Green-Marl

A DSL for Easy and Efficient Graph Analysis

Motivation

Issues with large-scale graph analysis

- Performance
- Implementation
- Capacity

Performance Issues

- RAM latency dominates running time for large graphs
- Solved by exploiting data parallelism

Implementation Issues

Writing concurrent code is hard

- Race-conditions
- Deadlock
- Efficiency requires deep hardware knowledge
- Couples code to architecture

Alternative: a DSL

Green-Marl and its compiler

- High level graph analysis language
- Hides underlying complexity
- Exposes algorithmic concurrency
- Exploits high level domain information for optimisations

Example

```
Procedure Compute BC (
   G: Graph, BC: Node Prop<Float>(G)) {
G.BC = 0; // initialize BC
Foreach(s: G.Nodes) {
  Node Prop<Float>(G) Sigma;
  Node Prop<Float>(G) Delta;
   s.Sigma = 1; // Initialize Sigma for root
   InBFS(v: G.Nodes From s) (v!=s) {
    v.Sigma = Sum(w: v.UpNbrs) {w.Sigma};
   InRBFS (v!=s) {
    v.Delta = Sum(w:v.DownNbrs) {
      v.Sigma / w.Sigma * (1+ w.Delta)
     };
    v.BC += v.Delta @s; //accumulate BC
```

$$g(v) = \sum_{s \neq v \neq t} \frac{\sigma_{st}(v)}{\sigma_{st}}$$





- Based on processing graph properties
- Mappings from a node/edge to a value
- e.g. the average number of phone calls between two people

Green-Marl is designed to compute

- scalar values from a graph and its properties
- new properties for nodes/edges
- selecting subgraphs (instance of above)

- Support for parallelism (fork-join style)
- Implicit
 - \circ G.BC = 0;
- Explicit
 - o Foreach(s: G.Nodes) (s!=t)
- Nested

Other characteristics

- Relaxed memory model (but atomic)
- Reductions
- Built-in graph and collection types
- Built-in operations: BFS, DFS, etc.



- Currently compiles to C++
- Semantic analysis checks for conflicts in parallel sections of code
- Generic and graph-specific optimisations
 9 in total

Architecture Independent Optimisations e.g. *Flipping Edges*

Foreach(t:G.Nodes)(f(t))

Foreach(s:t.InNbrs)(g(s))

t.A += s.B;

becomes

Foreach(s:G.Nodes)(g(s))

Foreach(t:s.OutNbrs)(f(t))

t.A += s.B;

Architecture Dependent Optimisations e.g. Saving BFS Children

InBFS(v:G.Nodes; s) { ... //forward }

InRBFS { // reverse-order traverse

Foreach(t: v.DownNbrs) { DO_THING(t); } }

becomes

```
_prepare_edge_marker(); // O(E) array for (e = edges ..) {
 for (e = edges ...) {
   index_t t = ...node(e);
   if (isNextLevel(t)) { edge_marker[e] = 1; } DO THING(t); } }}
```

- 1 machine, 5 algorithms, 2 graph generators
- 32 million nodes, 256 million edges
- Compared with the SNAP graph analysis platform (only 3 algorithms)



BC scaling across cores



Conductance

In a nutshell...

- At least as fast as SNAP
- Good speedup of up to ~16 threads
- Algorithms that are hard to parallelise do not scale so well (Amdahl's Law)

Usability

- Between 50% and 10% the lines of code of other implementations
- Does not require application rewriting
- Embedded foreign code
- Concise and intuitive descriptions of graph algorithms (in their opinion!)