Parallel Game Tree Search

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Abstract

- Use multiprocessor shared-memory or distributed memory machines to search the game tree in parallel.
- Questions:
 - Is it possible to search multiple branches of the game tree at the same time while also gets benefits from the searching window introduced in alpha-beta search?
 - What can be done to parallelize Monte-Carlo based game tree search?
- Tradeoff between overheads and benefits.
 - Communication
 - Computation
 - Synchronization
- Techniques
 - For alpha-beta based search algorithms.
 - Lockless transposition table.
 - For Monte-Carlo based search algorithms.
- Can achieve reasonable speed-up using a moderate number of processors on a shared-memory multiprocessor machine.

Comments on parallelization

- Parallelization can add more computation power, but synchronization introduces overhead and may be difficult to implement.
- Synchronization methods
 - Message passing, such as MPI
 - Shared memory cells
 - ▶ Avoid a record becoming inconsistent because one is reading the first item, but the last item is being written.
 - ▶ Memory locked before using.
 - It may be efficient to broadcast a message.
- Locking the whole transposition table is definitely too costly.
 - The ability to lock each record.
 - Lockless transposition table technique.
- A global transposition table v.s. distributed transposition tables.

Speed-up (1/2)

- Speed-up: the amount of performance improvement gotten in comparison to the the amount of hardware you used.
 - Assume the amount of resources, e.g., time, consumed is T_n when you use n when you use n processors.
 - Speed-up = $\frac{T_1}{T_n}$ using n processors.
- Speed-up is a function of n and can be expressed as sp(n).
 - Scalability: whether you can obtain "reasonable" performance gain when n gets larger.
- Choose the "resources" where comparisons are made.
 - The elapsed time.
 - The total number of nodes visited.
 - The scores.
 - • •
- Choose the game trees where experiments are performed.
 - Artificial constructed trees with a pre-specified average branching factor and depth.
 - Real game trees.

Speed-up (2/2)

- Three different setups for experiments.
 - Use the a sequential algorithm P_{seq} for the baseline of comparison.
 - Use the best sequential algorithm P_{best} for the baseline of comparison.
 - Use a 1-processor version of your parallel program $P_{1,par}$ as the baseline of comparison.
 - \triangleright It is usually the case that $P_{1,par}$ is much slower than P_{best} .
 - \triangleright It is often the case that $P_{1,par}$ is slower than P_{seq} .
 - Use an optimized sequential version of your parallel program $P_{1,opt}$ as the baseline of comparison.
 - \triangleright It is also usually the case that $P_{1,opt}$ is slower than P_{best} .
- Choose the game trees where experiments are performed.
 - Artificial constructed trees with a pre-specified average branching factor and depth.
 - Real game trees.

Amdahl's law

- The best you can do about parallelization [G. Amdahl 1967].
- \blacksquare Assume a program needs to execute T instructions and and x of them can be parallelized.
 - Assume you have n processors and an instruction takes a unit of time.
 - Parallel processing time is

$$\geq T - x + \frac{x}{n} + O_n \geq T - x.$$

where O_n is the overhead cost in doing parallelization with n processors.

Speed-up is

$$\leq \frac{T}{T-x}.$$

- If 20% of the code cannot be parallelized, then your parallel program can be at most 5 times faster no matter how many processors you have.
- Depending on \mathcal{O}_n , it may not be wise to use too many processors.

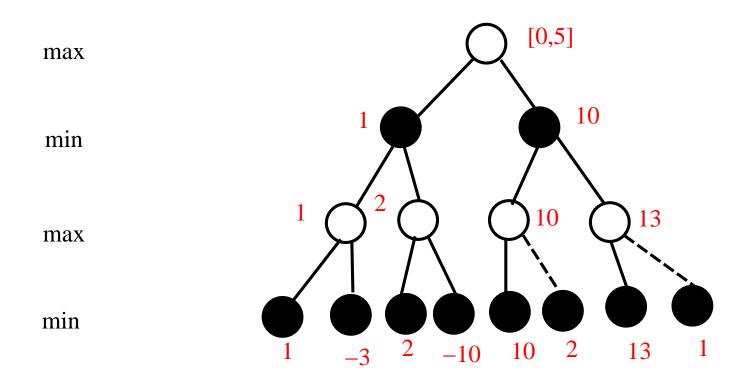
Load balancing and speed-up factor

Load balancing

- ▶ The ratio between the amount of the largest work on a PE and the amount of the lightest work on another PE.
- ▶ Good load balancing is a key to have a good speed-up factor.
- Speed-up factor: ratio between the parallel version with a given number of processors and the baseline version.
- Is it possible to achieve super linear speed-up?
 - \bullet Super linear speed-up means you can make the code to run N times faster using less than N times about of hardware.
 - > Yes, on badly ordered game trees.
 - ▶ Not in real game trees with a reasonable good algorithm.

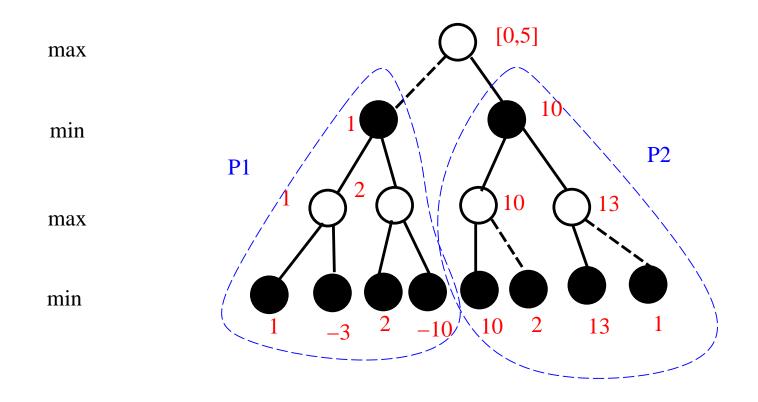
Super-linear speed-up (1/3)

- Sequential alpha-beta search with a pre-assigned window [0,5]:
 - Visited 13 nodes.



Super-linear speed-up (2/3)

- Parallel alpha-beta search with a pre-assigned window [0,5] on two processors:
 - P2: visited 5 nodes, and then the root performs a beta cut.
 - P1: being terminated by the root after 5 nodes are visited.



Super-linear speed-up (3/3)

- Total sequential time: visited 13 nodes.
- Total parallel time for 2 processors: visited 6 nodes.
- We have achieved a super-linear speed-up.

Comments on super-linear speed-up (1/2)

- Parallelization can achieve super-linear speed-up only if the solution is not found by enumerating all possibilities.
 - For example: finding an entry of 1 in an array.
- If the solution is found by exhaustively examining all possibilities, then there is no chance of getting a super-linear speed-up.
 - For example: counting the total number of 1's in an array.
- Overhead in parallelization comes from how much work should each processor "talks" to each other in order to decide the solution.
 - Trivially parallelizable: almost no need to talk to each other.

Comments on super-linear speed-up (2/2)

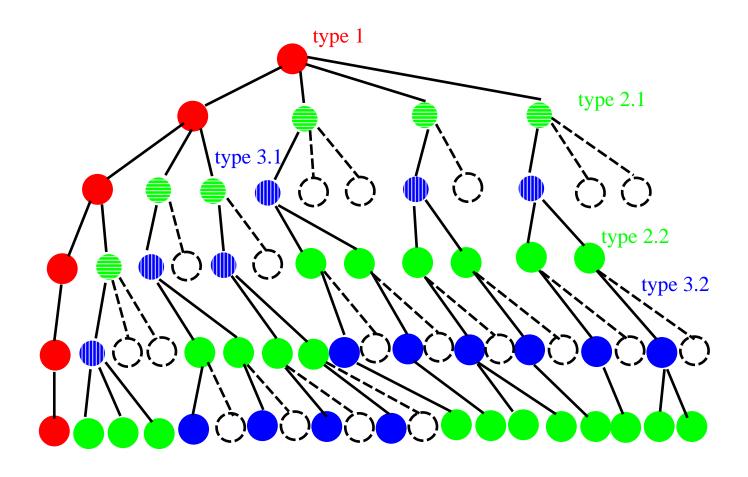
- Why is it possible to obtain a super-linear speed-up in searching a game tree using alpha-beta based algorithm?
 - Assume some cut-off happens during the execution.
 - Parallel algorithms offer a chance of getting a different "move ordering".
 - It is possible to find a solution faster.
- It is also possible to get poor speed-up if the "move ordering" of the parallel version is bad.
 - You may perform unnecessary work, e.g., searching a branch that will be cut in the future.
- For Monte-Carlo based search algorithm, super-linear speed-up may be obtained by trying out different PV branches at the same time.
 - Increase the chance of finding the right branch.

Parallel α - β search

- Three major approaches: depend on what tasks can be parallelized and the model of parallelism.
 - Principle variation splitting (PV split)
 - ▶ Central control or global synchronization model of parallelism.
 - Young Brothers Wait Concept (YBWC)
 - ▶ Client-server model of parallelism.
 - Dynamic Tree Splitting (DTS)
 - ▶ Peer-to-peer model of parallelism.

Classification of nodes (1/2)

 Classify nodes in a game tree according to [Knuth & Moore 1975].



Classification of nodes (2/2)

■ Type 1 (PV): principle variation.

- > Nodes in the leftmost branch.
- ▶ PV nodes needs to be searched first to established a good search bound.
- ▶ After the first child is searched, the rest of its children can be searched in parallel.

■ Type 2 (CUT): cut nodes.

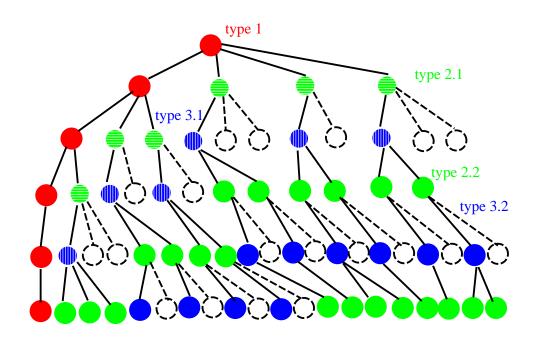
- ▶ Children of type-1 and type-3 nodes.
- ▶ Because children of a cut node may be cut, it is not wise to perform searches in parallel for children of a cut node.

Type 3 (ALL): all nodes.

- ▶ The first branch of a cut node.
- ▶ All children of an all node need to be explored.
- ▶ It is better to search these children in parallel.

Principle variation splitting

- Algorithm PVS:
 - Execute the first branch to get a PV branch $n_1, n_2, n_3, \ldots, n_d$ where n_d is a leaf node.
 - for i=d-1 down to 1 do
 - \triangleright Update the bound information using information backed-up from n_{i+1}
 - \triangleright for each non-PV branch of n_i do in parallel
 - ▶ A processor gets a branch and searches
 - ▶ Update the bounds when a branch is done



Comments for PV splitting

Comments:

- Parallelism is done on type-2 branches of a type-1 node.
- May not be able to use a large number of processors efficiently.
- Load balancing is not good.
 - ▶ The ratio between the amount of the largest work on a PE and the amount of the lightest work on another PE.
- Synchronization overhead is large.
- When the first branch is usually not the best branch, then the overhead is huge.
- Achieve a speed-up of 4.1 for 8 processors and 4.6 for 16 processors [Manohararjah '01].
 - ▶ Poor scalability.
 - ▶ Limited speed-up: within 5.

Improvements:

- ▶ When a processor is idle, it helps out a busy processor by sharing its tasks.
- ▶ Observe some improvements, but not much.

Young brothers wait concept (1/2)

- Concept: at each node, when the first branch is explored and a bound is obtained, then all the other branches can be executed in parallel.
 - Split point: a node whose value of the first branch is known.
 - Highest split point of a tree: a split point whose depth is the least.
 - A processor is assigned and owns a subtree rooted at a node.
 - ▶ This processor is the server of this subtree.
 - An idle processor asks a server for a subtree to search.
 - ▶ This processor is a **client** of this server.

Young brothers wait concept (2/2)

■ Algorithm *YBWC*:

- Let P_1 own the root of the game tree and begin to search using alpha-beta pruning until the tree is completely searched.
 - During searching, maintain the split point information.
- While the game tree is not searched completely, do In parallel for each processor P_i do
 - \triangleright If P_i is idle, it looks for server processors with split points.
 - \triangleright P_i gets a branch from a highest split point and owns this subtree.
 - \triangleright P_i begins to search using alpha-beta pruning and maintain the split point information.
 - \triangleright When a subtree owned by P_i has been searched, returns the information to the server processor where it gets the job from.
 - \triangleright P_i is idle again.

Comments for YBWC

Comments:

- Can utilize many processors.
- Parallelism is done on almost all nodes.
- It is possible to use non-shared-memory architectures.
 - ▶ For example: distributed memory machines.
 - ▶ Speed-up: 137 using 256 processors [Manohararjah '01].
 - ▶ Scalability is moderate.
 - ▶ Load balancing is not always good.
- The cost of splitting a node needs to be calculated to avoid splitting small trees.

Dynamic tree splitting (DTS)

Concepts:

- Peer-to-peer approach so that no one owns any subtree.
- The processor who finished last on a split point reports the value to the parent of the split point.
- More criteria for the selection of split points.

DTS: Classification of nodes

- D-PV: a node that has the same alpha and beta values as the root.
- D-CUT: a minimizing node with the same beta as the root or a maximizing node with the same alpha as the root.
 - On a MAX node,
 - ▶ if some branches are searched, then the returned values from the branches may update the lower bound.
 - ▶ If the lower bound is highered (updated), then it is possible to visit less nodes.
 - ▶ Hence it may not be cost effective to parallelize.
 - ▶ Note: It takes time to initialize a new job.
 - On a MIN node,
 - ▶ if some branches are searched, then the returned values from the branches may update the upper bound.
 - ▶ If the upper bound is lowered (updated), then it is possible to visit less nodes.
 - ▶ Hence it may not be cost effective to parallelize.
 - ▶ Note: It takes time to initialize a new job.
- D-ALL: any node that is neither D-PV nor D-CUT.
 - Nothing much is known here.

Split point: confidence

- A confidence factor is associated with each D-CUT and D-ALL node.
 - That is, the chance of a node being the specified type.
- If many moves (up to a limit, say 3) have been searched at a D-CUT node, then the confidence that it is a D-CUT node decreases.
- If several moves have been searched at a D-ALL node, then the confidence that it is a D-ALL node increases.

DTS: Split point

Criteria for a split point:

- The node must be of type D-PV, D-ALL with a high confidence or or D-CUT with a low confidence.
- If it is a D-PV node, its first branch must have been searched.
- Set thresholds for confidence factors.
 - ▶ A D-ALL node with a high confidence factor remains to be a candidate for a split point.
 - ▶ Can also fork a D-ALL node with the highest confidence factor first.
 - ▶ A D-CUT node with a low confidence factor may be a split point.

Note:

- ▶ Nodes that are higher up in the tree (closer to the root) represent more work.
- ▶ You want to fork a branch that are higher up and with a larger confidence factor for D-ALL, or with a smaller confidence factor for D-CUT.
- ▶ Use the above information to compute a global priority.

DTS: Algorithm

Algorithm DTS:

- Initialize a global job list with the root as the only available job.
- while the job list is not empty do
 - ▶ Idle processors look for jobs with the highest priority in the global job list.
 - ▶ A working processor maintains its own split point information at the global job list.
 - ▶ A working processor updates bounds when a job is finished and then becomes idle.

Comments:

- Used by several state-of-the-art chess programs.
- Spend a bit more time to decide whether a node is a split point or not.
 - ▶ Takes some time to tune for the best parameters.
- Speed-up factor is very good: 3.7 for 4 processors, 6.6 for 8 processors and 11.1 for 16 processors [Manohararjah '01].
- Load balancing is good.
- Scalability is reasonable.

Comments: parallel α - β search

- DTS is currently being used by most Chess-like programs.
- It also takes time to tune the system parameters for DTS to work well.
 - The threshold for confidence factors.
 - Dynamically adjusting of the confidence factors.

Memory issues (1/2)

- During searching, each process needs to maintain the following information.
 - Local data: such as the current depth, current best move.
 - Data that can be used later: such as the hash information.
- Distributed memory model.
 - Maintain each own data in a private memory area.
 - Exchange information when needed.
 - ▶ Using message passing to probe a hash entry.
 - ▶ Using message passing to return the value of a probe.
- Shared memory model.
 - Maintain each local data in a private memory area.
 - Maintain the re-used information in a global area.
 - ▶ Concurrent read is often allowed in the model.
 - ▶ Concurrent write is usually not allowed in the model.
 - ▶ Lock the cell when it needs to be written.

Memory issues (2/2)

- Advantage and disadvantage
 - Distributed memory model.
 - ▶ Coding is easy.
 - ▶ Slow response time.
 - Shared memory model.
 - ▶ Overhead in locking.
 - ▶ Fast response time when there is no extensive memory contention.
- Often used techniques: Lockless transposition tables.
 - Allow concurrent read, but not concurrent write.
 - Assume writing of an entry is atomic.
 - ▶ Records in data structures used consist of multiple entries.

Lockless transposition table

Scenario

- ullet Assume each entry of the transposition table H contains two parts where reading/writing each part is atomic.
 - \triangleright Position_signature: $H_1[]$ array, say each entry is 64-bit long.
 - \triangleright Data: $H_2[]$ array, say each entry is 64-bit long.
- Assume the hash key $hash_key$ is the rightmost h, say h=32, bits of $Position_signature$.
- ullet To read or write an hash entry given a position P, you do the followings.
 - Compute $Position_signature(P)$ and Data(P).
 - Let $hash_key(P)$ be the rightmost h bits of $Position_signature(P)$.
 - Read or write $H_1[hash_key(P)]$.
 - Read or write $H_2[hash_key(P)]$.
- Problem: The hash entry is corrupted if
 - ullet P is being visited at the same time by two processes C_1 and C_2 so that
 - $ightharpoonup C_1$ writes $H_1[hash_key(P)]$.
 - $ightharpoonup C_2$ writes $H_2[hash_key(P)]$.

Solution

- Algorithm for writing an entry
 - Compute $Position_signature(P)$ and Data(P).
 - Let $hash_key(P)$ be the rightmost h bits of $Position_signature(P)$.
 - write: $H_1[hash_key(P)] \leftarrow Position_signature(P)$ **XOR** Data(P).
 - write: $H_2[hash_key(P)] \leftarrow Data(P)$.
- Algorithm for reading an entry
 - Compute $Position_signature(P)$.
 - Let $hash_key(P)$ be the rightmost h bits of $Position_signature(P)$.
 - read: $W_1 \leftarrow H_1[hash_key(P)]$
 - read: $W_2 \leftarrow H_2[hash_key(P)]$
 - reconstruct: $W_1 \leftarrow W_1 \text{ XOR } W_2$
 - verify: check whether $W_1 = Position_signature(P)$
 - ▶ if they equal, then use this entry.
 - ▶ if they do not equal, then the entry is corrupted.

Why this works

- $H_1[hash_key(P)] = Position_signature(P)$ **XOR** Data(P).
- $\blacksquare H_2[hash_key(P)] = Data(P).$
- $H_1[hash_key(P)]$ **XOR** $H_2[hash_key(P)] = Position_signature(P)$.
- If $H_1[i]$ and $H_2[i]$ are written by two different processes with $Data(P_1)$ and $Data(P_2)$, then it will probably not produce the right position signature.
- Comments:
 - May have errors because of hash collisions.
 - It is not too difficult to extend this method to an hash table with more than 2 entries.

Parallel Monte-Carlo tree search

- Leaf parallelization.
- Root parallelization.
- Tree parallelization.
 - With global synchronization.
 - With local synchronization.

MCTS with UCT

- Algorithm MCTS:
- 1: Obtain an initial game tree
- 2: Repeat the following sequence N_{total} times
 - 2.1: Selection
 - ▶ From the root, pick a PV path to a leaf such that each node has the best UCB "score" among its siblings
 - ▶ May decide to "trust" the score of a node if it is visited more than a threshold number of times.
 - ▶ May decide to "prune" a node if its score is too bad now to save time.

2.2: Expansion

- ▶ From a best leaf, expand it by one level.
- ▶ Use some node expansion policy to expand.

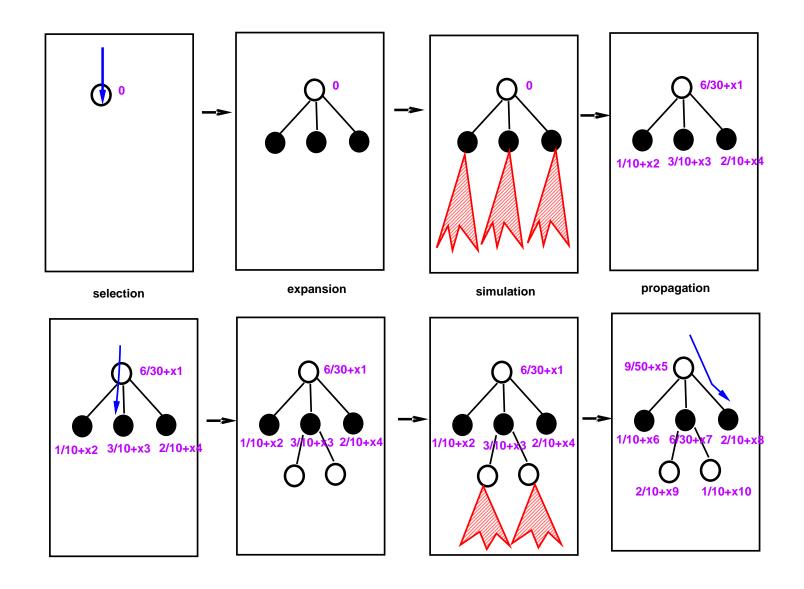
• 2.3: Simulation

- ▶ For the expanded leaves, perform some trials (playouts).
- ▶ May decide to add knowledge into the trials.

2.4: Back propagation

- ▶ Update the "scores" for nodes using a good back propagation policy.
- Pick a child of the root with the best score as your move.

MCTS: example



Leaf parallelization

• Algorithm PMCTS_{leaf}:

- loop:
- Select the best leaf and the PV path.
- Perform Expansion in sequential.
- Perform Simulation, i.e., multiple trials, in parallel on the same leaf.
- Perform Back propagation in sequential.
- if the terminating condition is not matched, go to loop;

Comments:

- Coding is very easy.
- Good parallelization for performing a large number of trials.
 - ▶ Speedup is good up to using a reasonable number of PE's.
- Can utilize a large number of PE's.
 - ▶ Scalability is reasonable.
- Problem: the best leaf may no longer be the best after only a few more trials.

Root parallelization

- Algorithm PMCTS_{root}:
 - loop:
 - Duplicate k copies of the current game tree.
 - for each copy do in parallel
 - ▶ Selection: pick the *i*th largest PV in the *i*th copy
 - ▶ Expansion: ...
 - ▶ Simulation: ...
 - Combine the copies into one copy by merging statistics on nodes and put the information into the current game tree.
 - Back propagation: ...
 - if the terminating condition is not matched, go to loop;

$PMCTS_{root}$: comments

Comments:

- Coding is easy.
- Can utilize as many PE's as available.
 - ▶ Scalability is good.
- Problem: the best leaf may no longer be the best after only
- May need to make sure that each tree does not pick the same best leaf.
- Problem: Need to have a mechanism to properly choose the best leaves among all trees.
 - ▶ Avoid duplicated efforts.

Tree parallelization — global synchronization

• Algorithm PMCTS $_{Tg}$:

- Use only one game tree.
- Perform Selection, Expansion and Simulation in parallel.
 - ▶ Different threads may work on different nodes in parallel.
 - ▶ Need a mechanism to ensure threads are not working on the same leaf.
- Use a global lock to make sure the game tree is writable by one thread during the Back propagation phase.

Comments:

Speed-up is bad.

Tree parallelization — local synchronization

• Algorithm PMCTS $_{Tl}$:

- Make every node of the game tree as a global variable.
- Perform Selection, Expansion, Simulation and Back propagation in parallel.
 - ▶ Different threads may work on different nodes in parallel.
 - ▶ Need a mechanism to ensure threads are not working on the same leaf.
- Use a lock to make sure each node is writable by one thread during Back propagation.

Comments:

- Heavy O.S. overhead.
- Unsure about the scalability.

Problems of parallel Monte-Carlo search

- Each iteration of a Monte-Carlo simulation is a Markov chain process.
 - You need to know the result of the previous trial to decide the current selection.
 - Making trials in parallel has a larger statistical error.
 - May explore the wrong branch if synchronization is done only after a lot of trials.
 - May not have too much parallelism if synchronization is done after only a few trials.
- The cost of synchronization.
 - Shared global variable.
 - Cost of lock and unlock.
 - Memory bandwidth.
 - Network bandwidth.
- The cost of coding.

Parallel Monte-Carlo search: Analysis

- ullet Amdahl's law: assume a program needs to execute T instructions and and x of them can be parallelized. can be parallelized.
 - ullet Assume you have n processors and an instruction take a unit of time.
 - Parallel processing time $\geq T x + x/n + p_n \geq T x$ where p_n is the cost for overhead in doing parallelization with n processors.
 - Speed-up $\leq T/(T-x)$.
 - ▶ If 20% of the code cannot be parallelized, then your parallel program can be at most 5 times faster no matter how many processors you have.
- Leaf and root parallelization both have a large portion that is not parallelizable.
- Global or local synchronization has a large overhead.
- Comments
 - Need a better parallel implementation.
 - Need a better way to deal with the increasing error in doing more samplings.

Concluding remarks

- Need to think about tradeoff between costs in doing parallelism and benefits of saving in searching efforts because of parallelism.
- May need to think how to maintain distributed transposition tables.
- May need to think about the machine architecture.
 - Shared-memory vs. distributed memory.
 - Fine grain or coarse grain.
 - Whether the parallel version is stable or not?
 - ▶ Ease of debugging.
 - ▶ Ease of coding.

References and further readings (1/2)

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