

Parallel Game Tree Search

Tsan-sheng Hsu

徐讚昇

tshsu@iis.sinica.edu.tw

<http://www.iis.sinica.edu.tw/~tshsu>

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
- 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 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.
 - ▷ *It is usually the case that $P_{1,par}$ is much slower than P_{best} .*
 - ▷ *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.
 - ▷ *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].
- Assume a program needs to execute T instructions 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 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.*

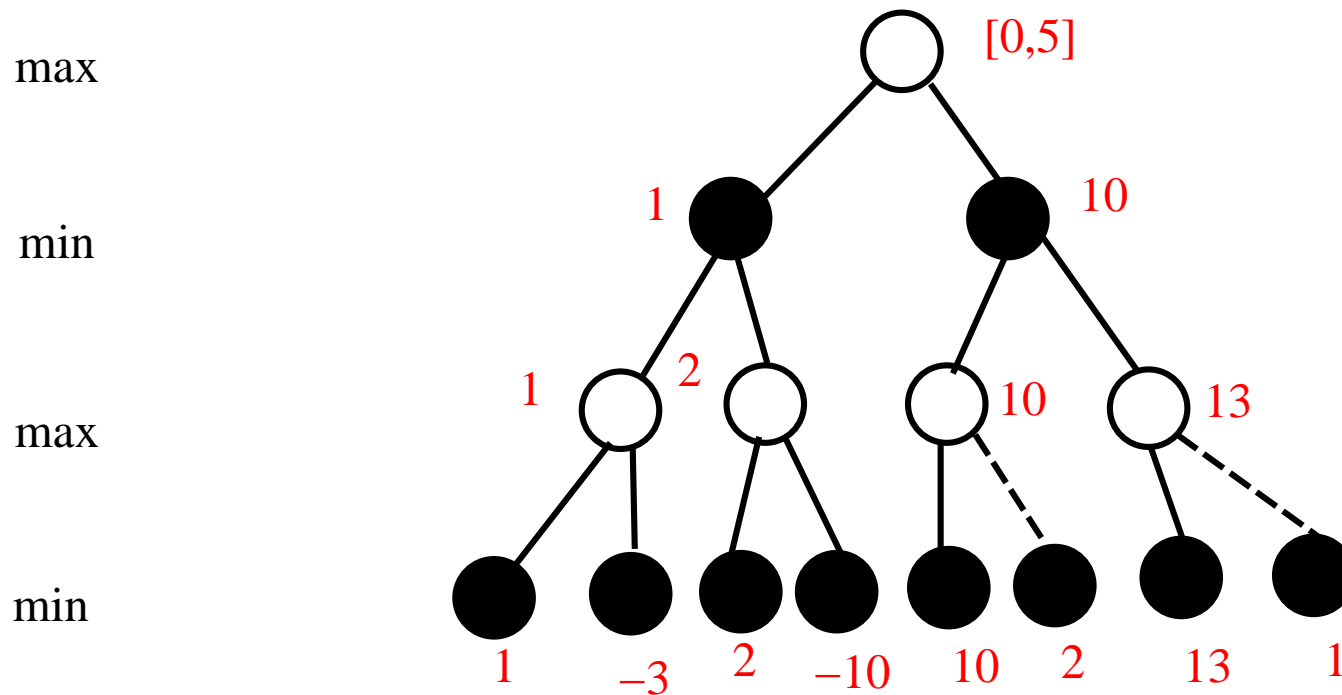
■ Speed-up factor: ratio between the parallel version with a given number of processors and the baseline version.

■ Is it possible to achieve **super linear** speed-up?

- 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.*

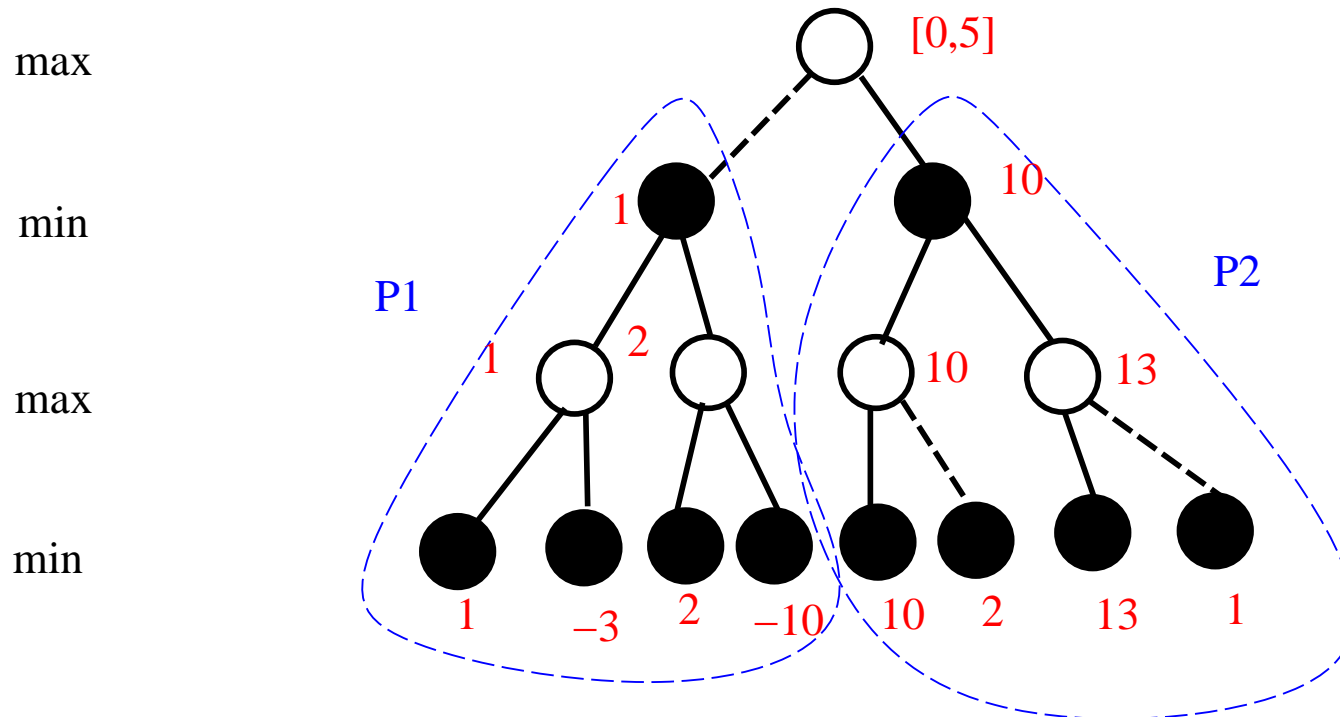
Super-linear speed-up (1/3)

- 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 in parallel.
- If the solution is found by exhaustively examining all possibilities, then there is no chance of getting a super-linear speed-up.
 - For example: the problem of 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)

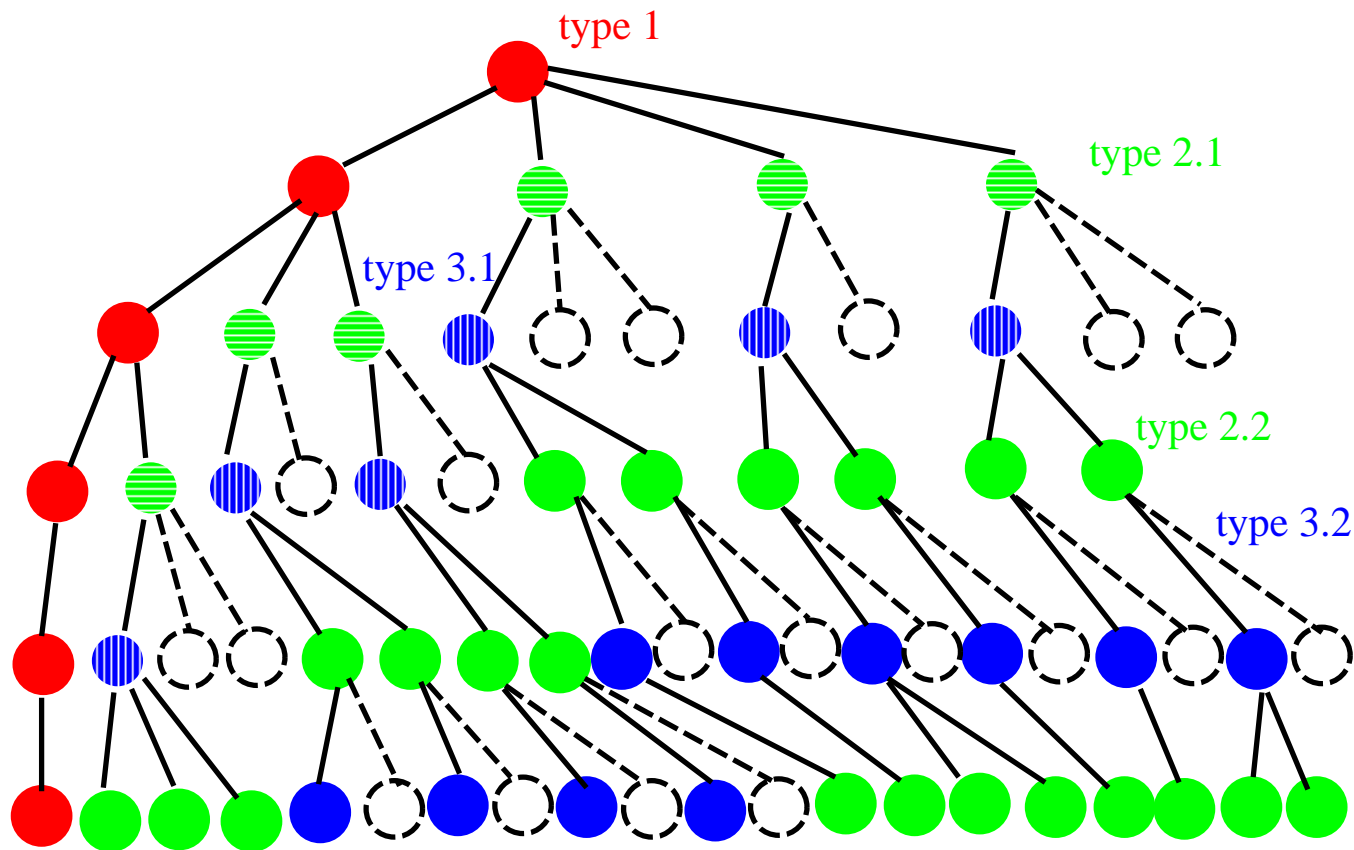
- Why is it possible to obtain a super-linear speed-up in searching a game tree using alpha-beta based algorithm?
 - Assume some cut-off happens during the execution.
 - Parallel algorithms offer a chance of getting a different “move ordering”.
 - It is possible to find a solution faster.
- It is also 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 maybe obtain 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].



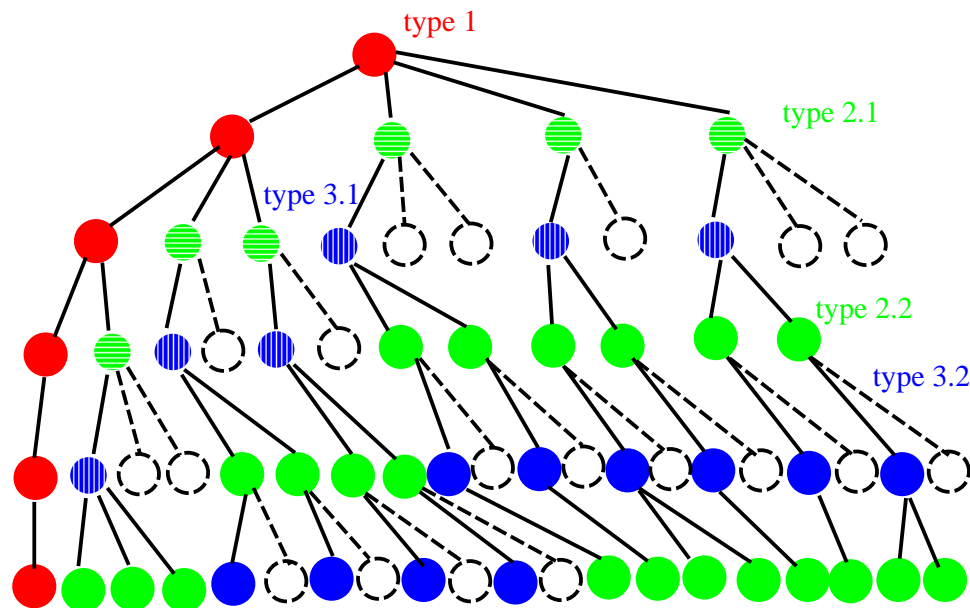
Classification of nodes (2/2)

- **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 $n_1, n_2, n_3, \dots, n_d$ where n_d is a leaf node.
- for $i = d - 1$ down to 1 do
 - ▷ Update the bound information using information backed-up from n_{i+1}
 - ▷ 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.
 - ▷ *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
 - ▷ *If P_i is idle, it looks for server processors with split points.*
 - ▷ *P_i gets a branch from a highest split point and owns this subtree.*
 - ▷ *P_i begins to search using alpha-beta pruning and maintain the split point information.*
 - ▷ *When a subtree owned by P_i has been searched, returns the information to the server processor where it gets the job from.*
 - ▷ *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.*
 - ▷ *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 selections 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.**
 - Means the chance of being a node of the specified type.
- **If many moves (up to a limit of 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.
- 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 find 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.
 - ▷ *Current read is often allowed in the model.*
 - ▷ *Lock the cell when it needs to write.*

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.

- Do not assume writing of an entry is atomic.

Lockless transposition table

■ Scenario

- Assume each entry of the transposition table H contains two parts where reading/writing each part is atomic.
 - ▷ *Position_signature*: 64 bits $\rightarrow H_1$.
 - ▷ *Data*: 64 bits $\rightarrow H_2$.
- Assume the hash key *hash_key* is the rightmost h , say $h = 32$, bits of *Position_signature*.

■ 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 $H_1(\text{hash_key}(P))$.
- Read or write $H_2(\text{hash_key}(P))$.

■ Problem: The hash entry is **corrupted** if

- P is being visited at the same time by two processes C_1 and C_2 so that
 - ▷ C_1 writes $H_1(\text{hash_key}(P))$.
 - ▷ C_2 writes $H_2(\text{hash_key}(P))$.

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: $H_1(hash_key(P)) \leftarrow Position_signature(P) \text{ XOR } Data(P)$.
- 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

- $H_1(\text{hash_key}(P)) = \text{Position_signature}(P) \text{ XOR } \text{Data}(P)$.
- $H_2(\text{hash_key}(P)) = \text{Data}(P)$.
- $H_1(\text{hash_key}(P)) \text{ XOR } H_2(\text{hash_key}(P)) = \text{Position_signature}(P)$.
- **If $H_1(i)$ and $H_2(i)$ are written by two different processes with $\text{Data}(P_1)$ and $\text{Data}(P_2)$, 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.

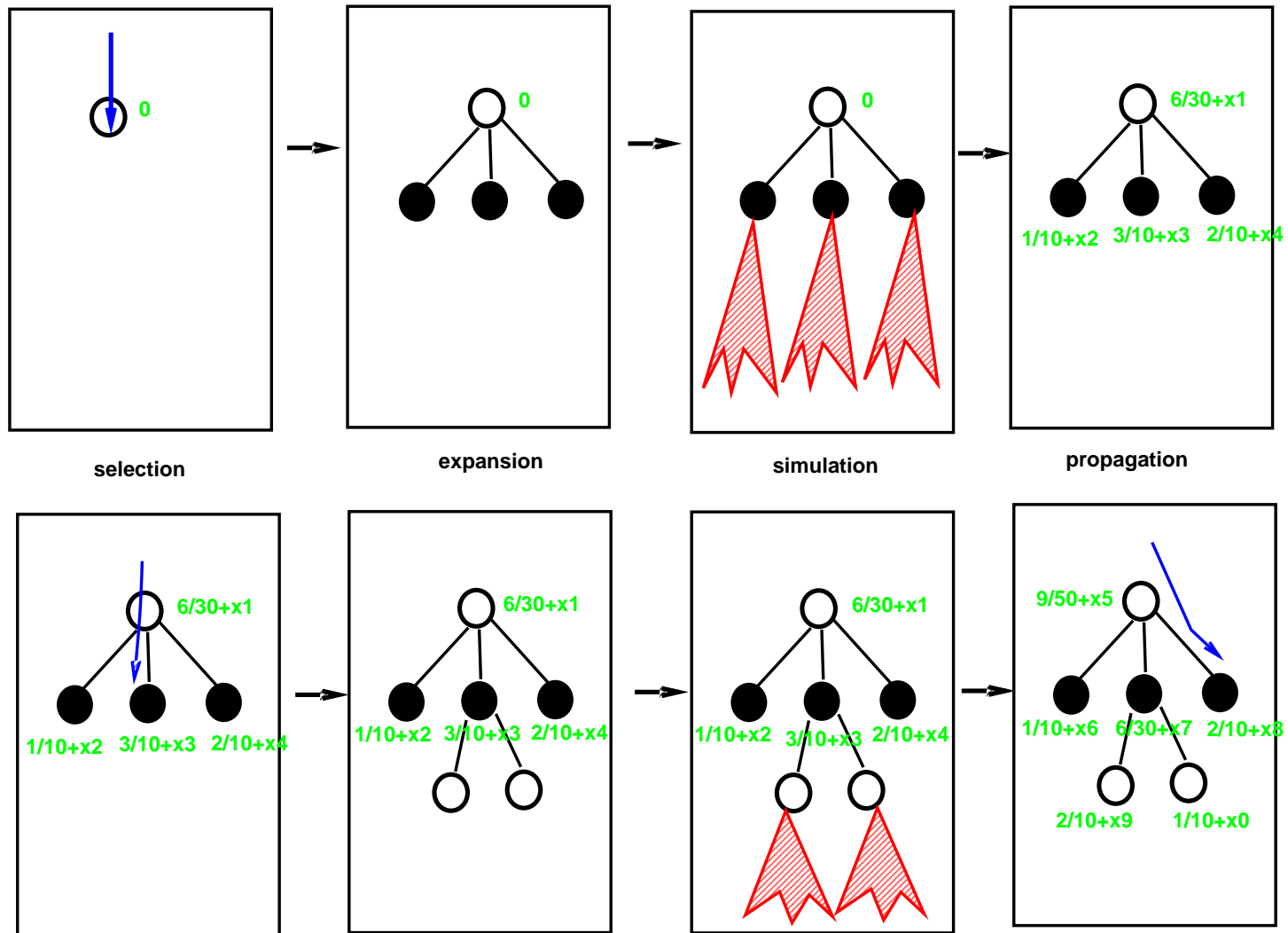
Parallel Monte-Carlo tree search

- Leaf parallelization.
- Root parallelization.
- Tree parallelization with global synchronization.
- Tree parallelization 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 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} :

- Select the best leaf.
- Perform Expansion in sequential.
- Perform Simulation, i.e., multiple trials, in parallel on the same leaf.
- Perform Back propagation in sequential.

■ Comments:

- Coding is very easy.
- Good parallelization for performing a large number of trials.
- Can utilize a large number of PE's.
- The best leaf may no longer be the best after only a few more trials.

Root parallelization

■ Algorithm PMCTS_{root} :

- Duplicate k copies of the current game tree.
- Perform Monte-Carlo tree simulation on each copy in parallel for a few trials.
- Combine the copies into one copy by merging statistics on nodes and put the information into the current game tree.

■ Comments:

- Coding is easy.
- Can utilize as many PE's as available.
- May need to make sure that each tree does not pick the same best leaf.
- 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 programming.**

Parallel Monte-Carlo search: Analysis

- **Amdahl's law:** assume a program needs to execute T instructions and x of them can be parallelized.
 - 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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