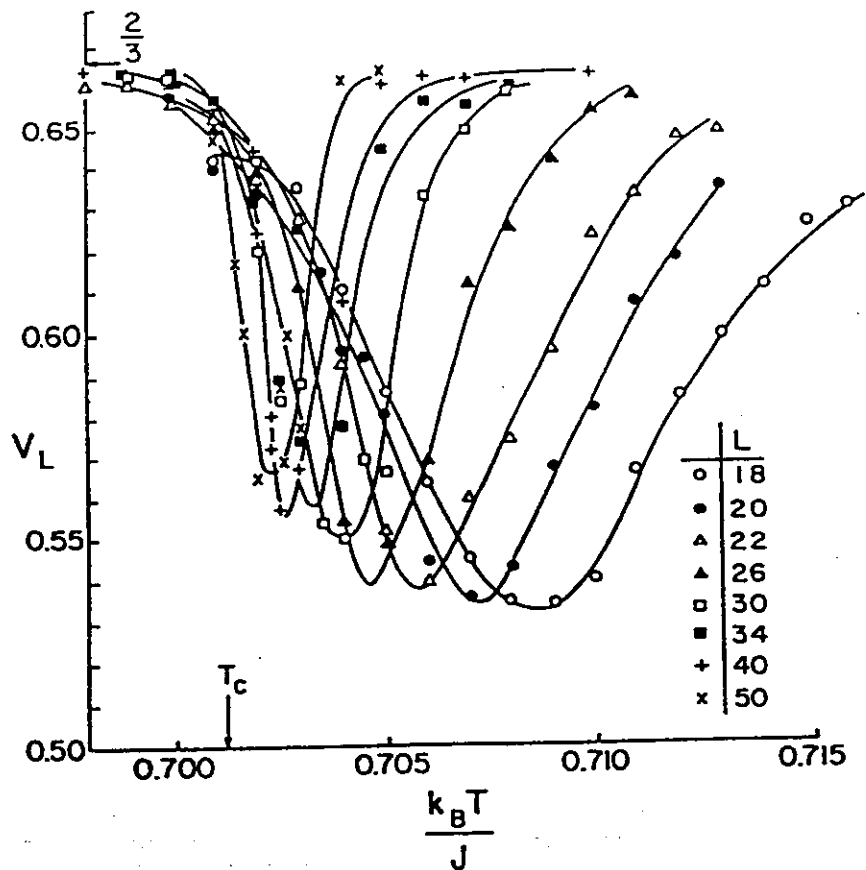


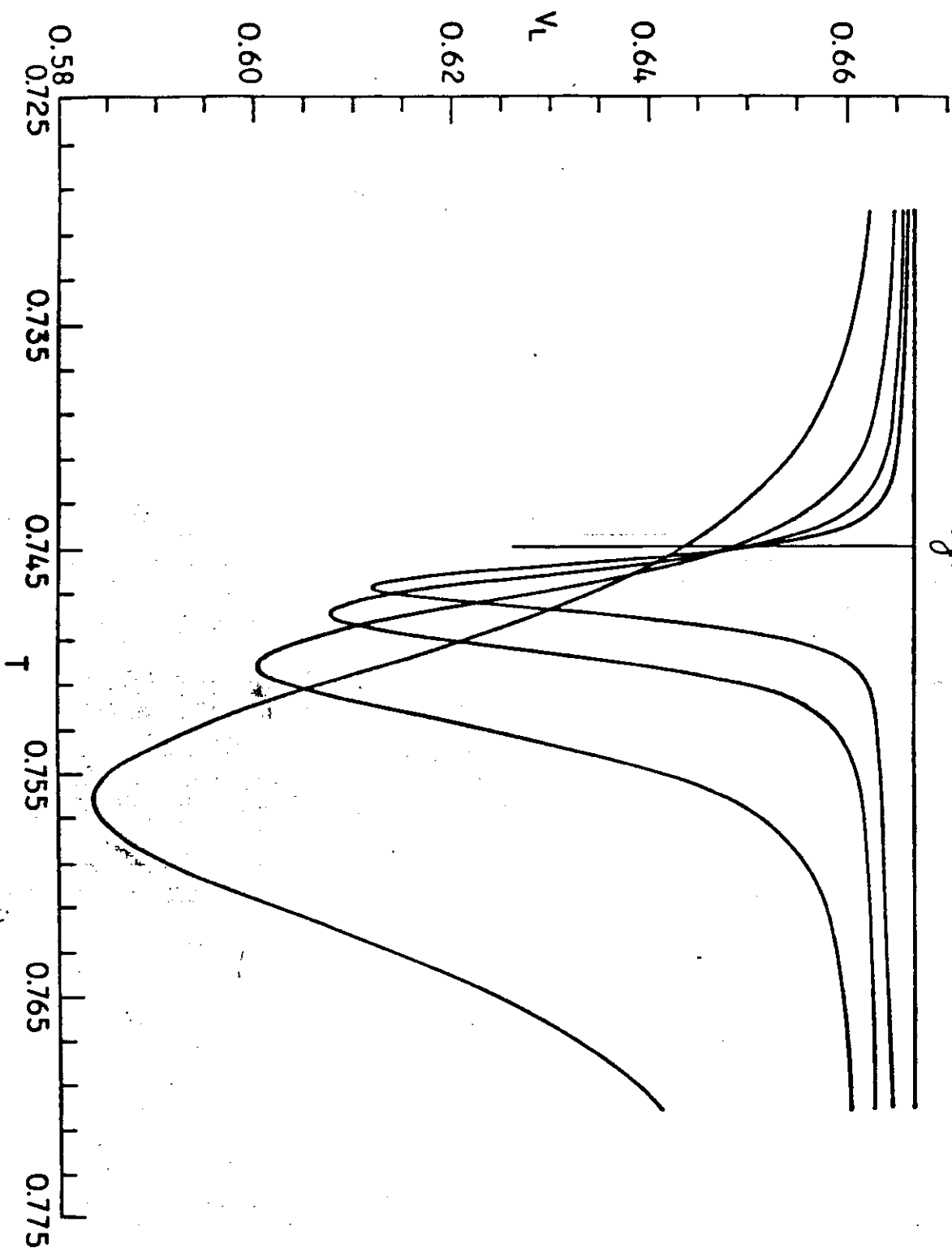
Fig. 1.4. Temperature variation of V_L , Eq. (1.11), of the two-dimensional 10-state Potts model for various lattice sizes. The transition temperature T_c and the trivial limit ($2/3$) for $V_L \rightarrow \infty$ are indicated. The nontrivial limit [$V_L|_{\min} \rightarrow 0.559$, Eq. (1.18)] is consistent with the data. (From [1.48])

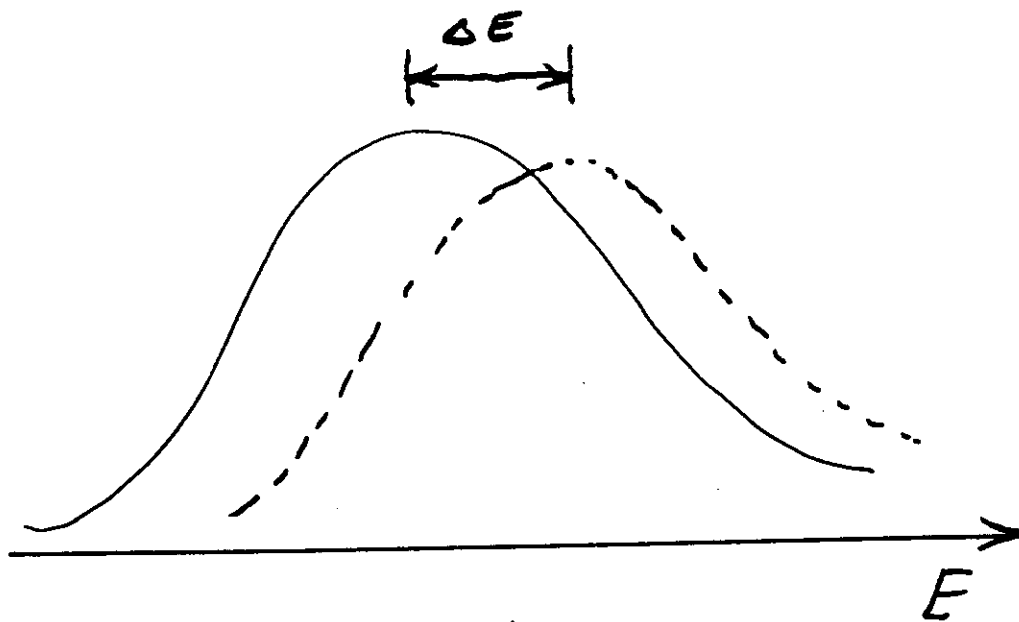
$$V_L = 1 - \frac{\langle E^4 \rangle_L}{3 \langle E^2 \rangle_L^2}$$

$$V_L^{\min} \rightarrow \begin{cases} 2/3 & 2^{\text{nd}} \text{ order} \\ \frac{2}{3} - \frac{1}{3} \left(\frac{E_+^2 - E_-^2}{2E_+ E_-} \right)^2 & 1^{\text{st}} \text{ order} \end{cases}$$

$l=8$ bits
 $d=2$

Fig. 3





$$\Delta E \approx [\langle (E - \langle E \rangle)^2 \rangle]^{\frac{1}{2}} = [L^d C]^{\frac{1}{2}}$$

$$\Delta E = \frac{\partial E}{\partial K} \Delta K = L^d C \Delta K$$

$$\Delta K \approx L^{-d/2} C^{-\frac{1}{2}}$$

$$\text{At } K_c \quad C \sim L^{\alpha/\nu}$$

$$\Delta K \sim L^{-\frac{1}{2}(d+\alpha/\nu)} = L^{-1/\nu} = L^{-\chi}$$

$$\alpha = 2 - d\nu$$

$$d + \alpha/\nu = 2/\nu$$

General Ensembles

$$P(E) \sim W(E) f(E)$$

Boltzmann

$$f(E) = e^{-\beta E}$$

$$\beta = 1/k_B T$$

"Multicanonical"

Berg + Neuhaus

$$f(E) \sim 1 / W(E)$$

$$\Rightarrow P(E) \sim \text{constant}$$

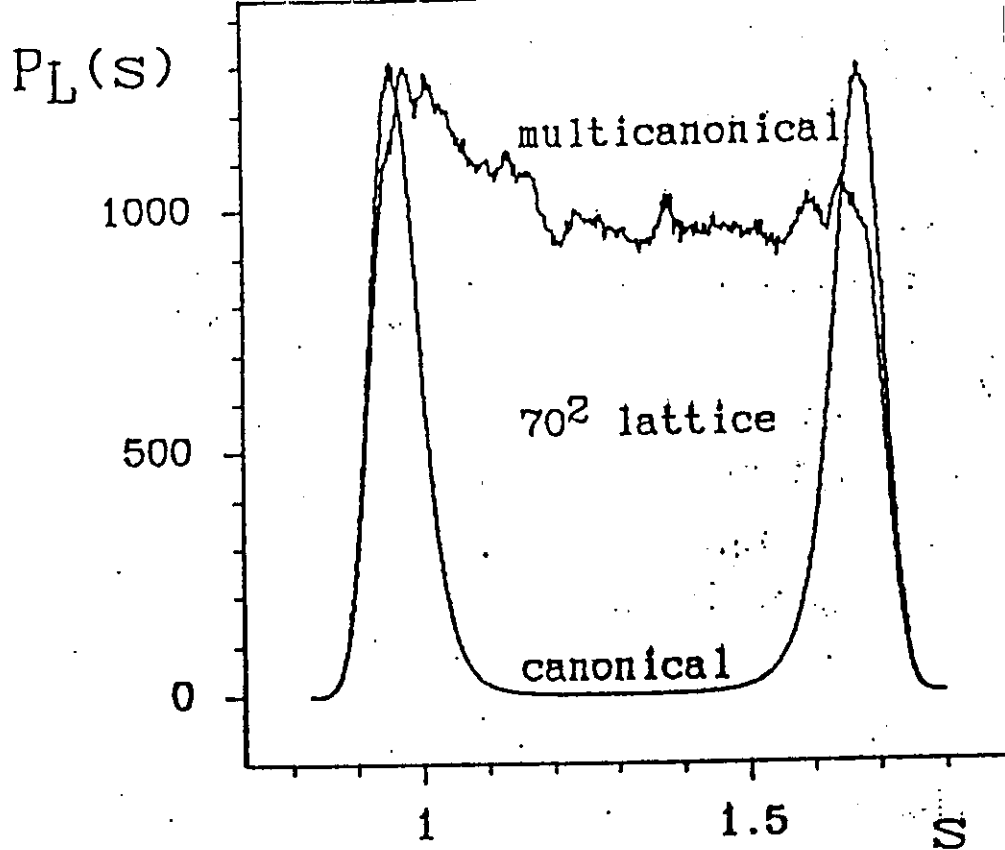
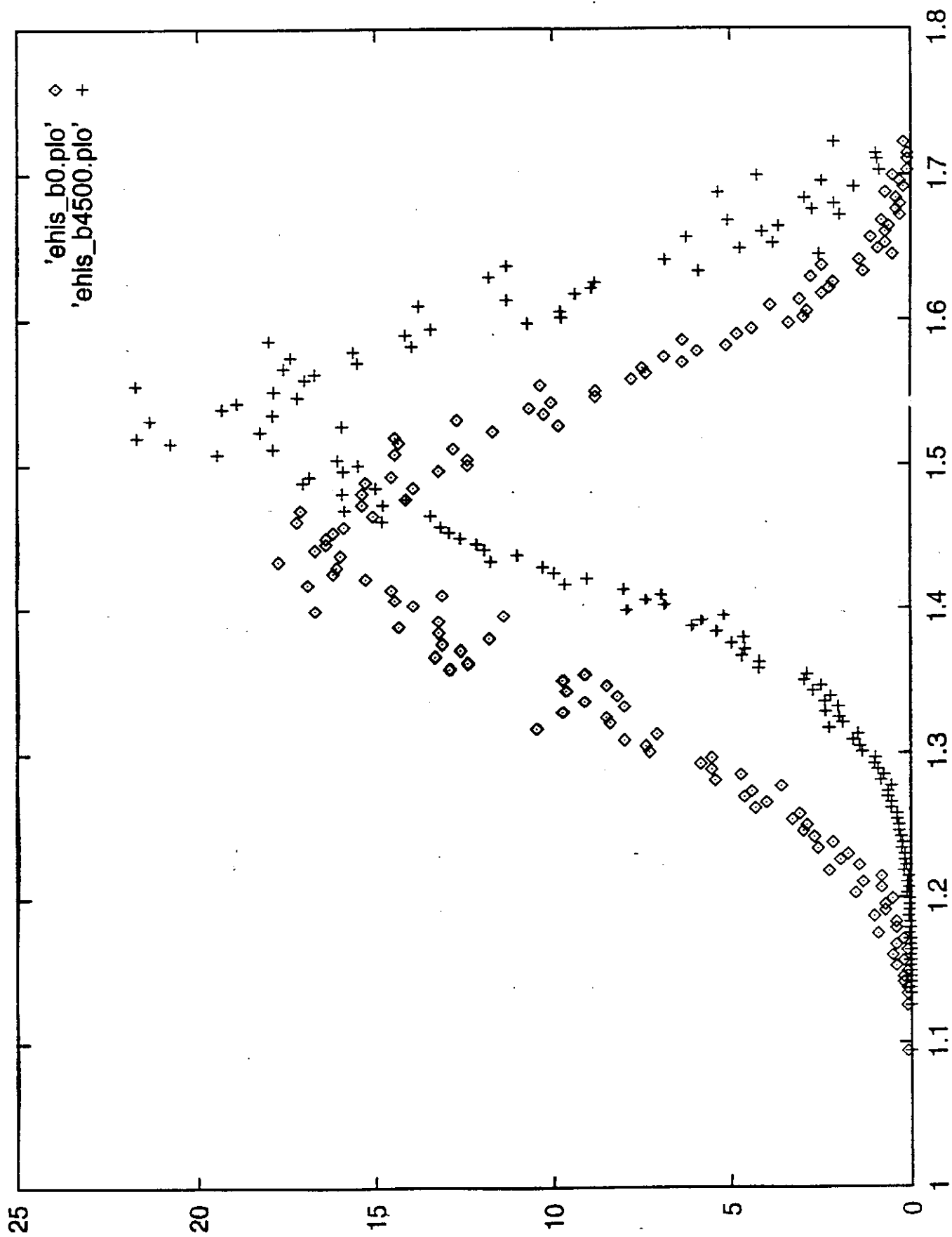


FIG. 2. Multicanonical action density distribution $\mathcal{P}_{70}(s)$ together with its reweighted distribution $P_{70}(s)$.

B. A. Berg and T. Nevhaus

Phys. Rev. Lett. 68, 9 (1992)



Multiple Histograms

$$W(E) = \sum_i p_i(E) n_i^{-1} N_i(E) \exp[\beta_i E - f_i]$$

$$f_i \equiv \beta_i F(\beta_i)$$

$$\sum_i p_i(E) = 1$$

Statistical error in $W(E)$

from statistical error in $\{N_i(E)\}$

Choose $\{p_i(E)\}$ to minimize error

Statistical Errors in Histograms

$N_i(E)$ independent of $N_j(E')$, $i \neq j$

Ferrenberg + Swendsen assume:

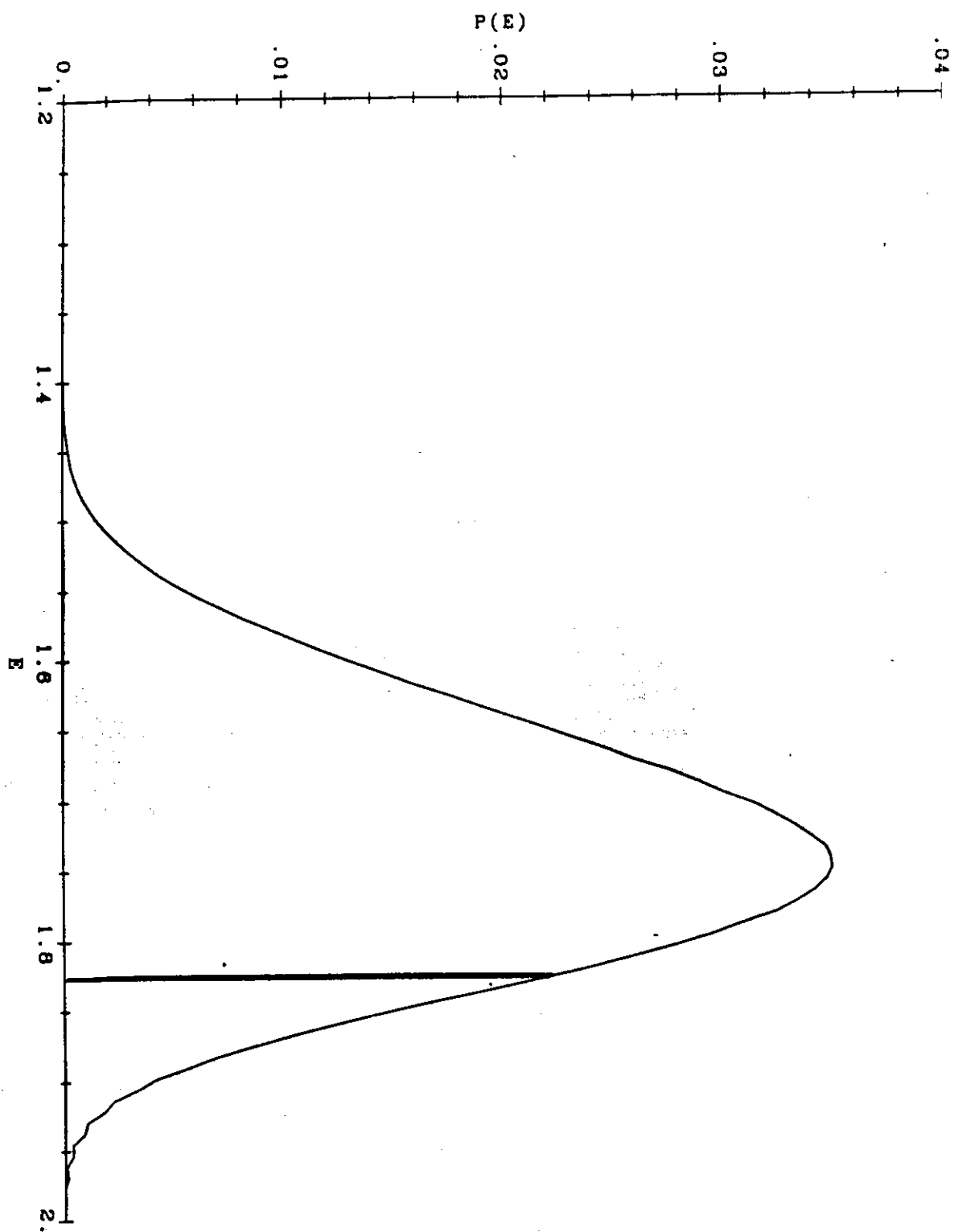
1) $N_i(E)$ independent of $N_j(E')$, $E \neq E'$

2) $S^2 N_i(E) = g_i \overline{N_i(E)}$

$$= g_i W(E) n_i \exp[-\beta_i E + f_i]$$

where $g_i = 1 + 2 \tau_i$

16x16



Multiple Histogram Equations

FS

$$W(E) = \frac{\sum_i g_i^{-1} N_i(E)}{\sum_j g_j^{-1} n_j \exp[-\beta_j E + f_j]}$$

$$e^{-f_i} = e^{-\beta_i F(\beta_i)} = \sum_E W(E) e^{-\beta_i E}$$

Error in density of states

$$\frac{\delta W(E)}{W(E)} = \left[\sum_i g_i^{-1} N_i(E) \right]^{-\frac{1}{2}}$$

Alternative treatment of error in $\{N_i(E)\}$

Huang, Moriarity, Myers, Potvin

Z. Phys. C 50, 221 (1991)

used $S^2 N_i(E) = g_i N_i(E)$

instead of

s) $S^2 N_i(E) = g_i W(E) n_i \exp[-\beta_i E + f_i]$

$N_i(E) = 0$ required special treatment

Using $W(E)$ includes information
from all histograms

Alternative treatment of error in $\{N_i(E)\}$

Alves, Berg, Villanova, PRB 41, 383 ('90)

64 replicas at each temperature

$\Rightarrow$ correct error for $\{N_i(E)\}$

(very nearly $S^2 N_i(E) = N_i(E)$)

But ignored correlations

between $N_i(E)$ and $N_i(E')$

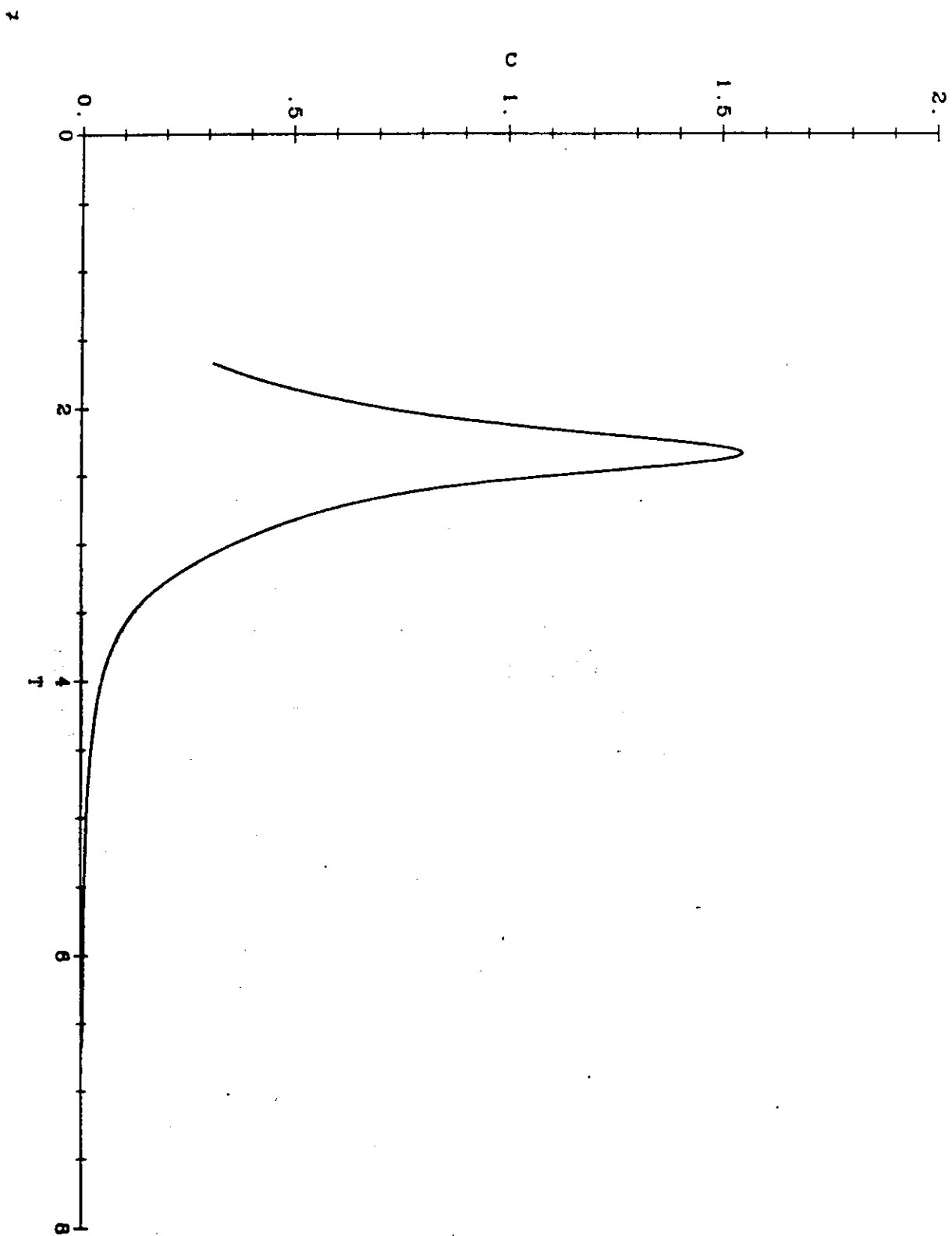
$\Rightarrow$ Total error underestimated

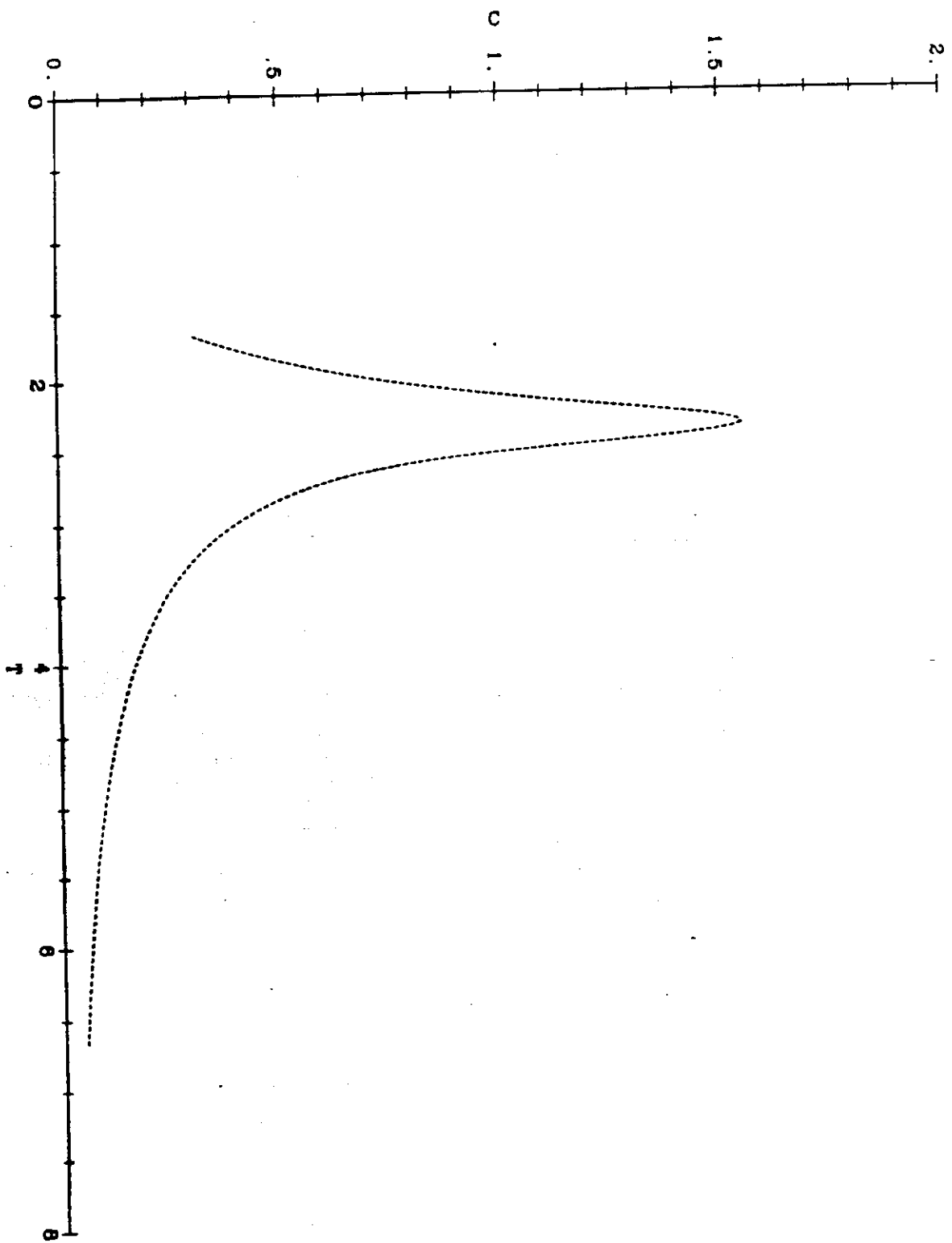
Different way of combining histograms

f_i 's determined separately for each E

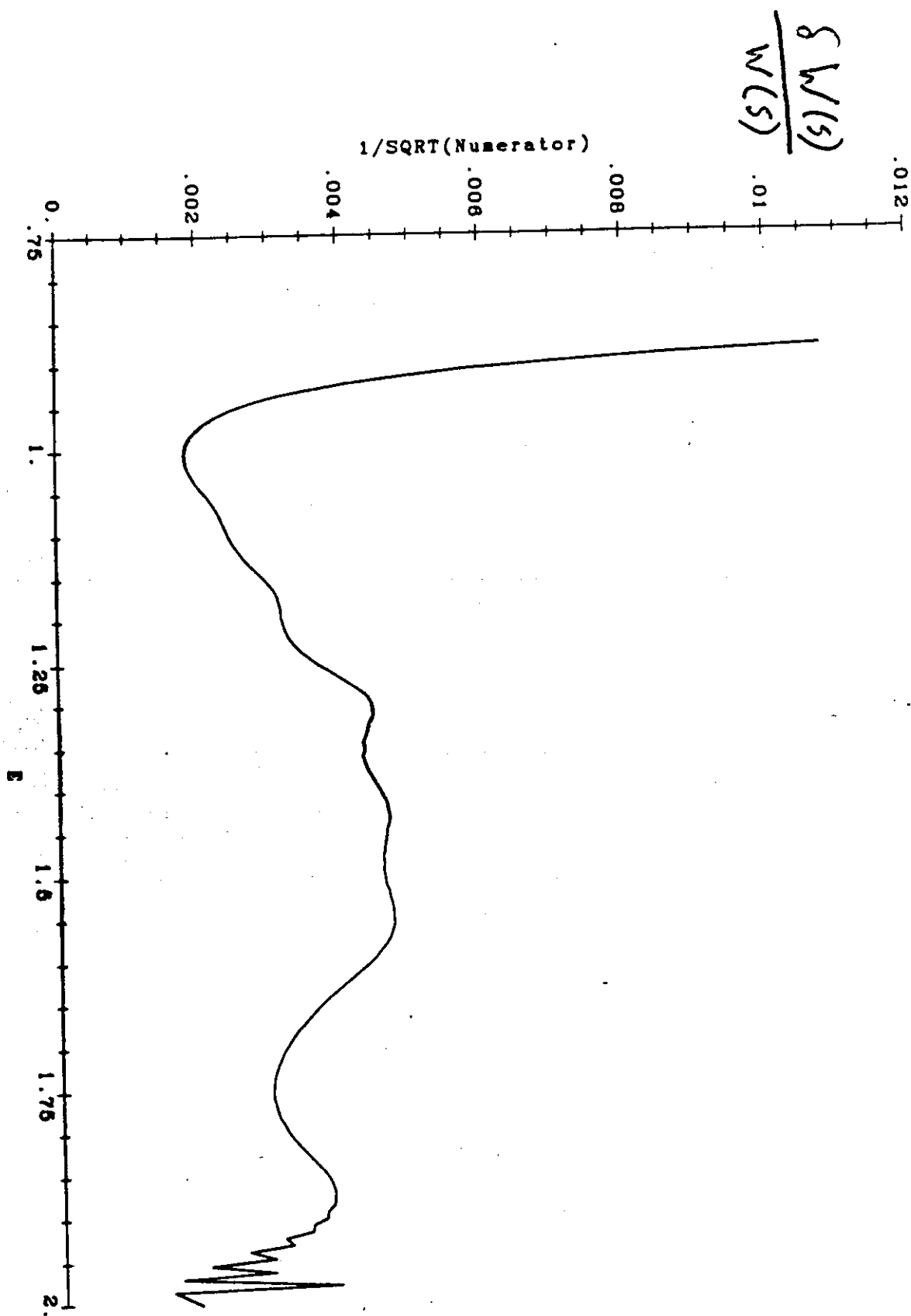
(not self consistently)

$$\{K_s\} = \{K_e\}$$

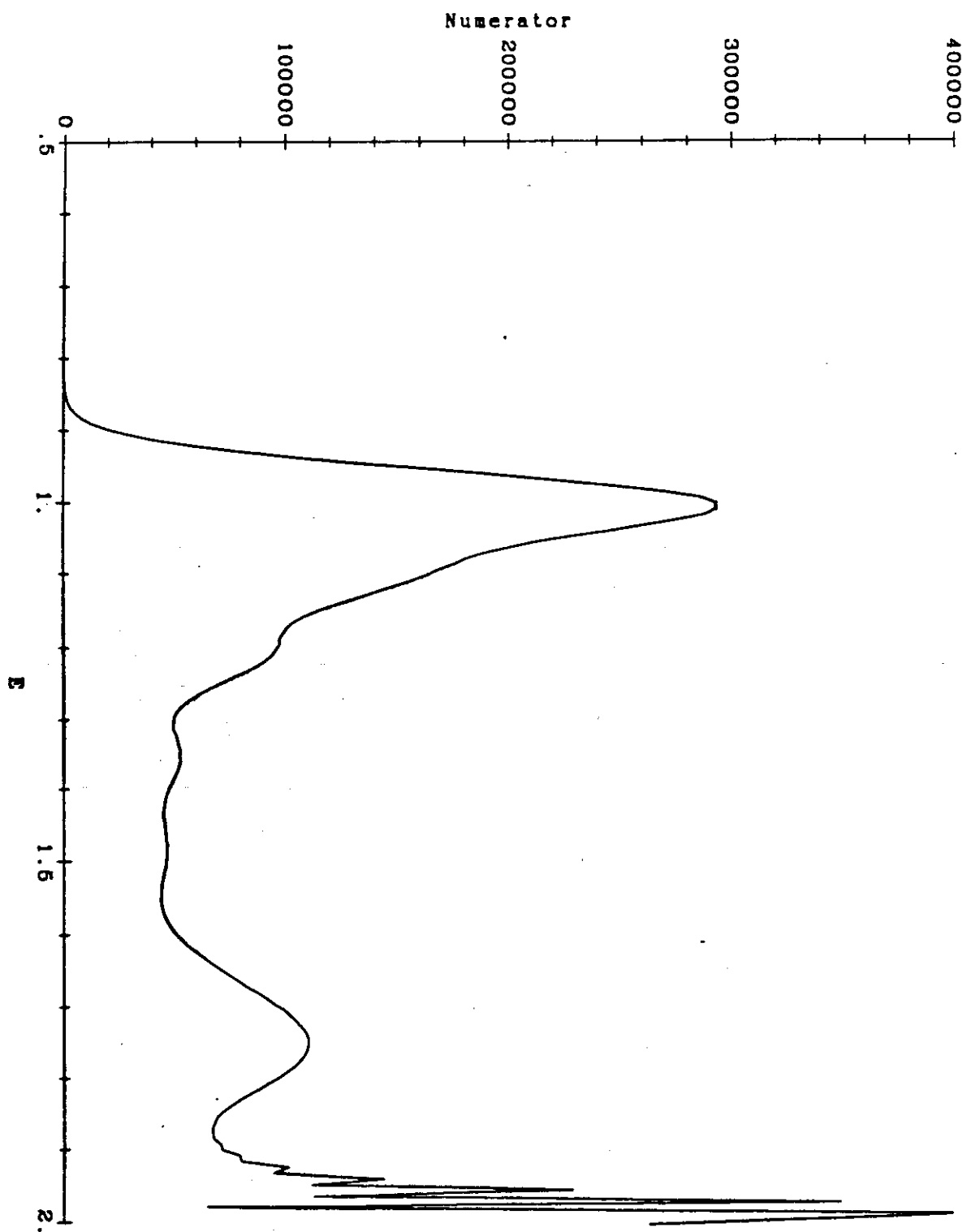


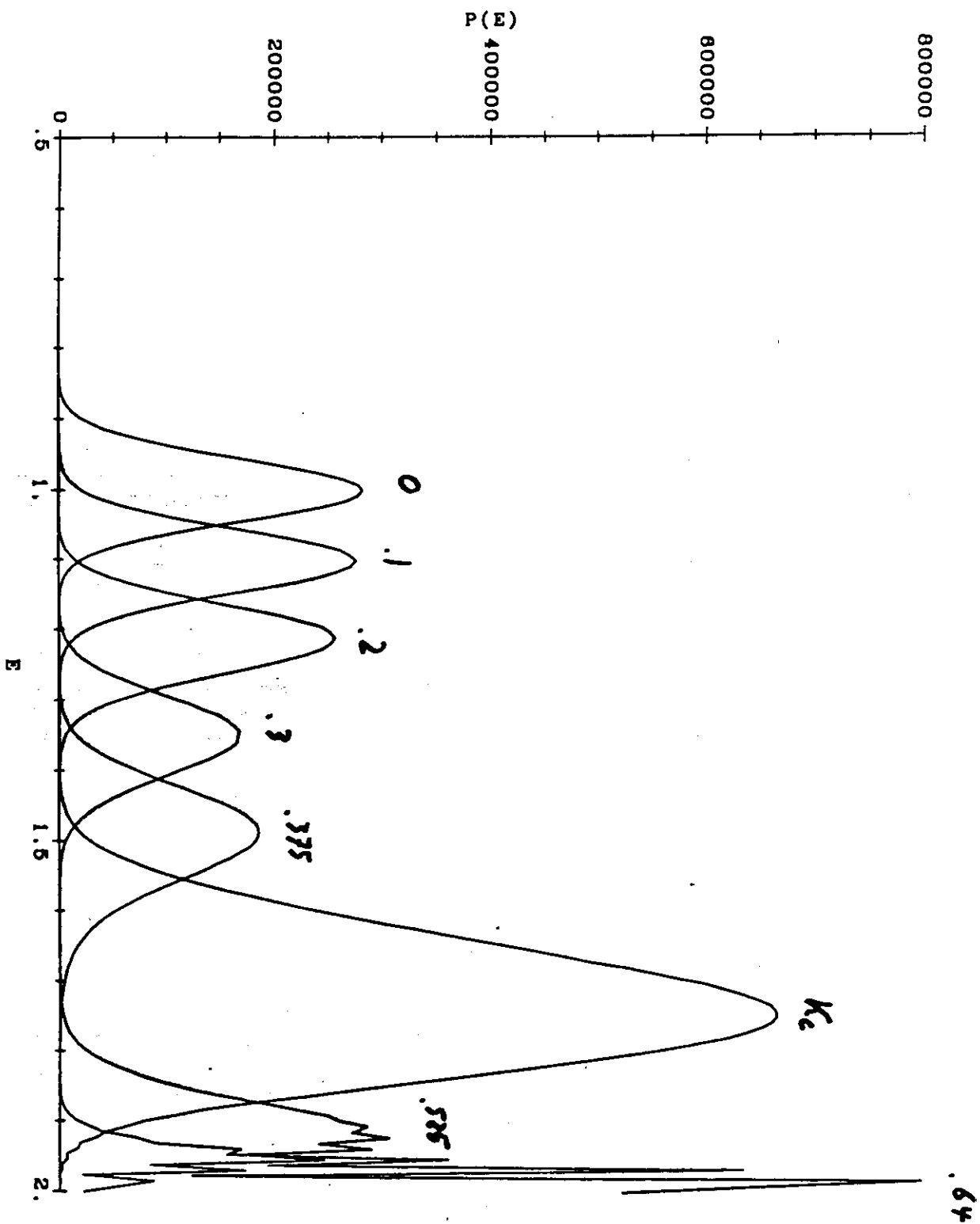


EXACT

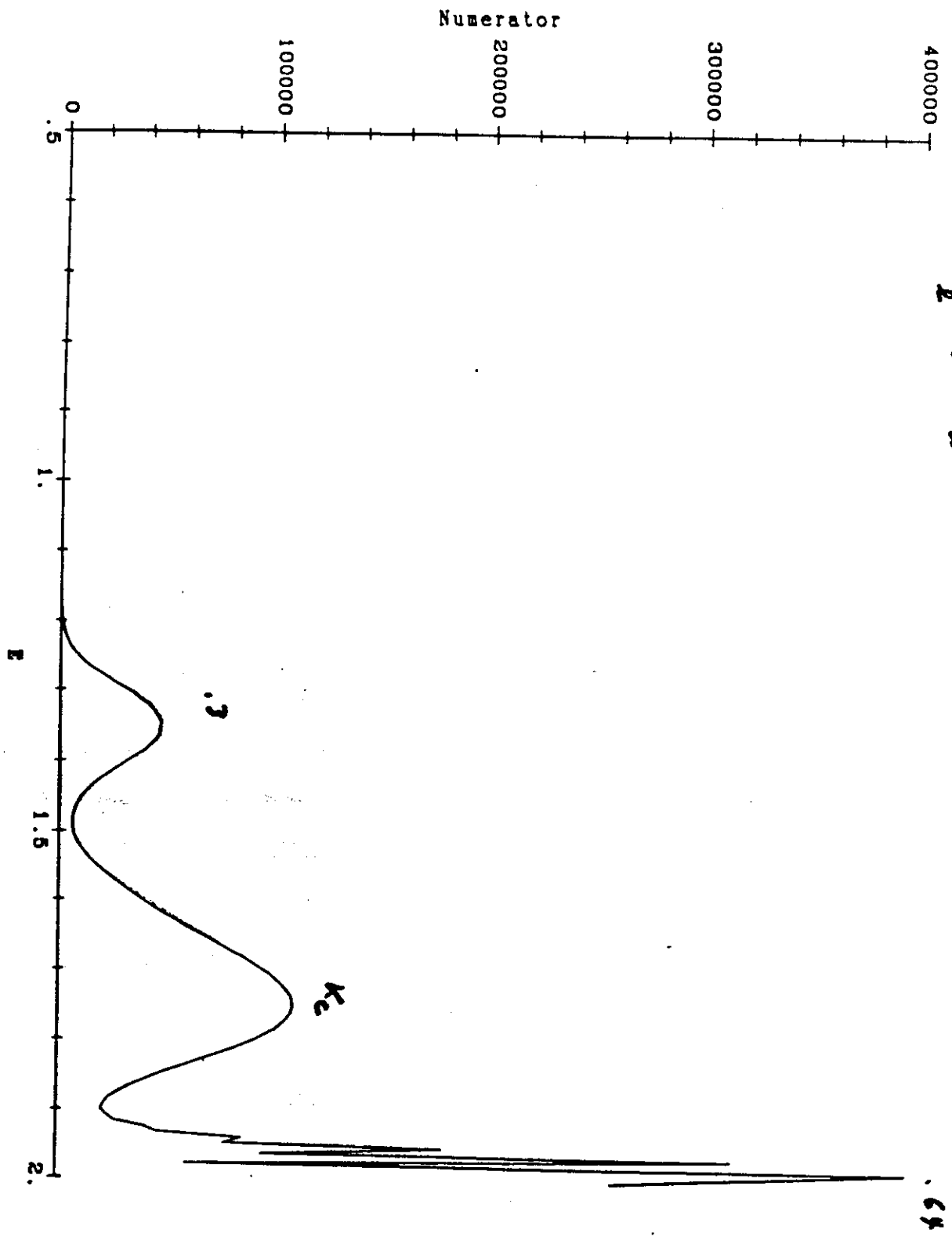


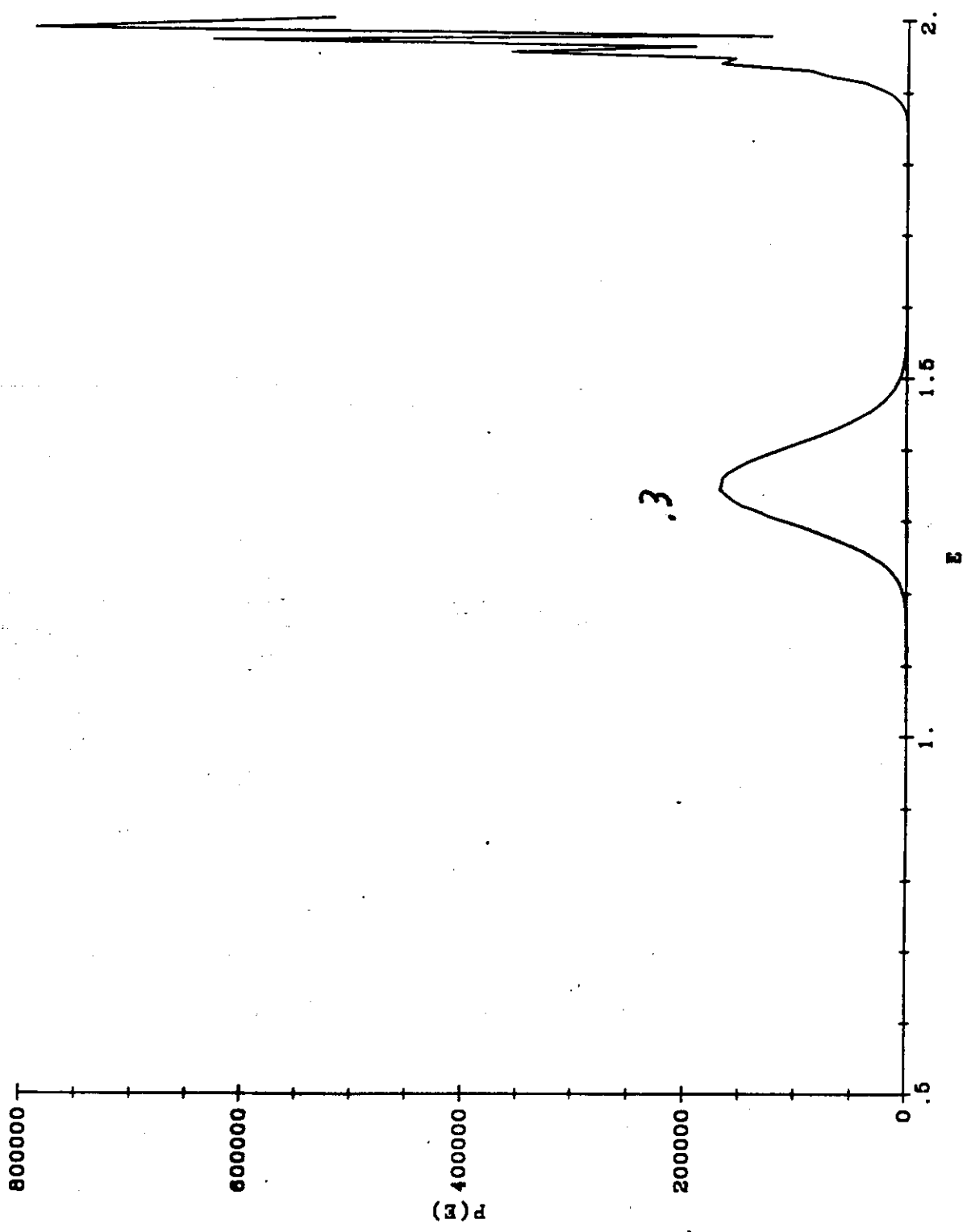
$$\sum_{k=1}^8 g_k^{-1} H_k(s)$$

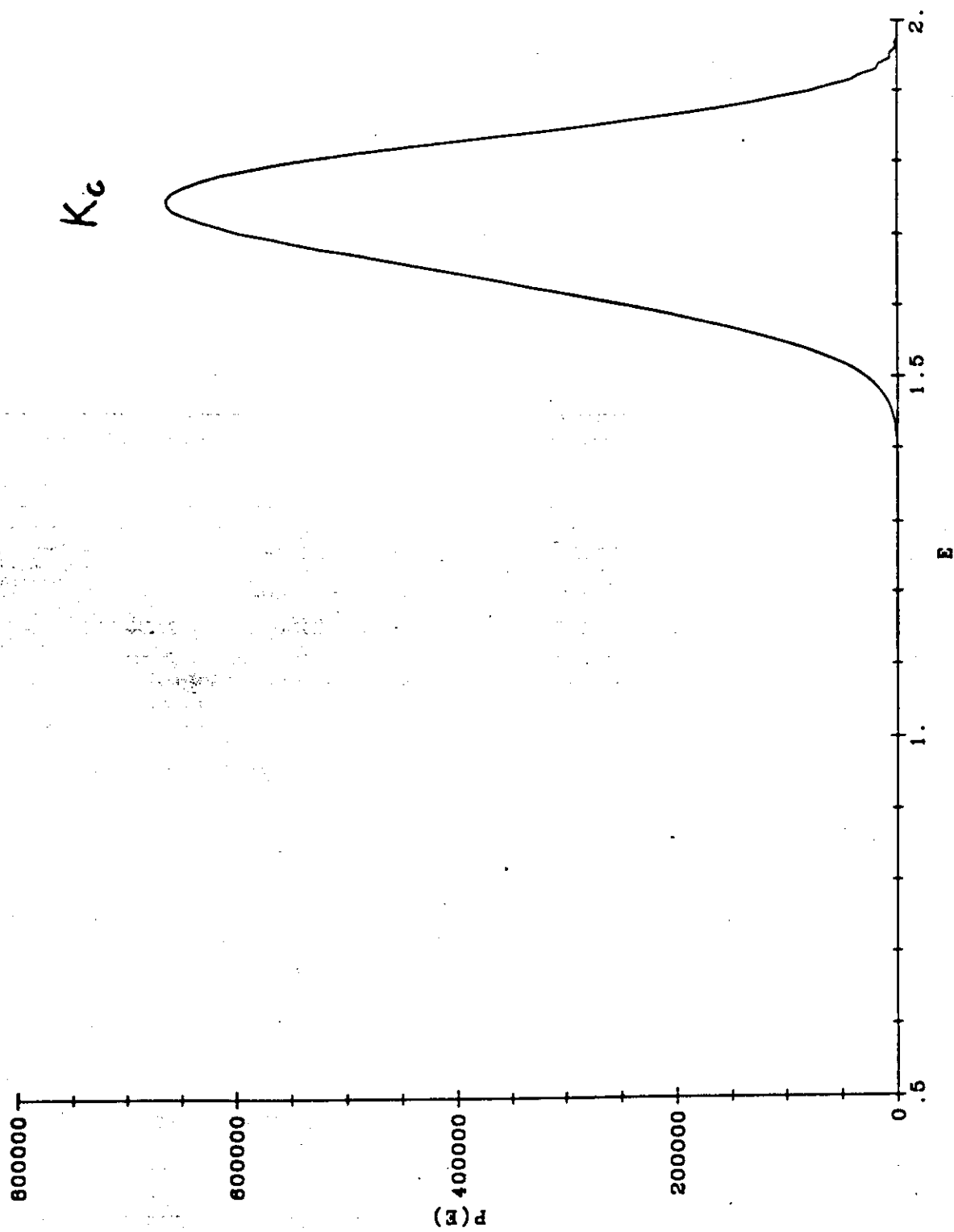




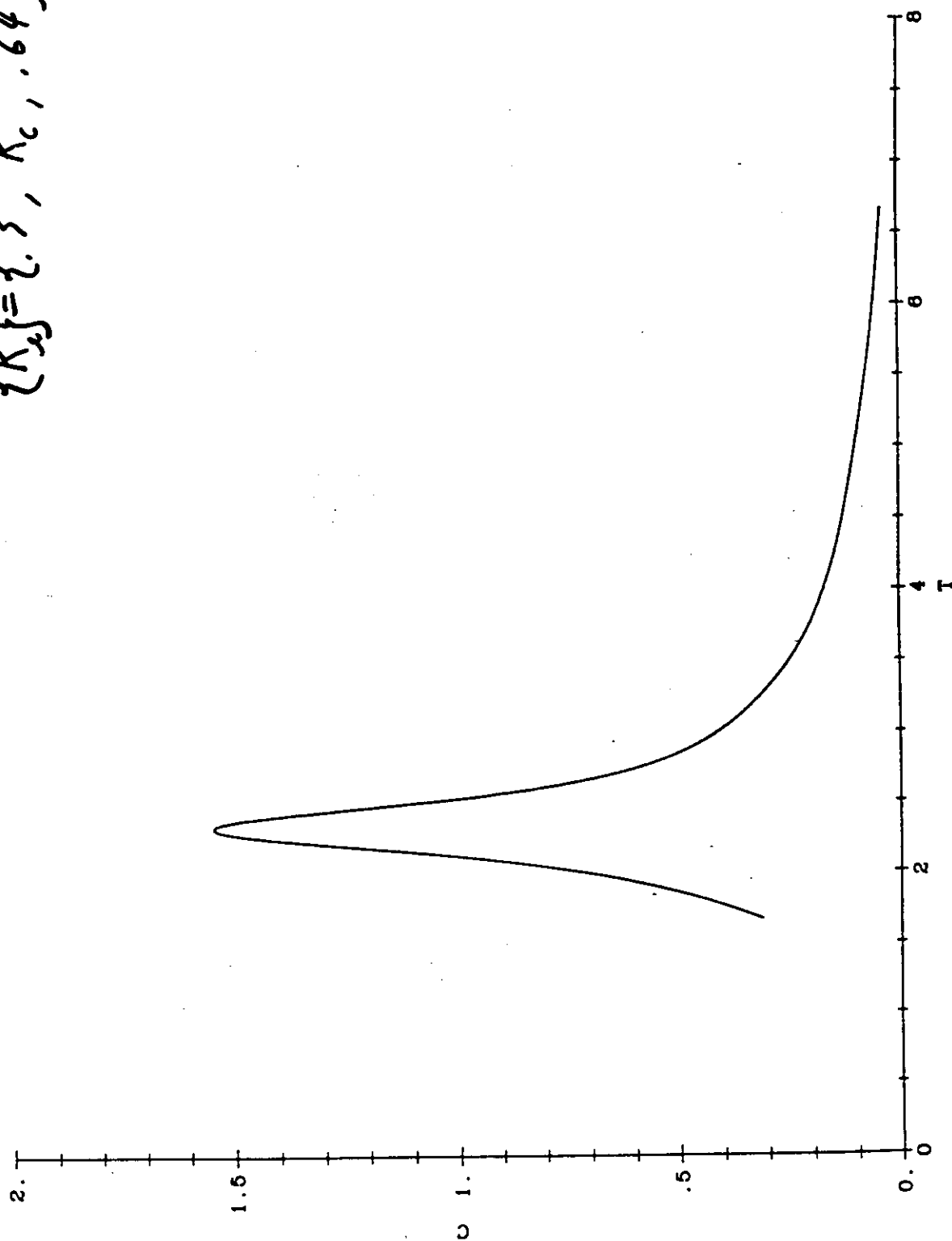
$$\sum g_n H_n(s)$$



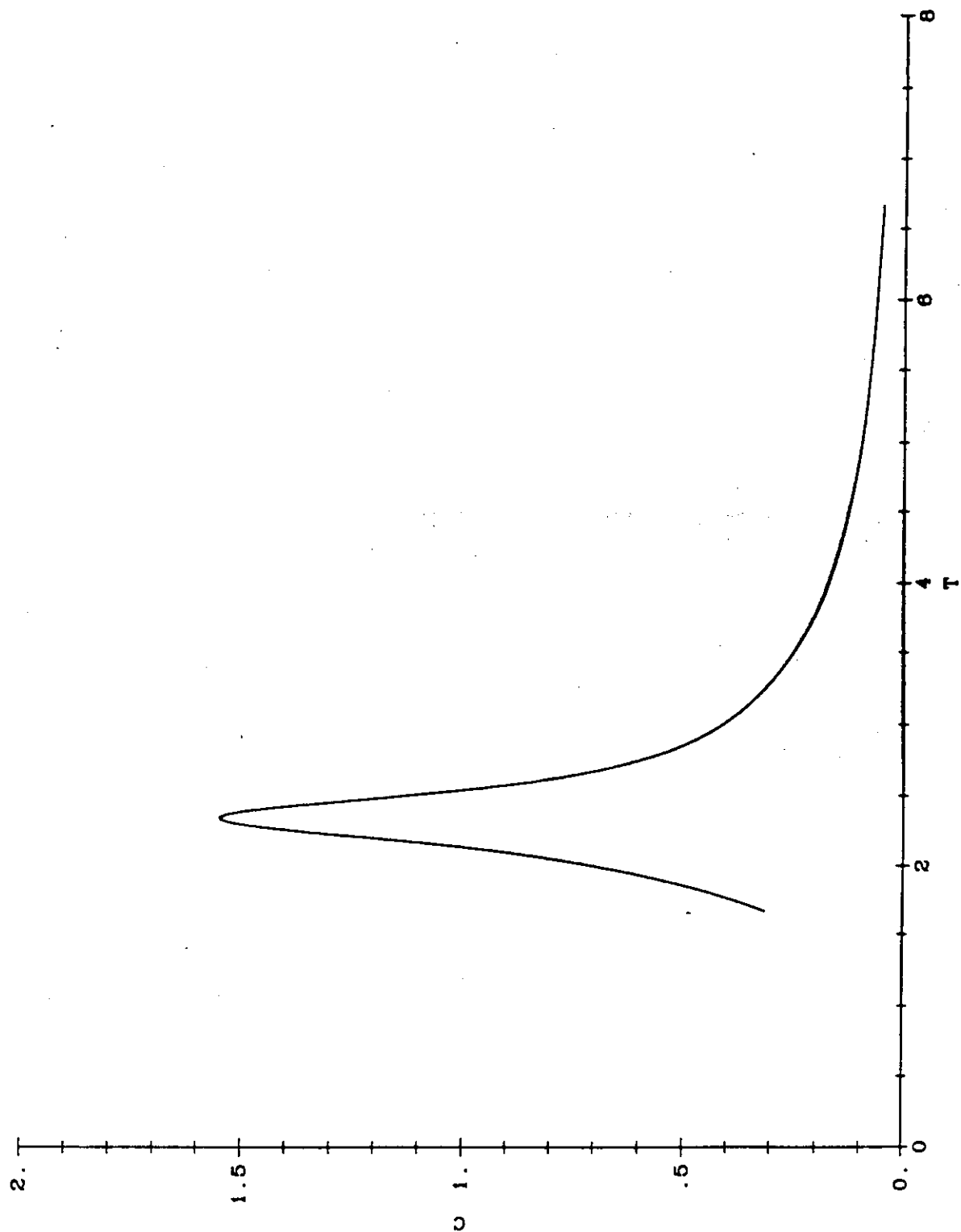


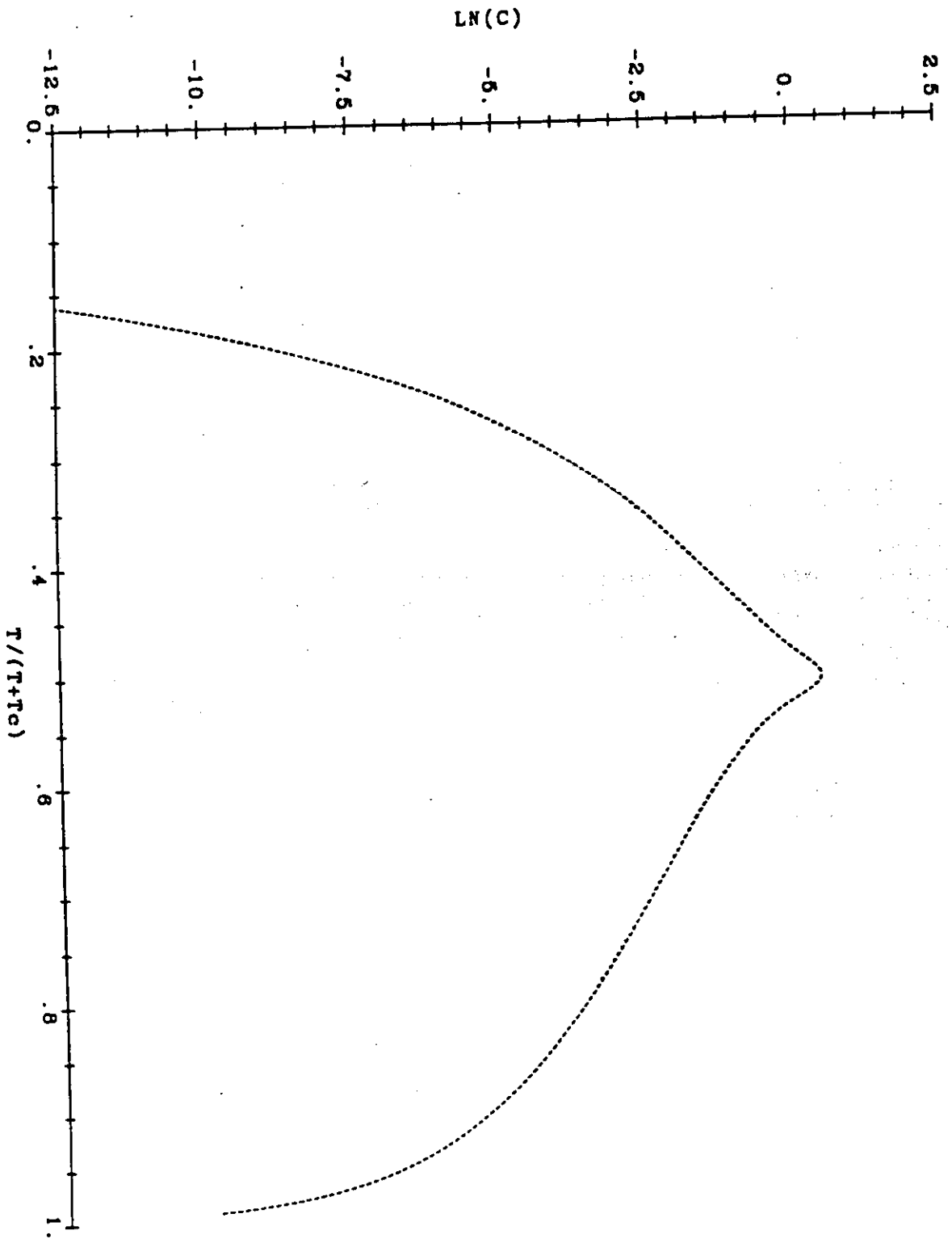


$$\{K_x\} = \{.3, K_c, .64\}$$

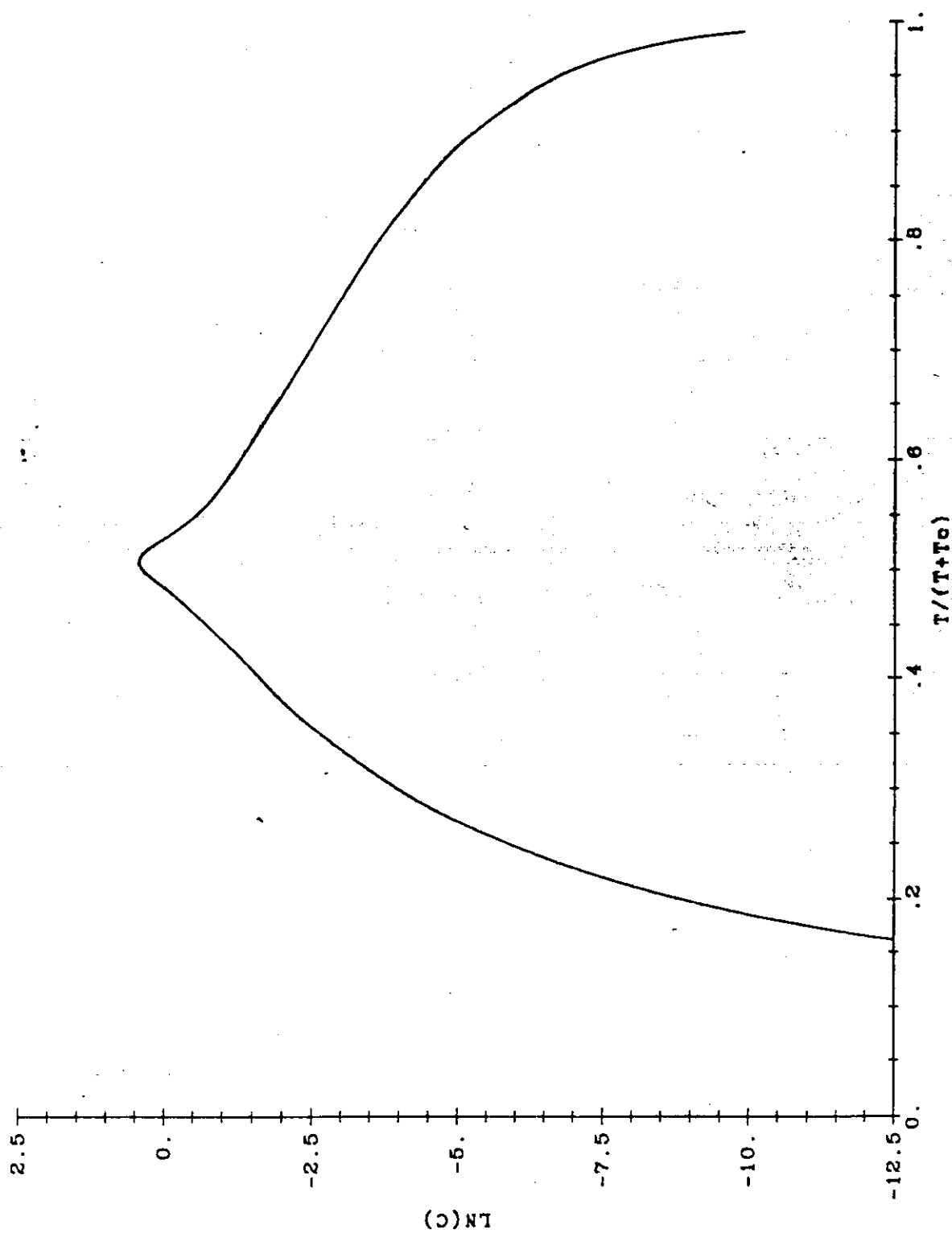


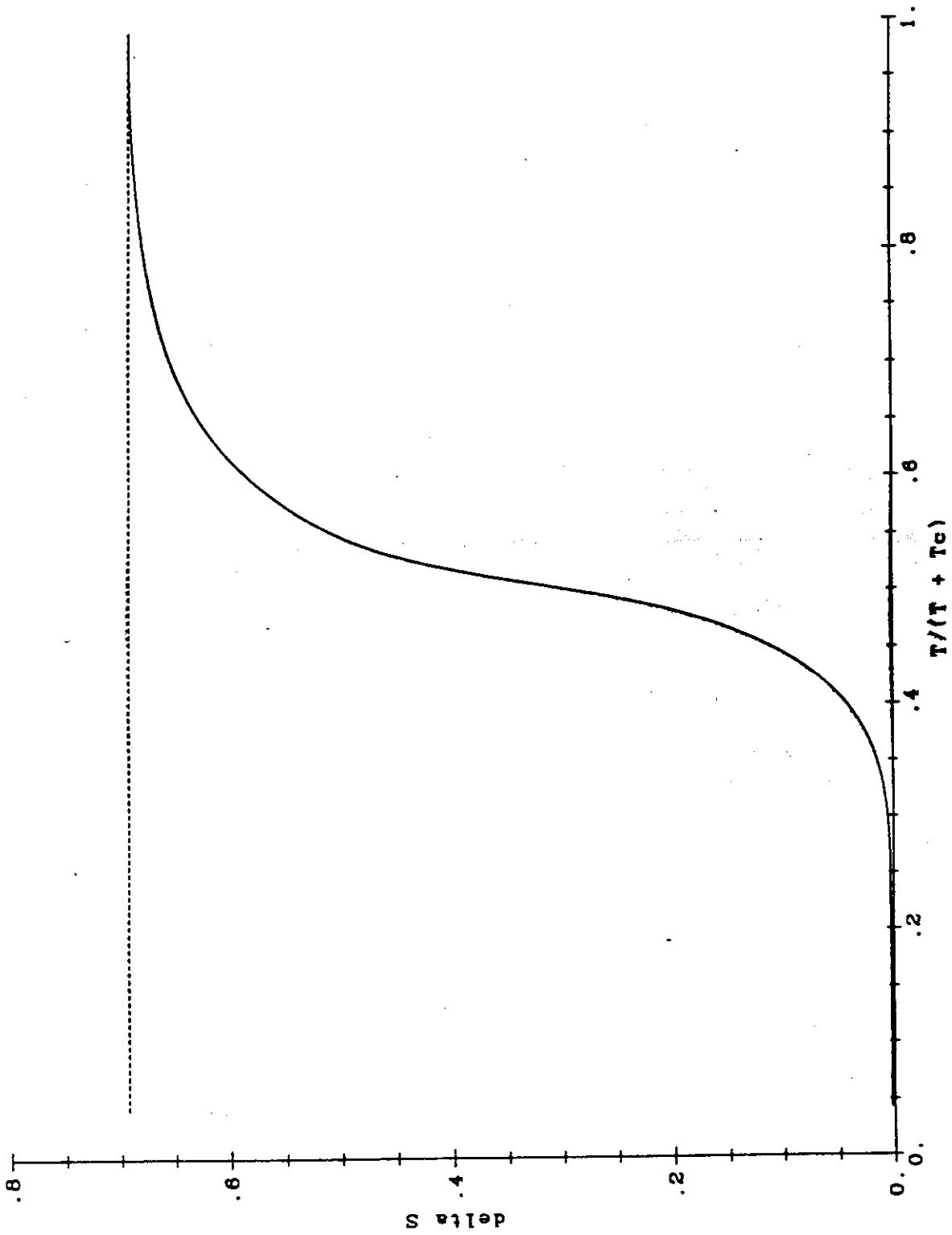
$$\{K_x | x = 1, \dots, 8\}$$

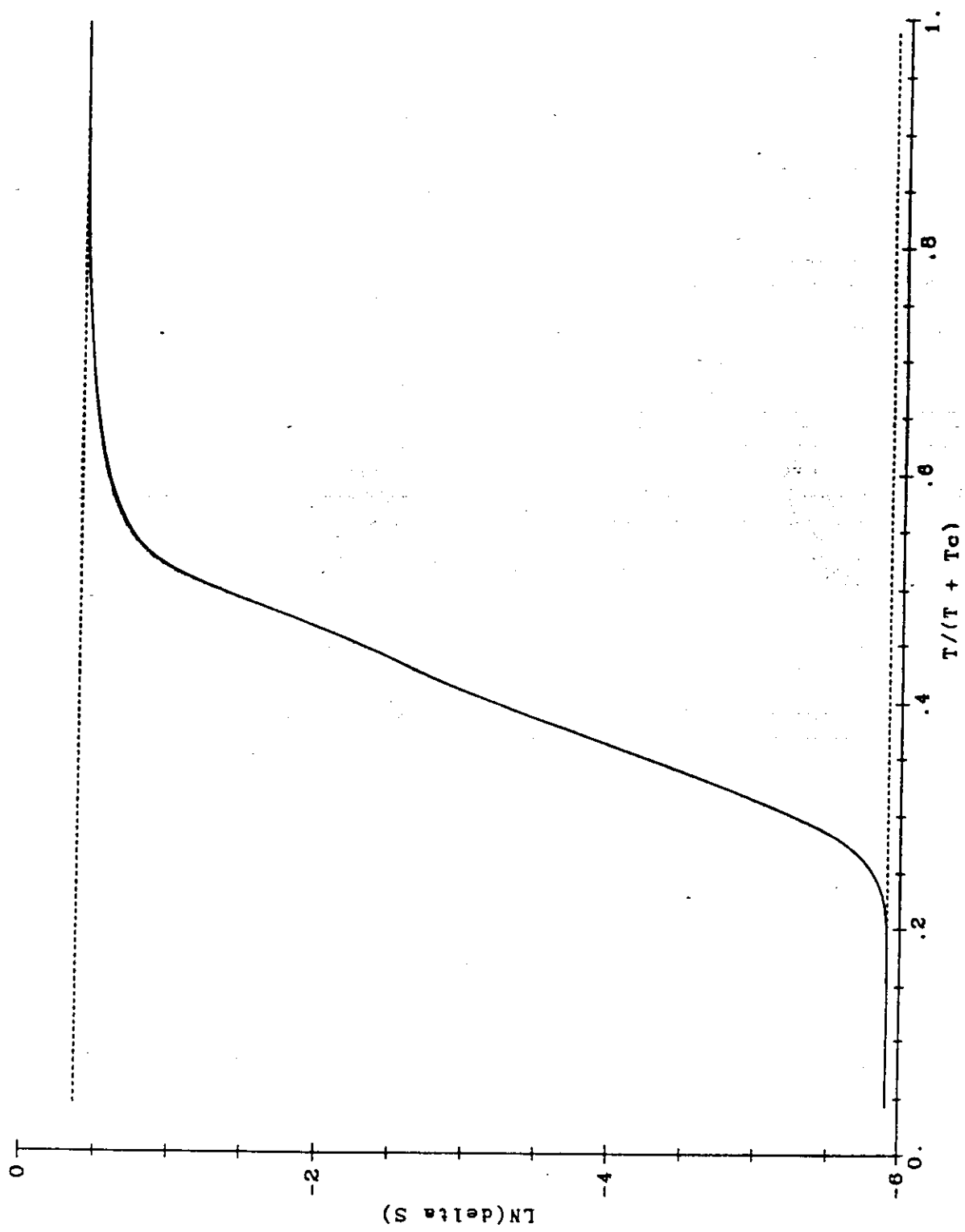




EXACT







DEFINITION(S) OF PROBABILITY

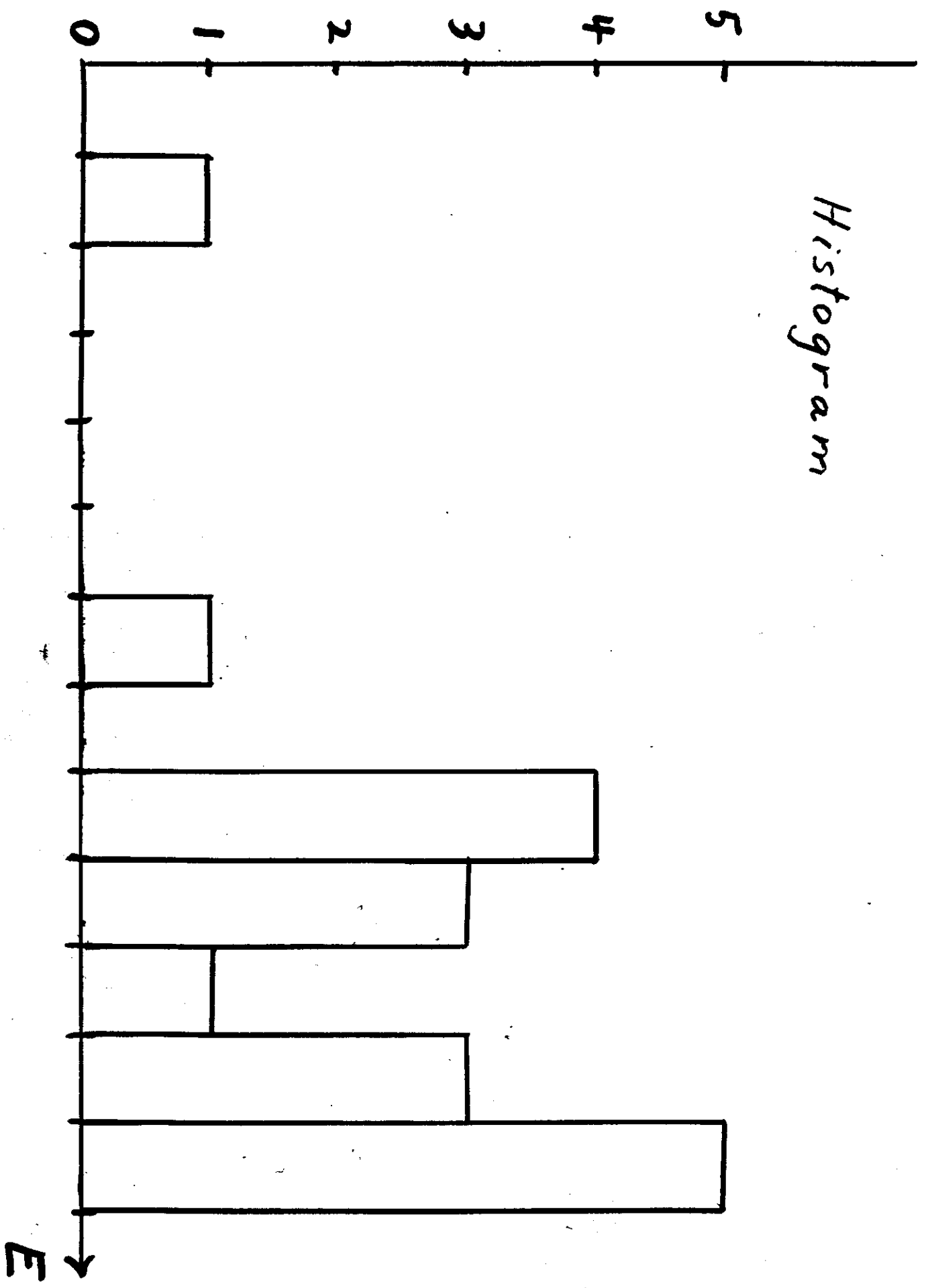
Frequentist

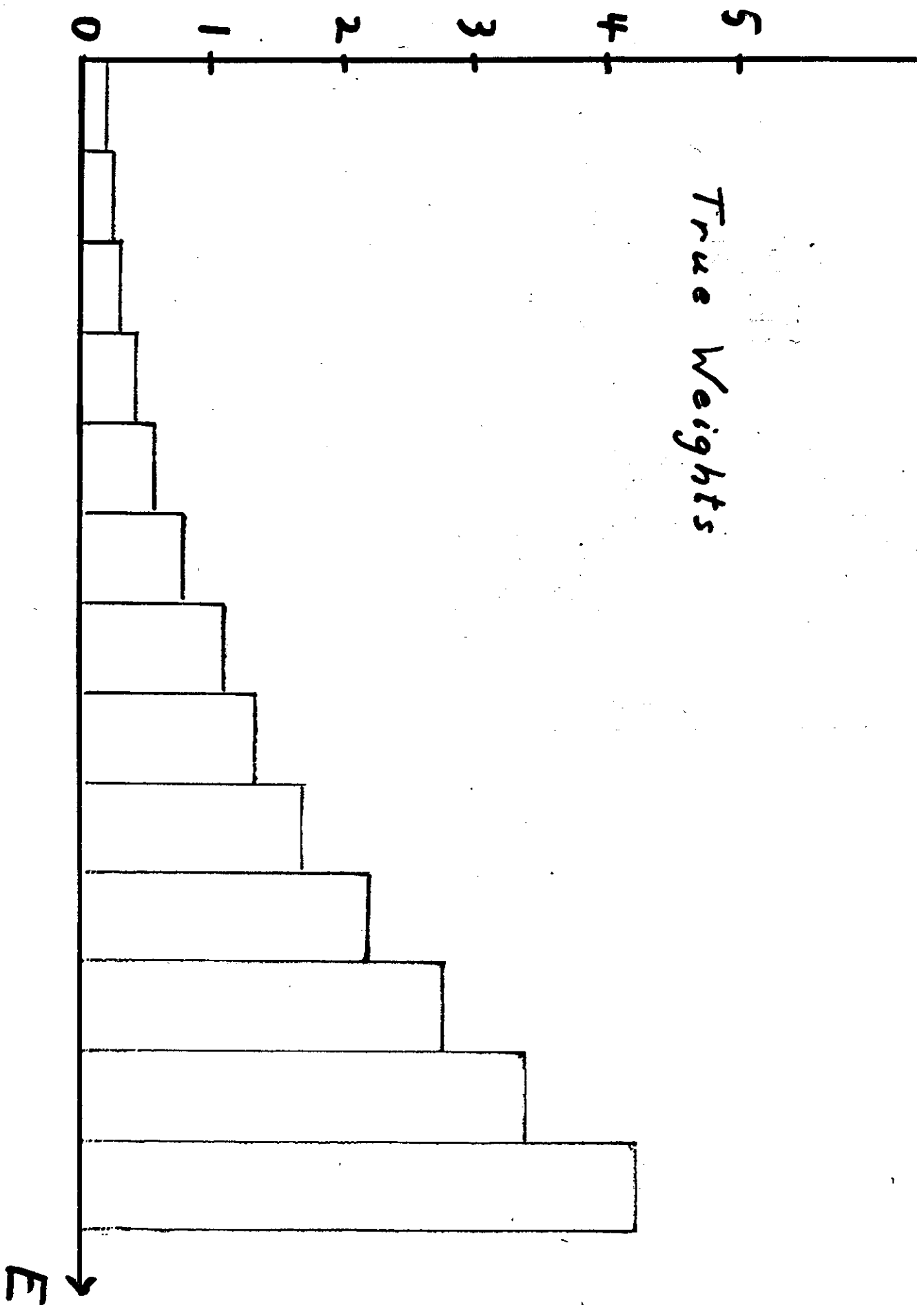
Ratio of "successes" to trials
in the limit of an infinite
number of trials

Bayesian

A description of your
knowledge of the result
of a trial

Histogram





Bayes' Theorem

$$P(\theta|x) = \frac{P(x|\theta) P(\theta)}{P(x)}$$

Interpretation:

x - experimental results

θ - theoretical parameters

$P(\theta)$ prior knowledge

$P(\theta|x)$ is what we know
about θ from both
our prior knowledge
and the
experimental results

n independent trials

θ = probability of success

$$P(k|\theta) = \binom{n}{k} \theta^k (1-\theta)^{n-k}$$

If we observe k successes,

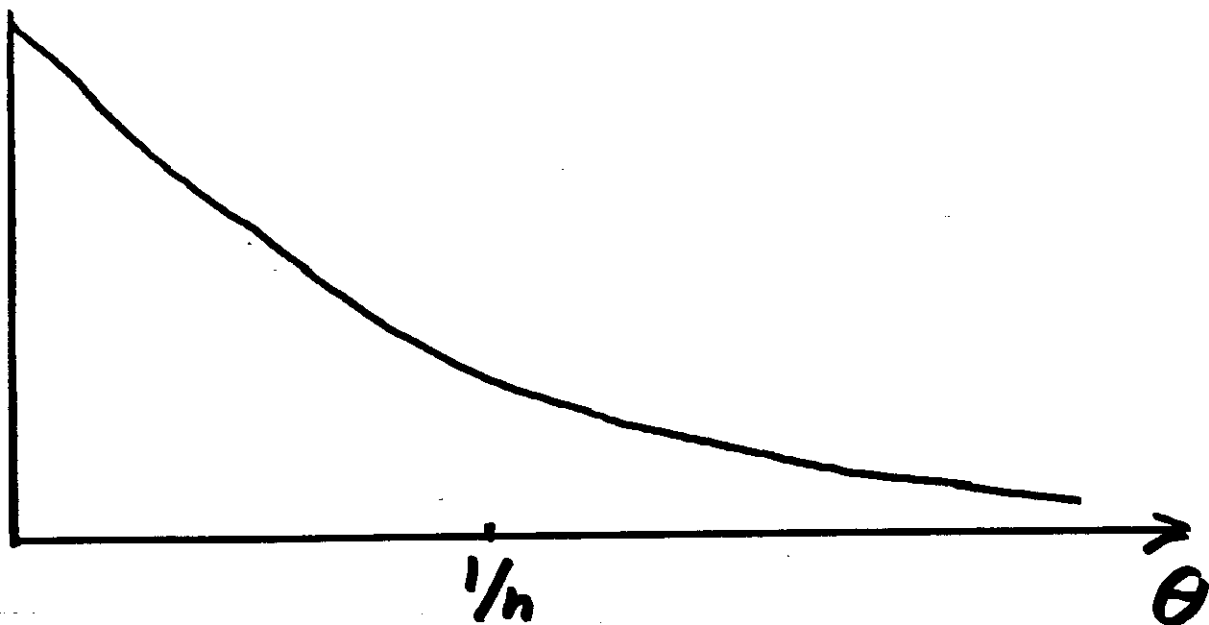
what do we know about θ ?

RJR: Assume $P(\theta) = 1$ for $\theta \in [0, 1]$

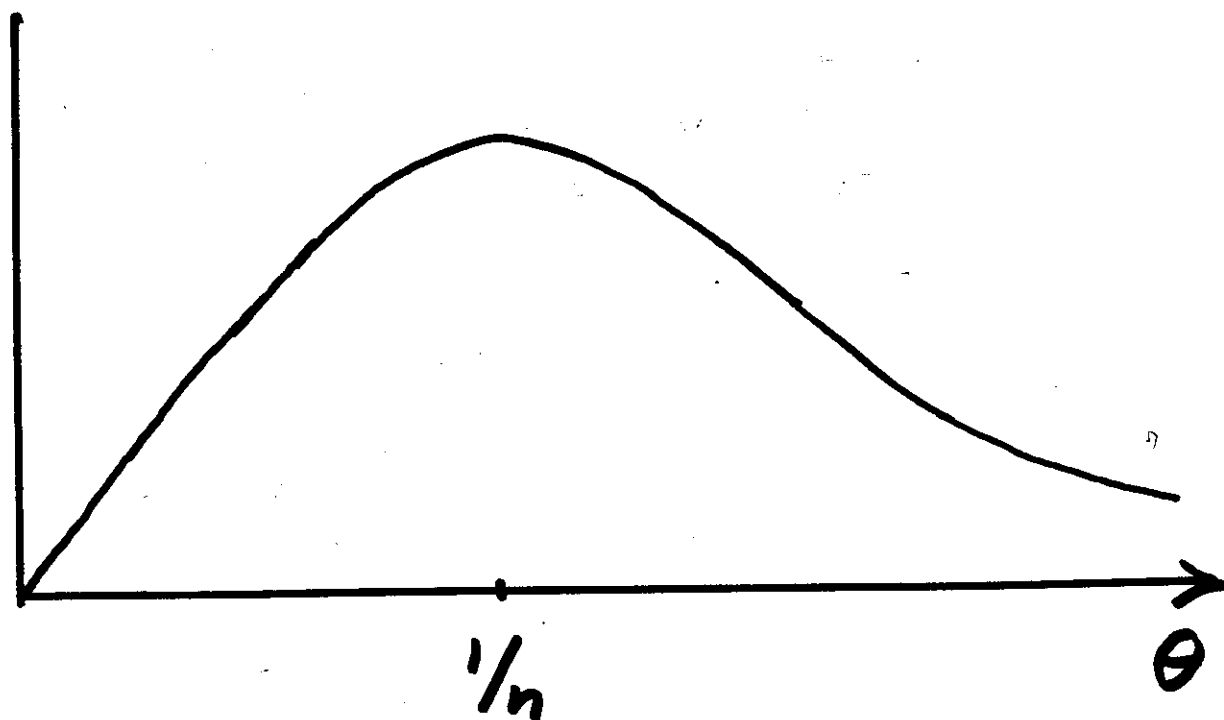
AYES: $P(\theta|k) \sim P(k|\theta) P(\theta)$

$$P(\theta|k) \sim \theta^k (1-\theta)^{n-k}$$

$$P(\theta | k=0)$$



$$P(\theta | k=1)$$



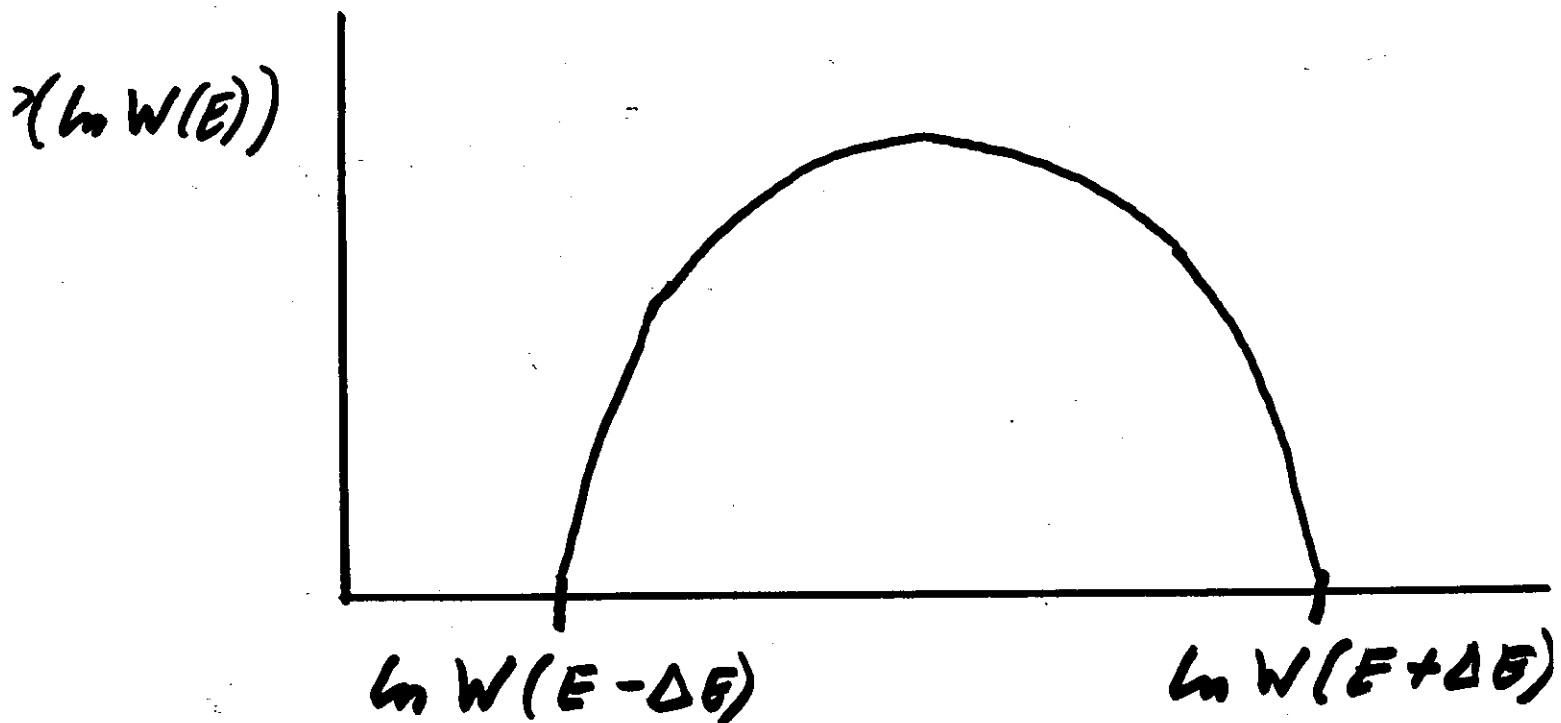
"Smoothness" of $W(E)$

$$W(E - \Delta E) < W(E) < W(E + \Delta E)$$

In critical region

$$\ln W(E) \sim \frac{1}{2} [\ln W(E - \Delta E) + \ln W(E + \Delta E)]$$

Can express this as a "prior" factor



Produces estimates for $W(E)$
that are qualitatively correct

Transition Matrix Monte Carlo

Shing-Te Li

Brian Diggs

J. Kadane

C. Genovese

"Broad histogram method"

P. M. C. de Oliveira

T. J. P. Penna

H. J. Herrmann

Brazilian Journal of Physics

26, 677 (1996)

Transition Matrix

$T = \infty$

$n_{\delta E}$ = number of spins that will
change the energy by δE
when $\sigma_i \rightarrow -\sigma_i$

$$T_{E, \delta E} \equiv \langle n_{\delta E} \rangle / N$$

$$N = L^d$$

gives the probability that
the next spin flip
will change the energy
from E to $E + \delta E$

$$\sum_{\delta E} T_{E, \delta E} = 1$$

Finite Temperature

Transition Matrix

$$T_{E, \delta E}^P \begin{cases} N^{-1} \langle n_{SE} \rangle & [\delta E \leq 0] \\ N^{-1} \langle n_{SE} \rangle e^{-\beta \delta E} & [\delta E > 0] \\ N^{-1} \langle n_0 \rangle + \sum_{SE > 0} N^{-1} \langle n_{SE} \rangle (1 - e^{-\beta \delta E}) & [\delta E = 0] \end{cases}$$

$$\sum_{\delta E} T_{E, \delta E}^P = 1$$

Probability of move being rejected

$$R = \sum_{\delta E > 0} \frac{\langle n_{SE} \rangle}{N} (1 - e^{-\beta \delta E})$$

Density of States

$W(E)$

Infinite temperature

$$\sum_{SE} T_{E-SE, SE} W(E-SE) = W(E)$$

Finite temperature

$$P_{\beta}(E) = \frac{1}{Z} W(E) e^{-\beta E}$$

$$\sum_{SE} T_{E-SE, SE}^{\beta} P_{\beta}(E-SE) = P_{\beta}(E)$$

The TTT Identity

Detailed balance: $W_1 T_{12} = W_2 T_{21}$

Transitions among three states:

$$W_1 T_{12} \quad W_2 T_{23} \quad W_3 T_{31} = W_1 T_{13} \quad W_3 T_{32} \quad W_2 T_{21}$$

$$\Rightarrow T_{12} T_{23} T_{31} = T_{13} T_{32} T_{21}$$

$$T_{E-\Delta E, \Delta E} \quad T_{E, \Delta E} \quad T_{E+\Delta E, -2\Delta E} =$$

$$T_{E, +2\Delta E} \quad T_{E+2\Delta E, -\Delta E} \quad T_{E+\Delta E, -\Delta E}$$

where ΔE is the interval between energy levels

Transition Matrix

Combining data from simulations
at different temperatures
- or different ensembles -
is trivial

Used with cluster MC
histograms and transition matrix
give same accuracy

N-fold Way

For each configuration,

$$R = \sum_{\delta E > 0} \frac{n_{\delta E}}{N} (1 - e^{-\beta \delta E})$$

is the probability of rejected moves

Rejected moves retain current state

$\Rightarrow$ weight current configuration with $(1 - R)^{-1}$

and make a δE spin flip with

$$\frac{n_{\delta E}}{N} (1 - R)^{-1} \begin{cases} 1 & \delta E \leq 0 \\ e^{-\beta \delta E} & \delta E > 0 \end{cases}$$

Transition matrix

with N -fold way

- Same accuracy as with cluster Monte Carlo
- Also works for models with frustration

Correlation time

given by second eigenvalue

$$\sum_{SE} T_{E-SE, SE}^P \phi_i^P(E-SE) = \lambda_i^P \phi_i^P(E)$$

$$\bar{\tau}(\beta) = -1 / \ln \lambda \quad [\text{MCS}]$$

$$\text{Expect} \quad \bar{\tau} \sim L^3 \quad [\text{MCS/s}]$$

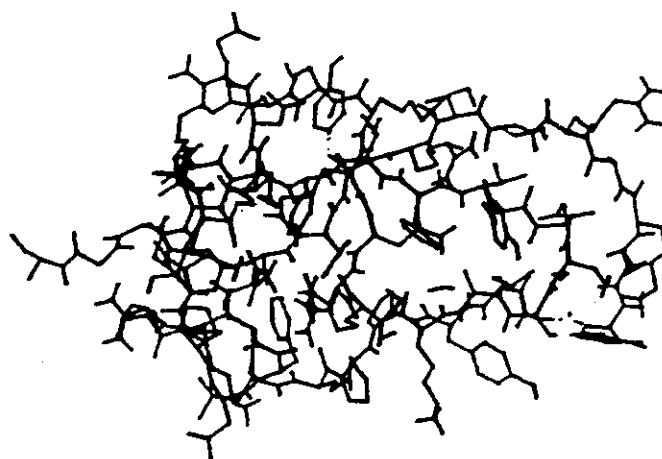
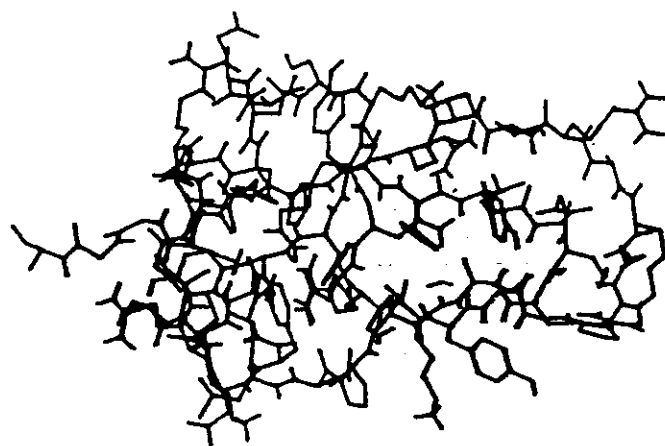
$$\text{Get} \quad \bar{\tau} \sim \ln L$$

Computer Simulation of Biological Molecules

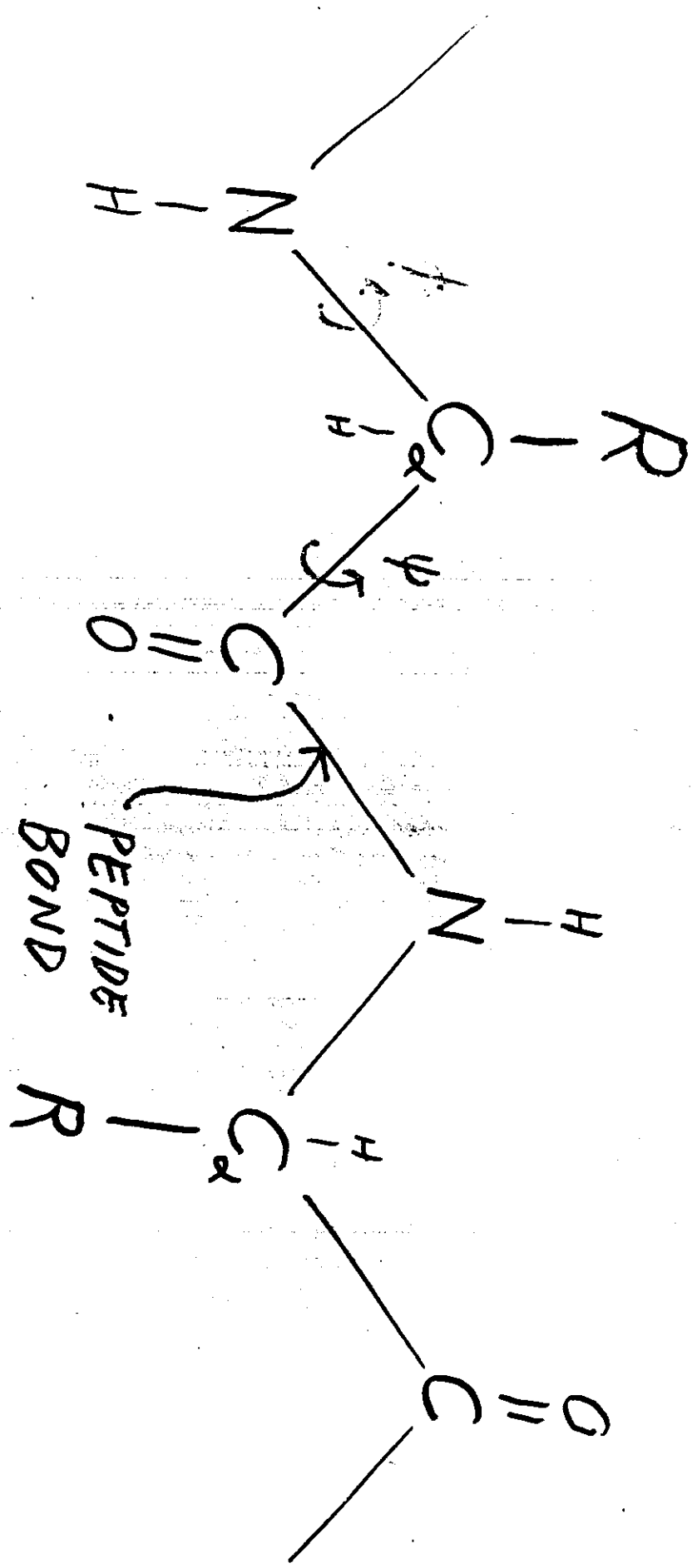
Robert H. Swendsen

Djamal Bouzida

Shankar Kumar



BPTI



Empirical potential energy function

- Bond Stretching:

$$\frac{1}{2} \sum_{bonds} K_b (b - b_{eq})^2$$

- Angle Bending:

$$\frac{1}{2} \sum_{angles} K_\theta (\theta - \theta_{eq})^2$$

- Torsion:

$$\frac{1}{2} \sum_{dihedrals} V_\varphi [1 + \cos(n\varphi - \delta)]$$

- Lennard-Jones, and electric:

$$\sum_{non-bonds} \left[\frac{A}{r^{12}} - \frac{B}{r^6} + \frac{q_i q_j}{\epsilon r} \right]$$

Monte Carlo

MC is a *stochastic* method:

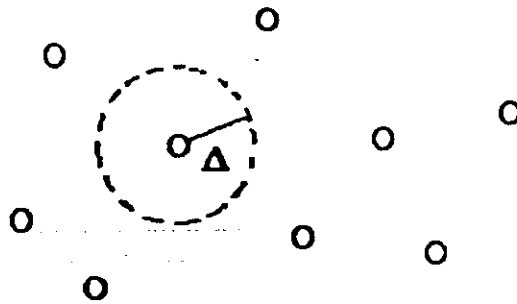
- Generate a sequence of system configurations which is distributed according to the Boltzmann distribution.
- Equilibrium properties are found by computing appropriate averages over the resulting set of configurations.
- **Metropolis Algorithm:**
 - Pick an atom, make random move.
 - Compute ΔE .
 - If $\Delta E < 0$, the move is accepted.
 - If $\Delta E > 0$, the move is accepted with probability $\exp(-\beta\Delta E)$.
- Ideal acceptance ratio is about 50%.

Classical Paper:

Metropolis et. al., Jour. Chem. Phys., Vol. 21 (6), 1087 (1953)

Simulation of liquids

How does one simulate a liquid?



Atoms are moved randomly inside spheres of equal radii.

System is:

- homogeneous.
- isotropic.

Differences in simulating liquids and molecules

- **Inhomogeneity**

- Each atom of the molecule sees a different environment.
- Wide range of variation in the atomic fluctuations.

- **Anisotropy**

- Highly anisotropic potentials.
- Local effective potentials **change** during simulation.

$d=1$ SHO

$$E = \frac{1}{2} k x^2$$

Monte Carlo move Δx

$$\delta \geq \Delta x \geq -\delta$$

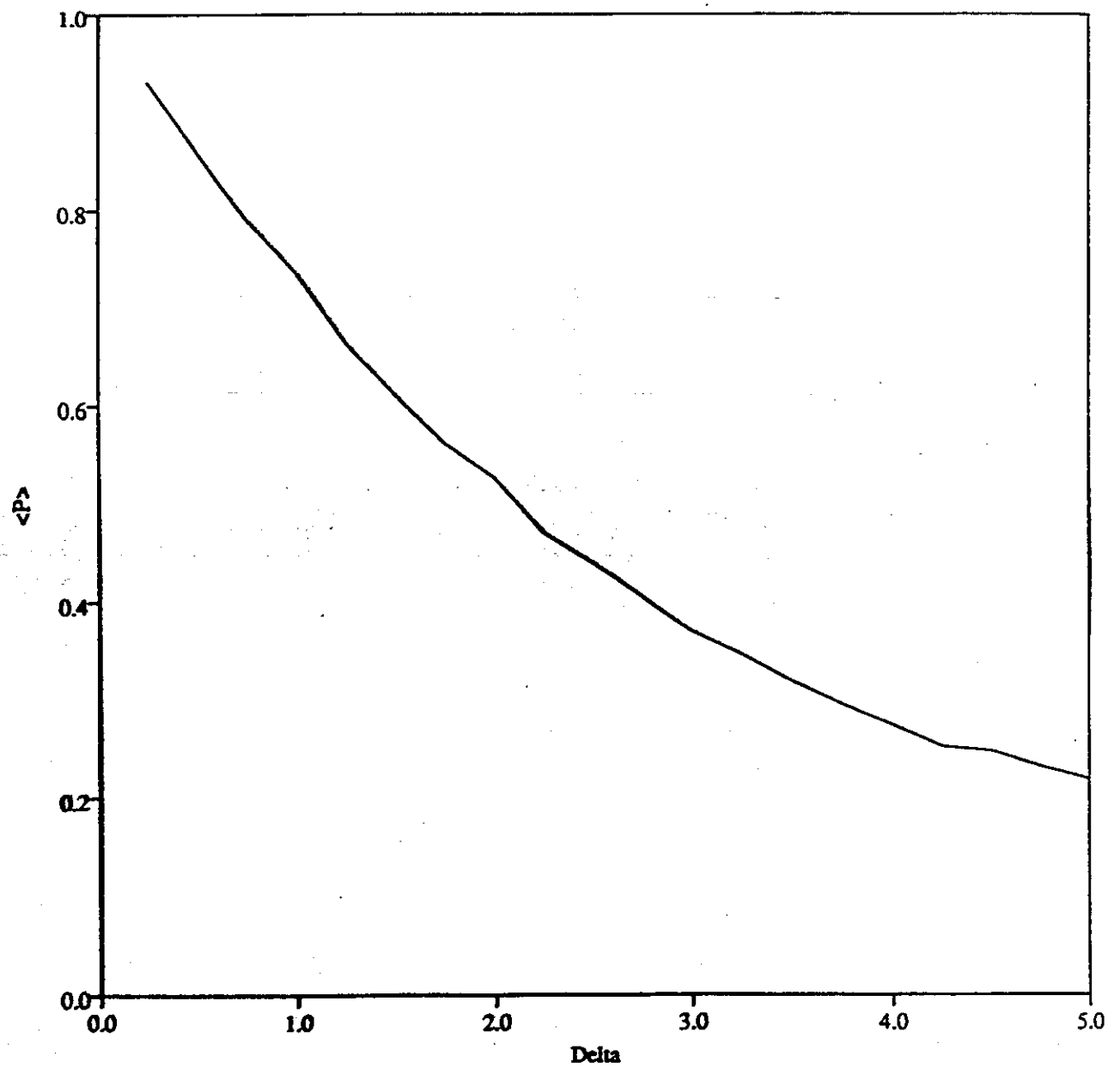
δ = optimal step size

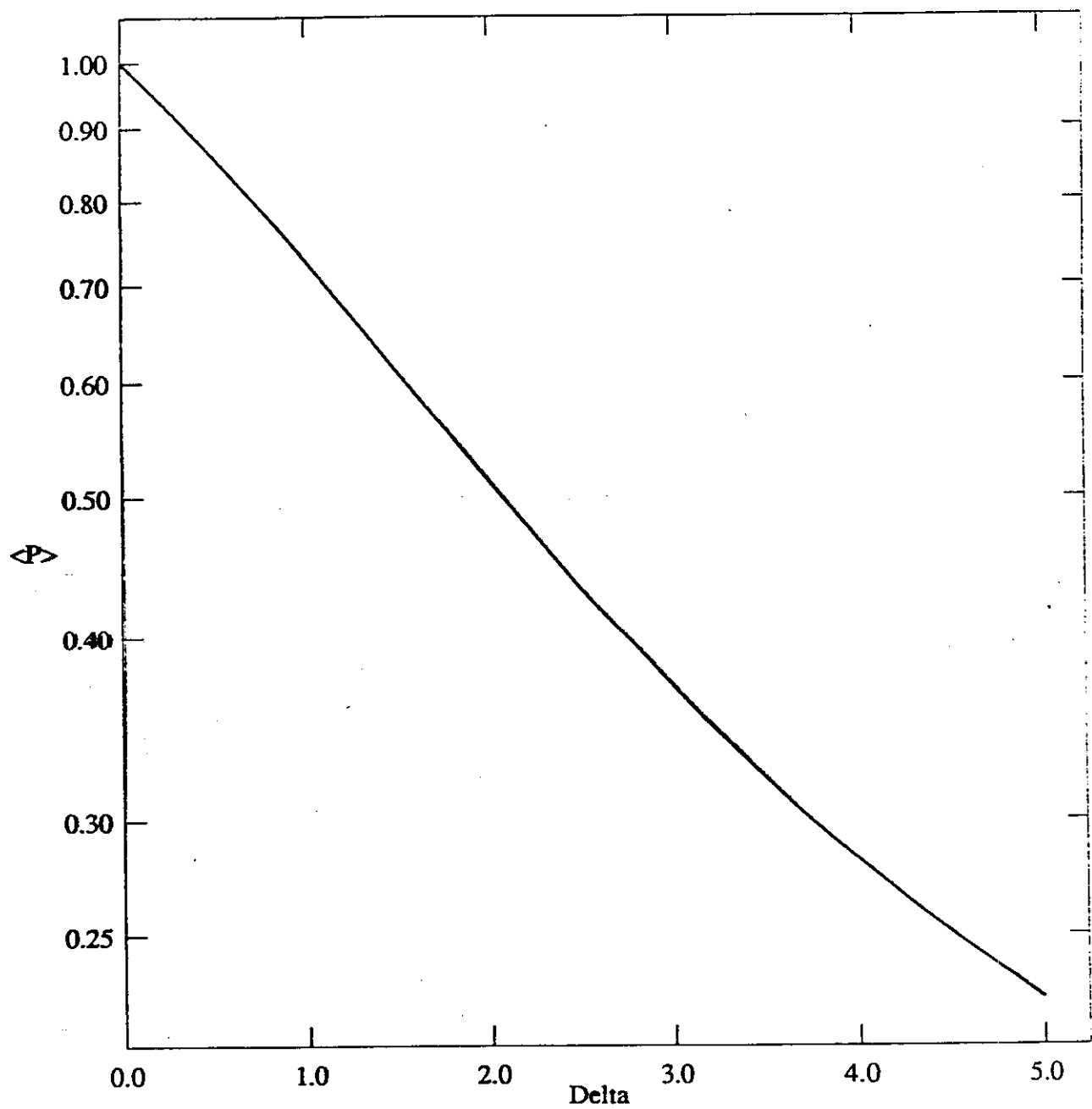
Scaling relation:

$$\frac{1}{2} \beta k \delta^2 = F^2$$

$$V(x) = x^2 \quad T = 1.0$$

MC

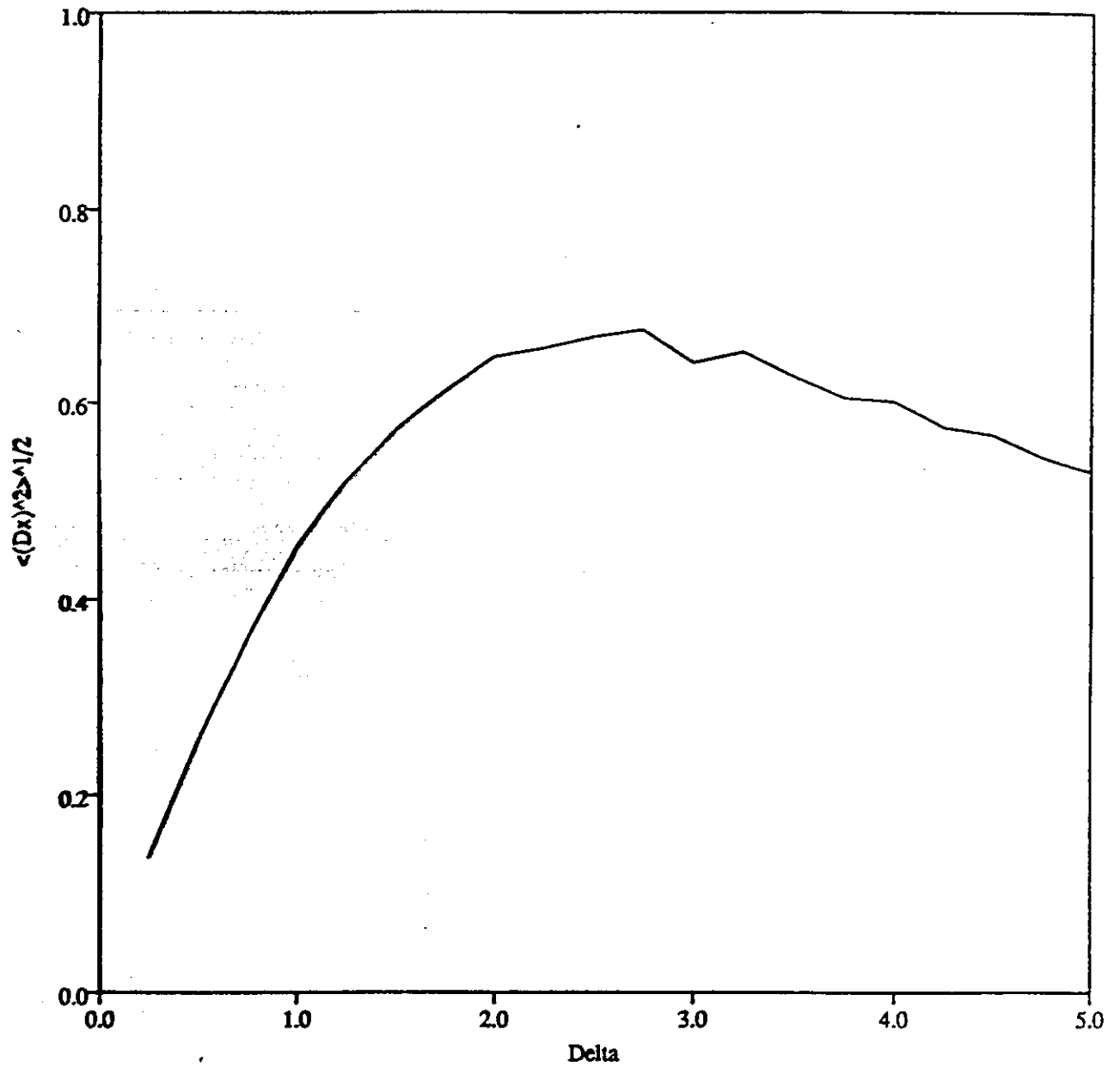




$$V(x) = x^2$$

$$T = 1.0$$

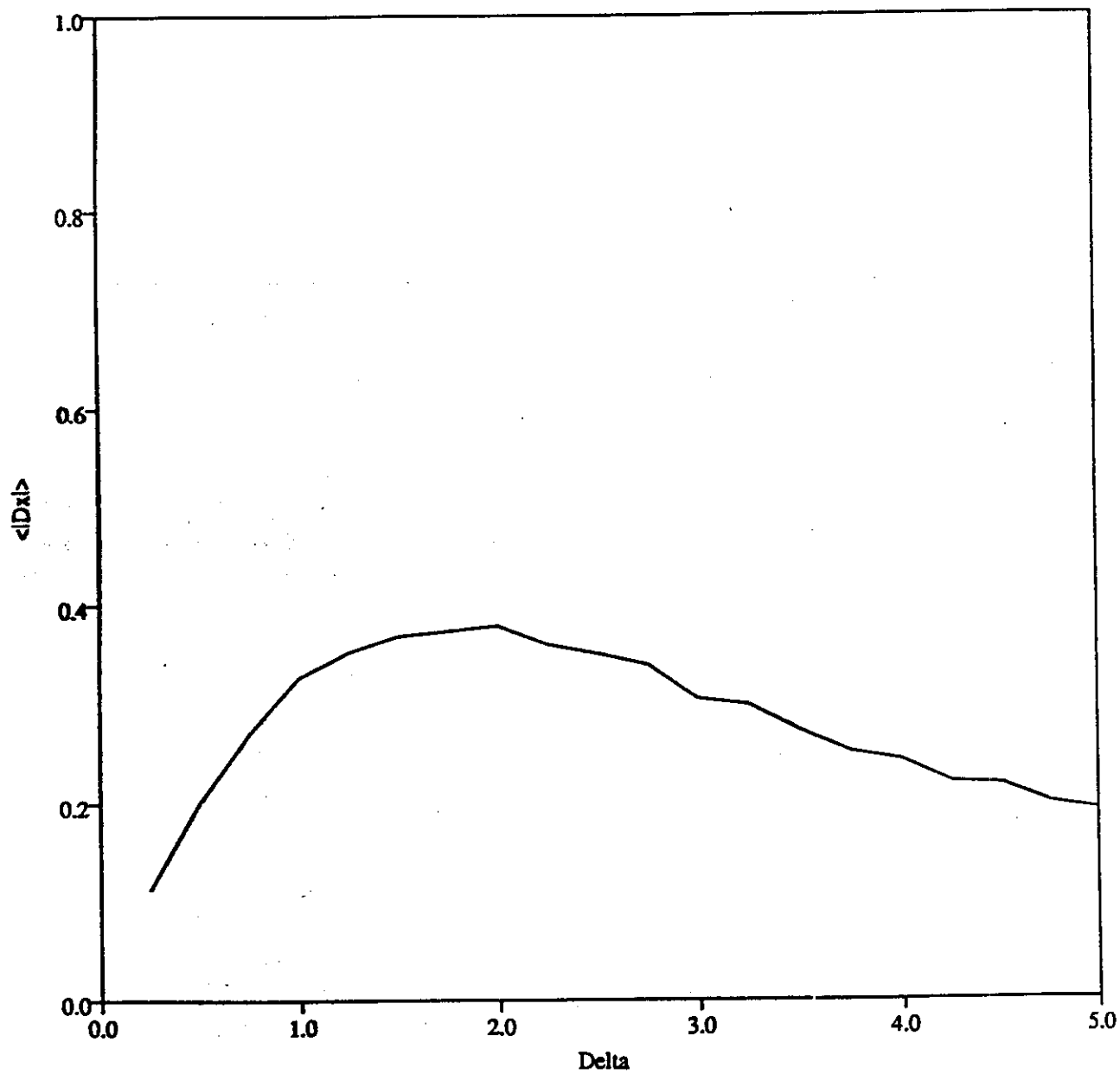
MC

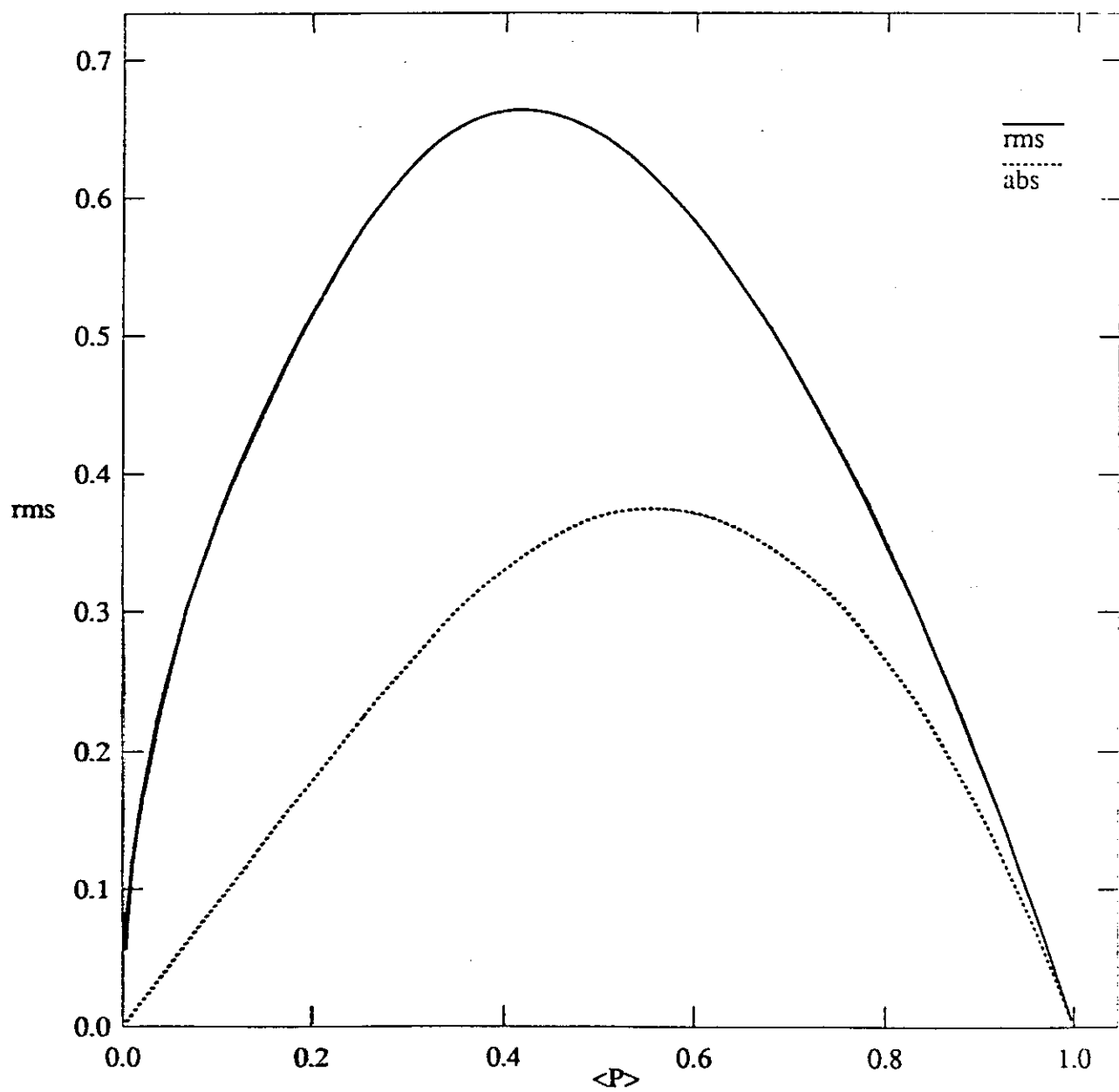


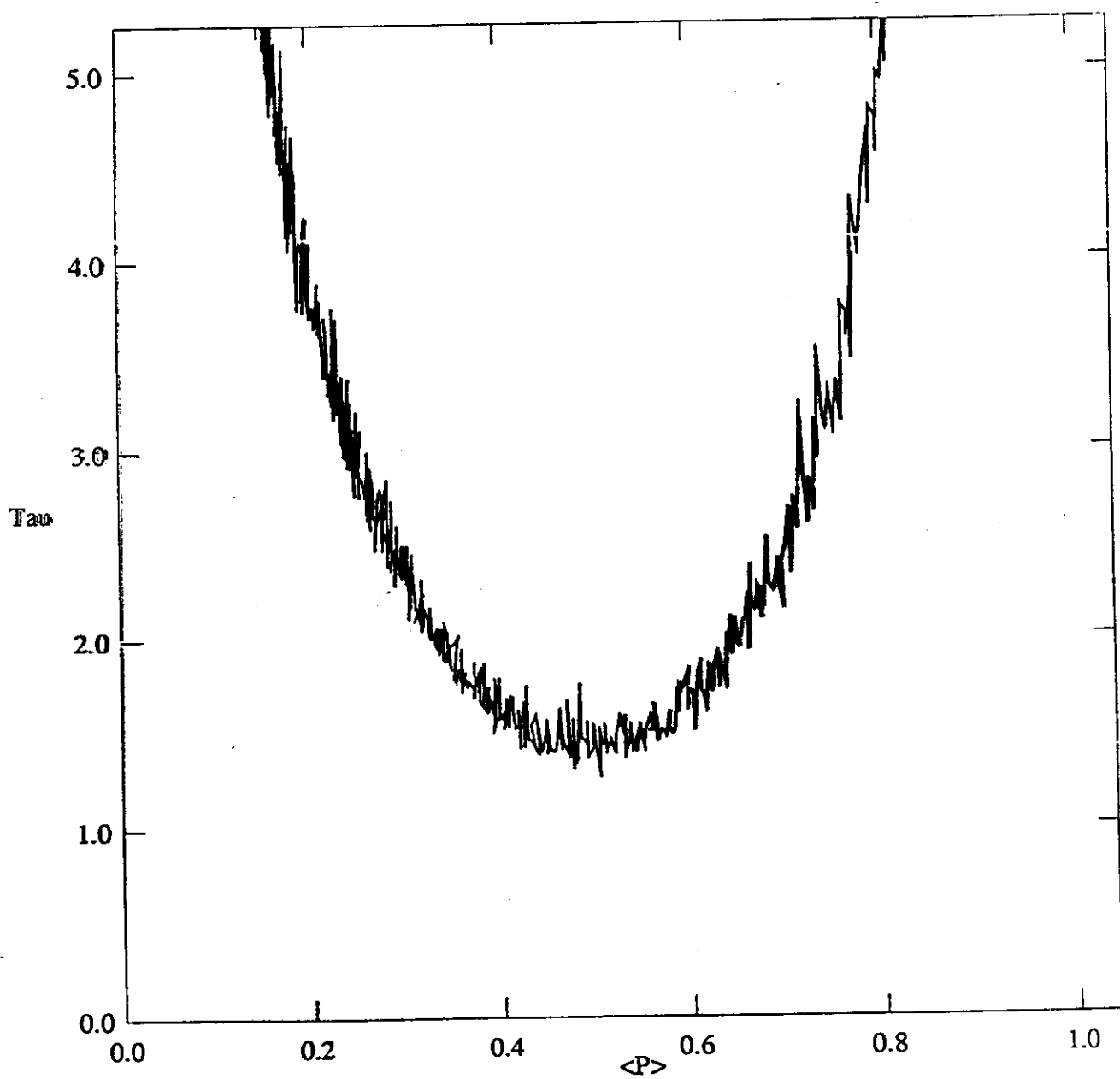
$$V(x) = x^2$$

$$T = 1.0$$

MC







Acceptance Ratio Method

$$P \approx \exp[-\Delta/\Delta_0]$$

Ideal ratio $\leftrightarrow$ ideal step size

$$P_i = \exp[-\Delta_i/\Delta_0]$$

$$\Rightarrow \Delta_{\text{new}} = \Delta_{\text{old}} \ln P_i / \ln P$$

Actually use:

$$\Delta_{\text{new}} = \Delta_{\text{old}} \frac{\ln(a P_i + b)}{\ln(a P + b)}$$

Dynamically Optimized Monte Carlo

DONC

$$E = \frac{1}{2} k x^2$$

$$\Delta E = \frac{1}{2} k (x + \Delta x)^2 - \frac{1}{2} k x^2$$

$$\Delta E = k x \Delta x + \frac{1}{2} k (\Delta x)^2$$

$[\dots]$ $\equiv$ average over all attempted moves

$$[\Delta x] = 0$$

$$[\Delta E] = \frac{1}{2} k [(\Delta x)^2]$$

$$\frac{1}{2} \beta k \delta^2 = F^2$$

$$\delta = F \left(\frac{[(\Delta x)^2]}{\beta [\Delta E]} \right)^{\frac{1}{2}}$$

DOMC for $d > 1$

$$E = \frac{1}{2} \sum_{i,j} k_{ij} x_i x_j$$

MC moves $\{z_i\}$ in an ellipsoid

$$z_i = \sum_{j=1}^d D_{ij} \xi_j$$

$\{\xi_i\}$ is a random vector from the unit sphere.

Scaling relation:

$$F^2 = \frac{1}{2} \rho \bar{D}^T \cdot \bar{k} \cdot \bar{D}$$

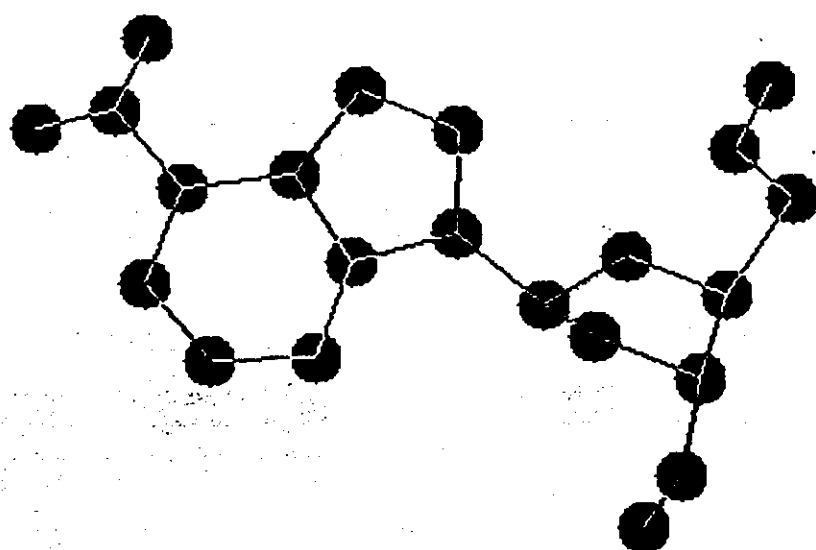
$$[\Delta E z_\ell z_m] = \frac{1}{2} \sum_{i,j} k_{ij} [z_i z_j z_\ell z_m]$$

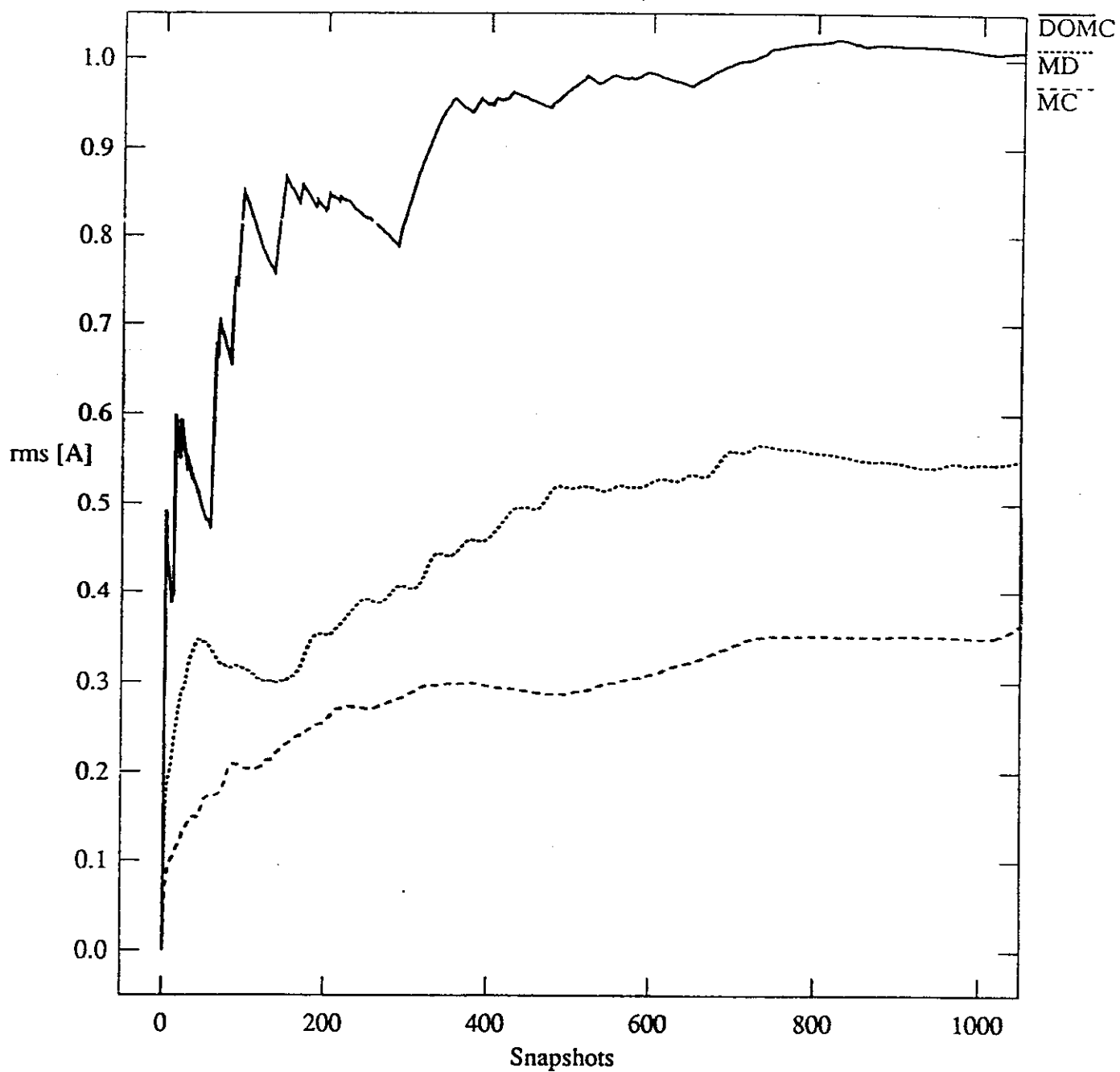
Solve for $\{k_{ij}\}$

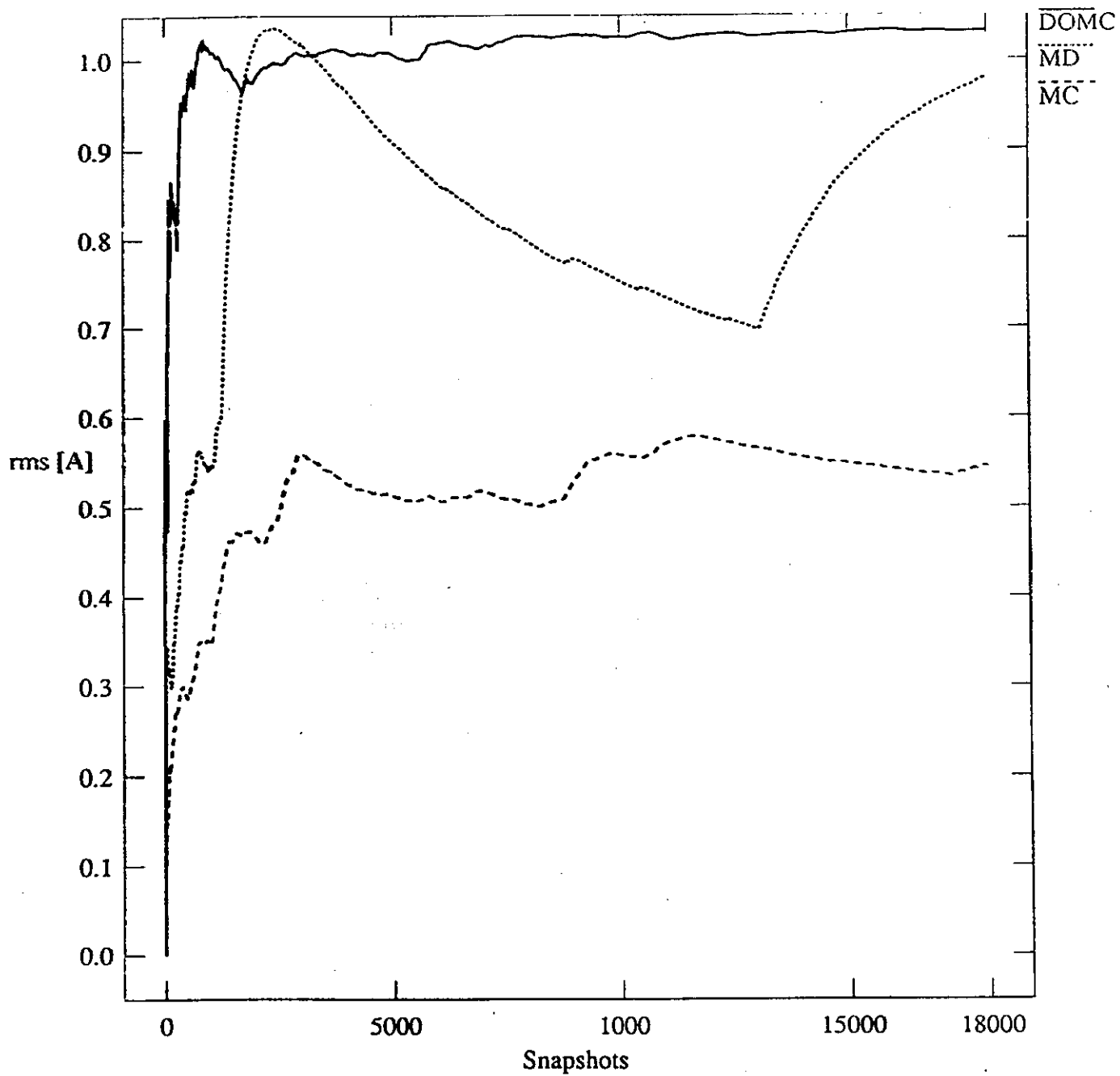
λ_ℓ = eigenvalue of $\{k_{ij}\}$

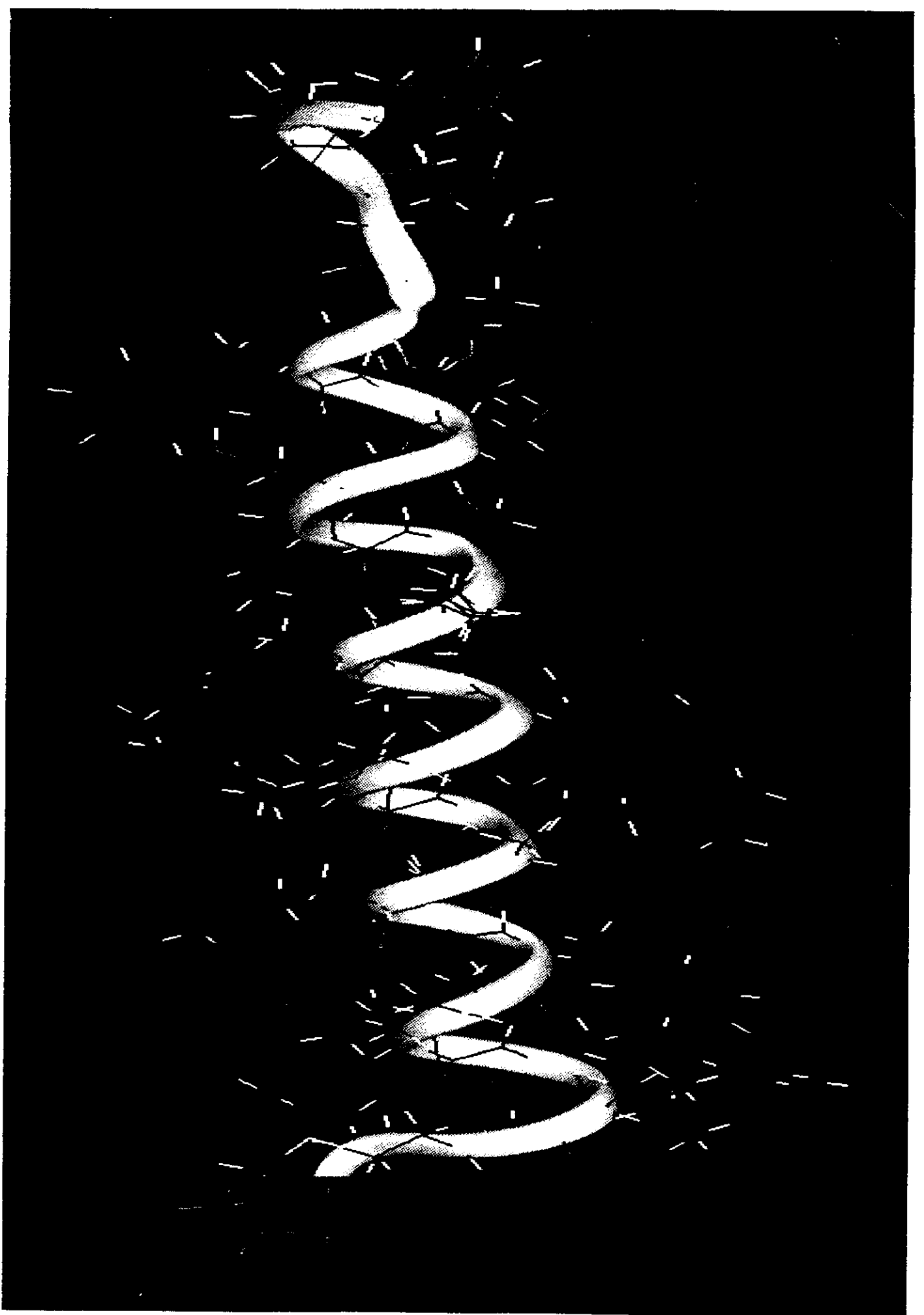
$V_{i\ell}$ = eigen vector of "

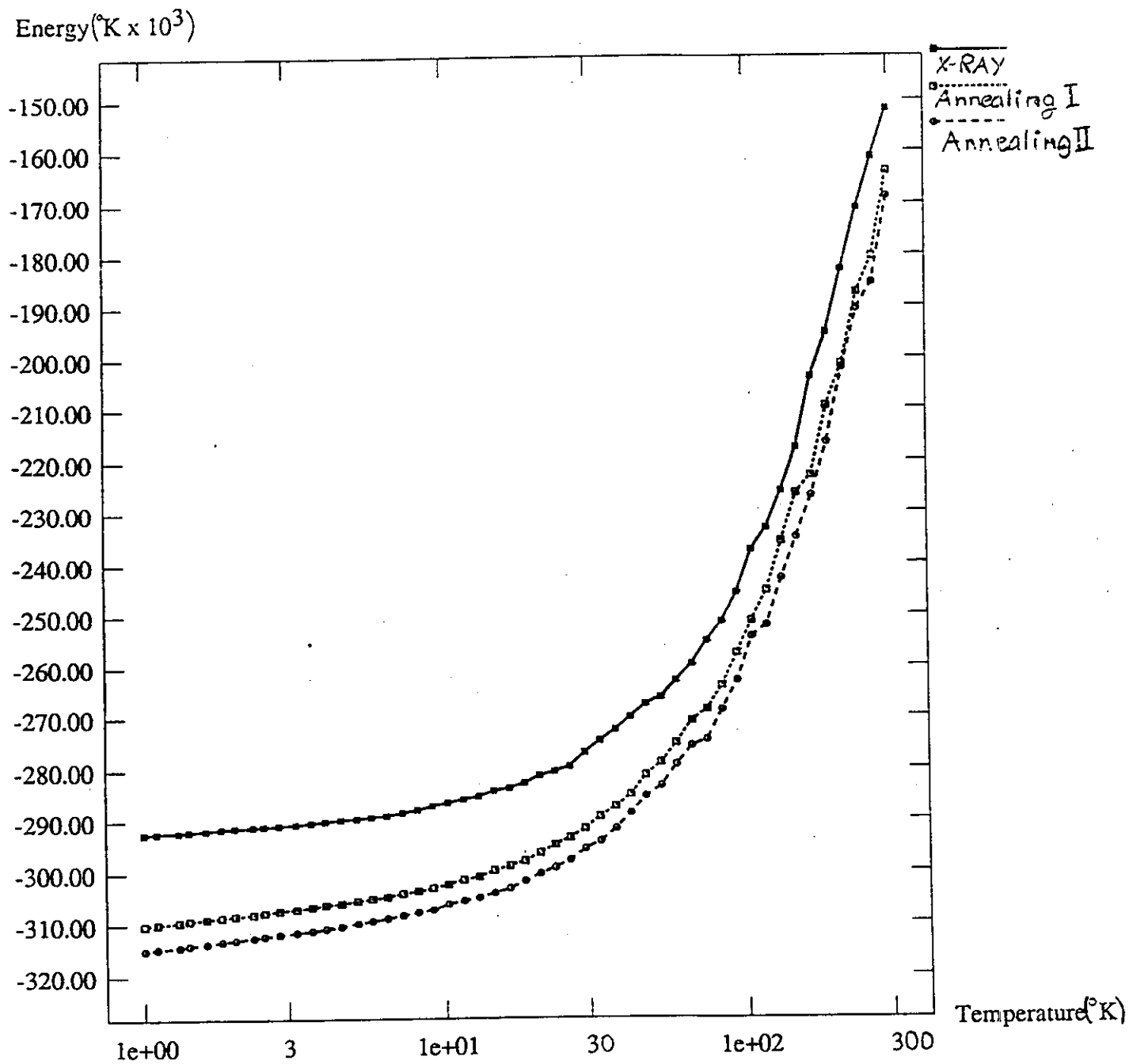
$$D_{i\ell} = F \left(\frac{2}{\rho \lambda_\ell} \right)^{\frac{1}{2}} V_{i\ell}$$

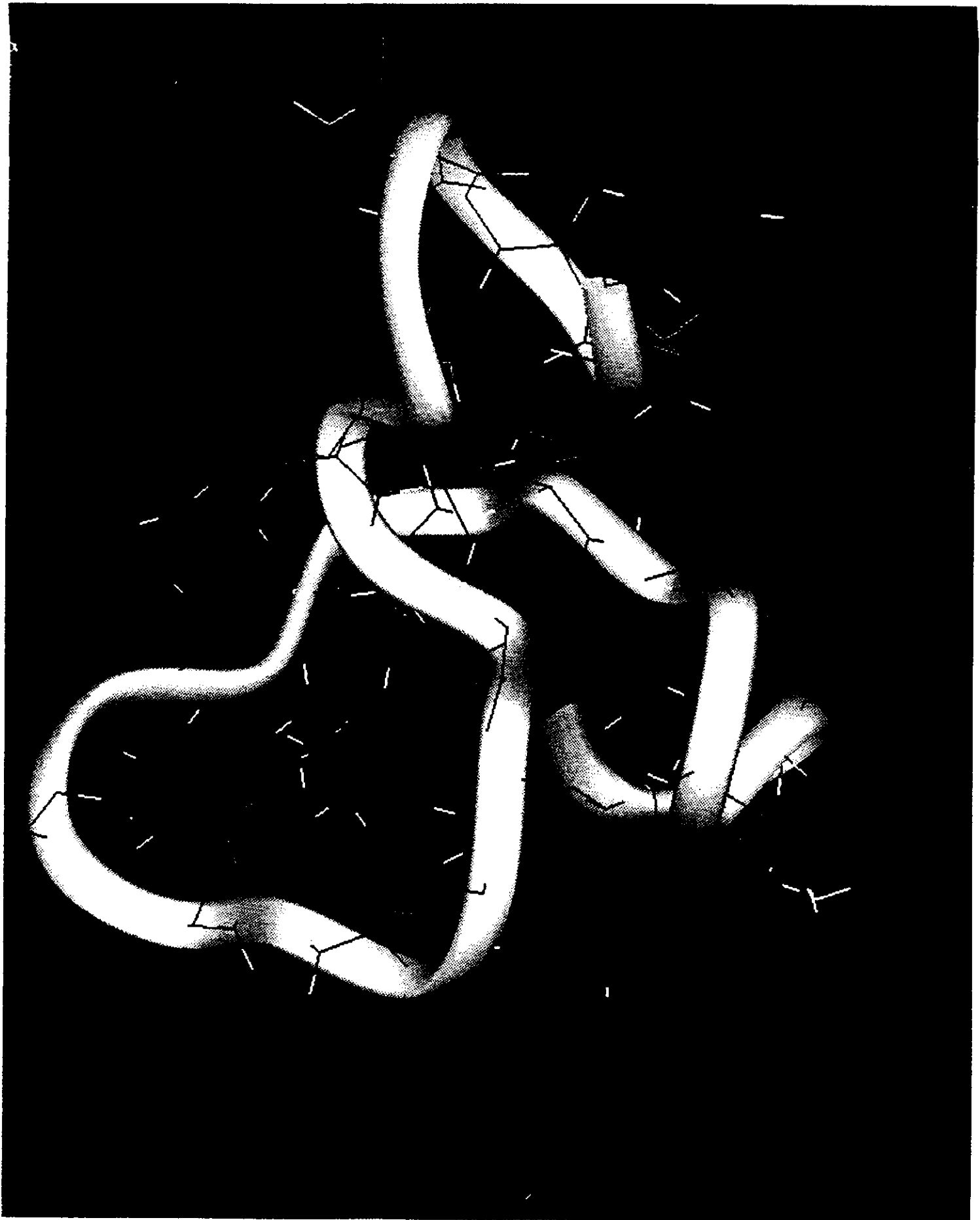












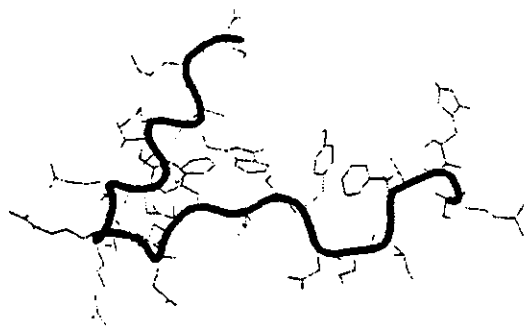


Figure 24: structure of glucagon after annealing

molecule. In the remaining cases the structure analysis program detected no helical regions. Figures 25 and 27 show the results of one of these runs.

Energetically, the helical structure lies about 20 kcal/mol below the energy of the partially folded structures. The random coils had an even higher energy, about 30 kcal/mol above the helical structure.

Even though the program did not reproduce the X-ray structure the results found agree rather well with experimental measurements by Panijpan and Gratzer [27] and Gratzer et al. [28]. From NMR studies, measurements of the temperature dependence of optical rotary dispersion and thermal difference measurements they conclude that that in dilute solution, glucagon is largely a random chain which contains about one turn of α -helix. They predict that the surmised α -helical segment is formed by residues 21 or 22 to 26 near the C-terminal. This is the region in which the simulated molecule showed the largest probability of forming an α -helical structure. The experiments also support a hypothesis brought forth by the authors of [28] that the molecule at room temperature is at the high end of a transition between a helix and a random coil state. The experiments show that the apparent helix content of glucagon in dilute solution increases as the temperature is lowered. The authors make a prediction that glucagon should fold into the helical form