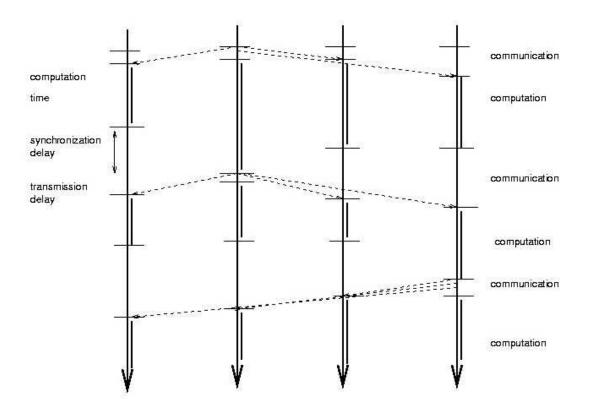
A model for Entropy of Parallel Execution

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5TH ANNUAL LLU ALGORITHMS IN PARTICLE IMAGING AND TREATMENT WORKSHOP

Irregular execution and communications



Parallel execution:

Definition 2.1. A parallel state is a pair $S_{G,J} = (x_J, D_J^G)$, indexed by a step counter J which is a multi-index $J = J_G = \{(j_p) \mid p \in G\}$.

Definition 2.2. A parallel transition is a relation $S_{G,J} \rightarrow S_{H,K}$ between a pair of parallel states such that

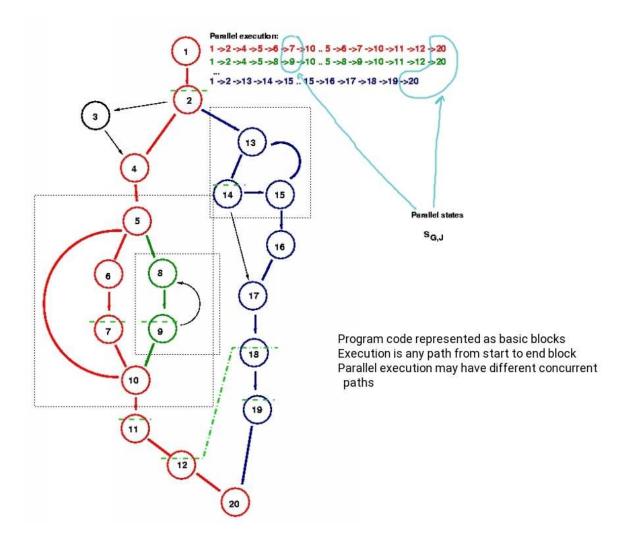
- G ∩ H ≠ ∅
- for each $p \in G \cap H$, we have $k_p j_p \le 1$, and for at least one $p \in G \cap H$ we have $k_p j_p > 0$.

Definition 2.3. Let $T=(S_{G^1,J^1}\to S_{G^2,J^2}\to \ldots)$ be any sequence of parallel transitions relating parallel states. (Note that $G_i\cap G_{i+1}$ needs to contain at least one element, but there is no other restriction on the G_i). The p-trace of T is the list of serial state indices $\tau_p = (j_p^{i_1}, j_p^{i_2}, \dots, j_p^{i_k})$ in the order in which those states appear in T. Note that, since p is not necessarily found in every G_i , the superscripts i_{ℓ} are not necessarily consecutive. We will say that a p-trace τ_p is time-like if $j_p^{i_{\ell+1}} \ge j_p^{i_{\ell}}$ for all ℓ .

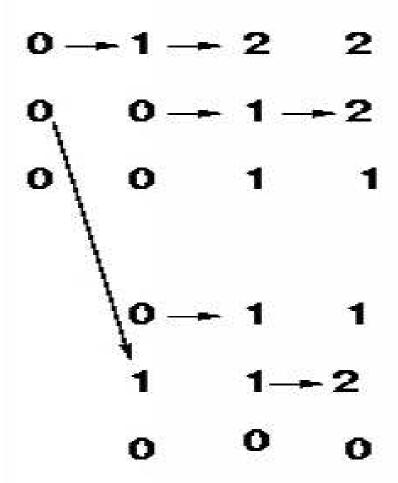
Definition 2.4. An SPMD parallel execution E is a directed graph of transitions starting at $E_{G_{tot}} = S_{G_{tot},0}$ all of whose p-traces (def: 2.3) are time-like.

Motivation:

- 1. Experimentally explore non-determinism in parallel state and execution
- 2. Verify that synchronization is detectable in execution, find other features that consistently appear in visualization
- 3. Characterize hardware-independent costs of synchronization (and other constructs) by analysis of entropy



non-deterministic transition



Parallel state is list of concurrent blocks -> multiple possible successor states

SPMD model used for simpler representation, but not restricted to this model.

Represent code as Control Flow Graph CFG

Process q state is basic block x and memory contents M at step j

$$\sigma_{q,j} = (x_j, M_j)$$

Execution is a series of states representing a walk through CFG

$$S_{\Gamma,J} = \{\sigma_{i,j}\}$$

A parallel state is the set of sequential states ocurring concurrently

Gamma is the ordered set of processes and J the ordered set of step numbers - a parallel transition occurs when 1 or more processes advance - even if processes are deterministic, parallel transition is not Representing each state as a P-tuple of concurrent block numbers gives us a P dimensional hypercube with side=number of blocks.

Some features in the phase space repeat (synchronization code in specific blocks). Most states are not deterministic and won't repeat.

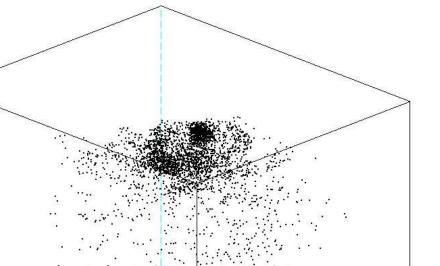
Given a set of points corressponding to one or a set of executions, Gibbs-Shannon entropy is:

$$-\sum p_i \log_2(p_i)$$

We use standard combinatoric properties to caculate the probabilities:

$$p_i = \frac{P!}{N^P \prod (b_k)!}$$

Where the $\mathbf{b}_{\mathbf{k}}$ are subsets in the state with the same block number and we normalize to the total phase space hypervolume



MPI Synchronous, 17 processes

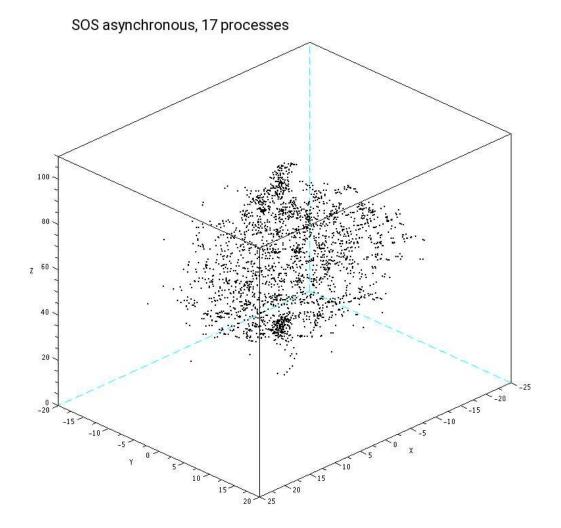
110

90 -

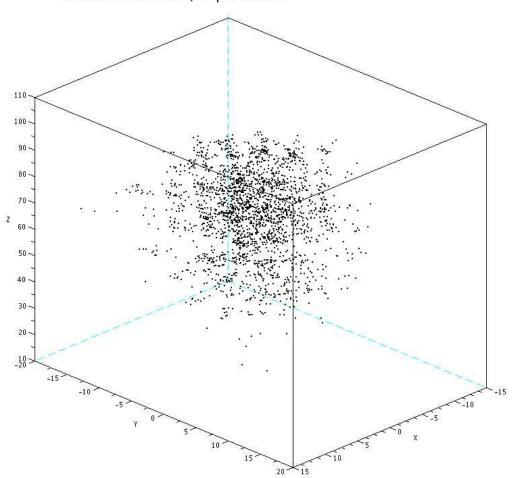
70 -

40 ×

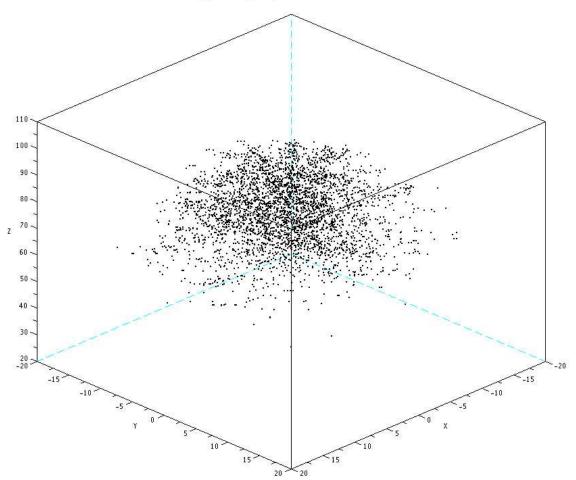
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No communication , 17 processes



No communications, guided, 17 processes

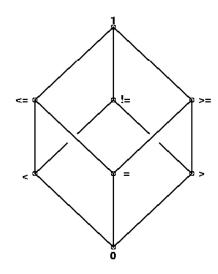


Algebra

 $S=\{1,\geq,\leq,=,>,<,0\}$ is complete

Lattice closed under intersection, union and complement, with 0 and 1 is a Boolean Algebra

S with set operations is **B**³



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phase space reduction:

">" synchronization removes 1 quadrant - R=.75 of phase space per dimension

"=" synchronization removes 2 quadrants - R=.5 of phase space per dimension

given unsynchronized entropy S, then entropy with synchronization S' is given by

S' = \log(\exp(S) \times R^N)) (given interval between synch longer than relaxation time)

Time prediction - dt/t = \log(N) \times dS/S (the \log N factor assumes collective communication tree)
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AMOEBA - time: _____ TARDIS: comparison of amoebampi and amoebaM (no synch) 17 amoebasos.mat entropy = 12.06215727 mean state size 5.340149838 time 2550742 17 amoebampi.mat entropy = 11.40358976 mean state size 5.660535992 time 1842581 17 amoebaN.mat entropy = 12.53261805 mean state size 5.178458642 time 991982 17 amoebaM.mat entropy = 14.42974945 mean state size 4.930923558 time 603365 dt/T = 0.763455104436278 (expected) dt/T = 0.46486557378577 (actual RAVEN: comparison of amoebampi and amoebaM (no synch) 17 amoebasos.mat entropy = 11.79212385 mean state size 5.21425634 time 13013195 17 amoebampi.mat entropy = 10.85077704 mean state size 5.616563899 time 6372684 17 amoebaN.mat entropy = 12.78105428 mean state size 4.997991238 time 3577237 17 amoebaM.mat entropy = 16.32720129 mean state size 4.387070064 time 981748 dt/T = 0.924557497217248 (expected) dt/T = 0.78695924005089AMOEBA - entropy prediction RAVEN: 11.4094047683197 - expected 11.79 TARDIS: 9.40940476831973 - expected 12.06215727

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