FPGA-Based Supercomputing: An Implementation for Molecular Dynamics

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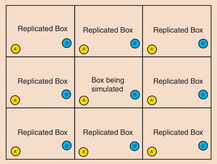
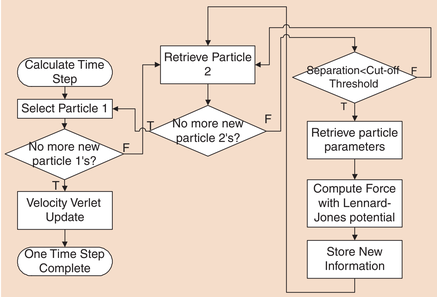
Molecular Dynamics

- Simulates atom and molecule interactions with Newtonian mechanics
- Applications - defect analysis, protein folding
- Any interesting volume has far too many particles to simulate
- Use periodic boundary conditions to reduce number of particles that must be tracked



- a cube of particles is simulated
- cube is replicated in every dimension
- a particle on the edge of the cube interacts with its *neighbour* on the other side of the cube

Simulation Process



- Model intermolecular forces using Lennard-Jones potential

$$F = -\nabla \phi_{r,LJ}(r) \qquad \phi_{LJ}(r) = 4\varepsilon\left[\left(\frac{\sigma}{r}\right)^{12} - \left(\frac{\sigma}{r}\right)^{6}\right]$$

- Forces must be calculated for each pairwise interaction between particles
  - problem becomes $O(n^2)$
  - for simplicity insignificant interactions can be discarded
- Acceleration due to net force is calculated using Newton's second law F=ma
- Must perform time integration to get position and velocity updates.
- Use Velocity Verlet Update rule to perform this Integration

$$v\left(t+\frac{\delta t}{2}\right) = v(t) + \frac{\delta t}{2}a(t) \qquad v(t) = v\left(t-\frac{\delta t}{2}\right) + \frac{\delta t}{2}a(t)$$

$$r(t+\delta t) = r(t) + \delta t v(t) + \frac{\delta t^2}{2}a(t)$$

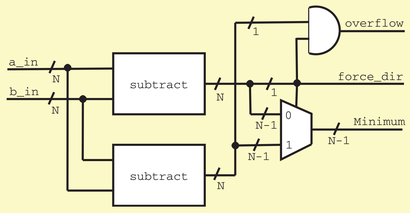
Previous Hardware

- no known previous implementations used programmable logic
- MODEL chip - floating point throughout
  - 76 MODEL chips are 50 times faster than a 200 MHz Sun Ultra 2
- MD-GRAPE - fixed and floating point design
  - Communication with host computer slows simulation speed

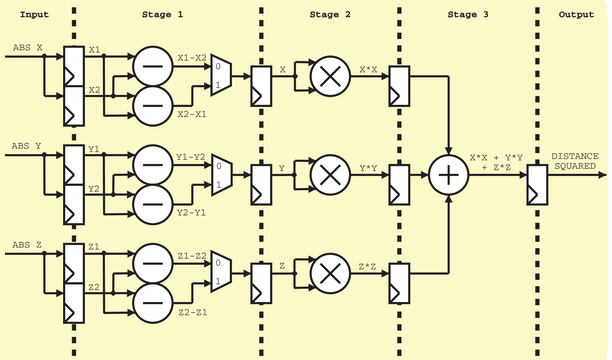
Pair Generator

- Retrieves 3-D co-ordinates for all pairs of particles from memory.
- Computes the distance squared between all pairs of particles and the force direction.
- Distance squared between two particles calculated using the formula:

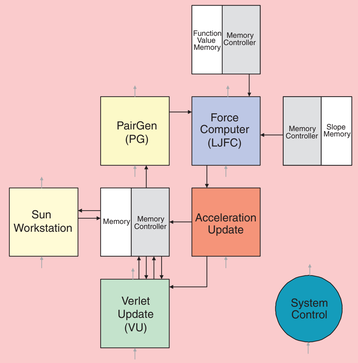
$$\min(x1-x2, x2-x1)^2 + \min(y1-y2, y2-y1)^2 + \min(z1-z2, z2-z1)^2$$



- By using unsigned numbers in the Minimum operation the Periodic Boundary Conditions are accounted for



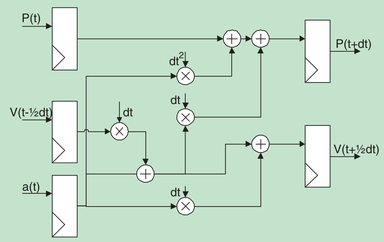
System Overview



For every timestep in the simulation

1. PG examines a pair of particles and determines the distance between the pair.
2. LJFC computes the force between the pair
3. AU computes the total acceleration on a particle from the series of incremental forces.
4. VU uses the total acceleration to update the position and velocity

Verlet Update



- Uses acceleration to perform time integration and update velocity and position
  - Performed once per particle
- Conventionally two separate steps are performed to obtain a half timestep velocity
- For hardware implementation this is inefficient Instead a single combined operation is performed
  - however, velocity stored in memory is a half timestep ahead of current time
- Multiplication is simplified by pre-multiplying acceleration in lookup table by the timestep

Software

- Constants Package
  - Generates shared constants for hardware and software
  - Includes precision, scaling factors, number of particles

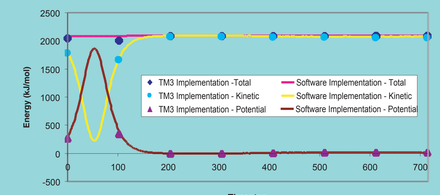
Scaling Factors and Precision of Various Fields

Field	Used in Block	Scaling Factor	Precision
r_x	PG	2^{-64}	38
r^2	PG, LJFC	2^{-128}	76
Slope	LJGC	2^{77}	36
Function	LJFC	2^{16}	47
Residual x slope	LJFC	2^{15}	48
Pseudo-acceleration	LJFC, AU	2^{-14}	50
Velocity	VU	2^{-13}	51
Velocity after step 2 of Verlet	VU	2^{-13}	51
Acceleration	VU	2^{-64}	37
Velocity x dt	VU	2^{-64}	37
Representation of t	VU	2^{-70}	22

- Top Level VHDL Generator
  - Uses the generated constants to create a wrapper

Validation

- Compared against academic C-based simulator:
  - MD3DLJ
  - Potential and Kinetic Energy track
  - Difference in energy for software and hardware was on the order of 1% RMS.

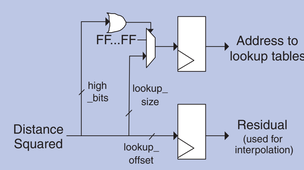


Comparison of Energy of TM3 vs Software Model

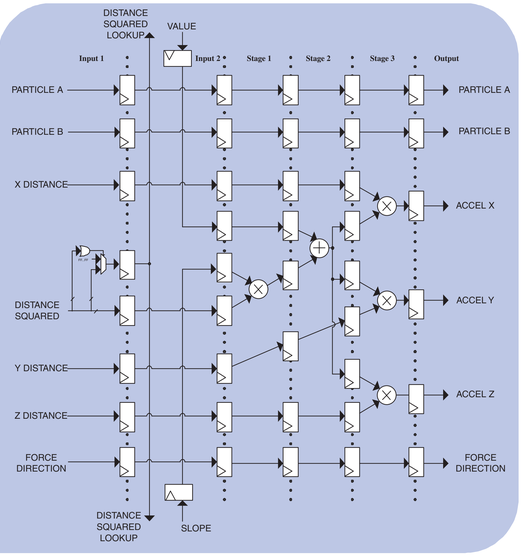
Time Step	Kinetic Energy (KJ/mol)	Potential Energy (KJ/mol)	Total Energy (KJ/mol)
Initial	1794	293.4	2087
Software – End	2078	14.4	2093
Hardware -End	2068	13.52	2081

Lennard-Jones Force Generator

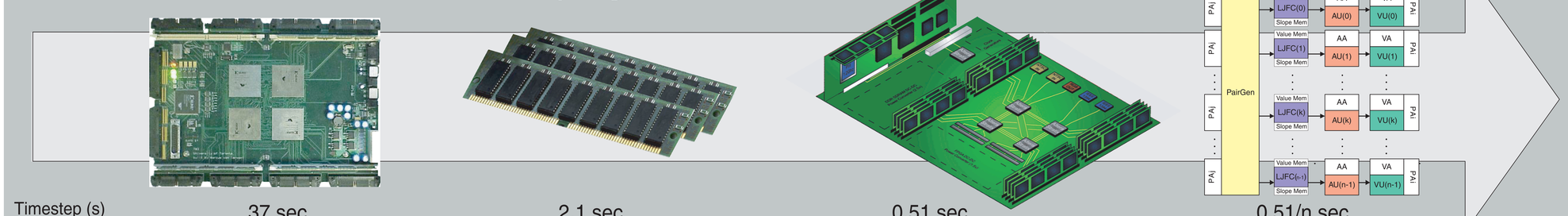
- Receives distance from PG
- Uses middle-order bits of r^2 to lookup value and slope of LJ potential function
- Since LJ potential does not change much at large distances, the higher-order bits are only used to limit the maximum distance
  - Allows for finer granularity of lookup where function is rapidly changing



- Uses lower bits of r^2 to multiply with slope to interpolate
  Adds function value to interpolated amount to obtain pseudo-acceleration
- Not true acceleration because it includes division by r and multiplication by dt, which were included in the look up table to simplify hardware
- Final multiplication by distance components to obtain acceleration components



Results



	TM3	Better Memory Organization	Modern FPGA	Multiple FPGAs
Timestep (s)	37 sec	2.1 sec	0.51 sec	0.51/n sec
Relative Speedup to Software (10.7s)	0.29X	5.1X	21X	21*nX

Comparison of MD Simulator and State-of-the-Art Processor

Per Board	2.4GHz P4 + 1GB DRAM	4 FPGAs + 24 MB SRAM
Performance	1X	Min 40X
Cost	~$1500	~$1500
Power	106W	40W

Metric	Improvement of MD System
Performance/Power	106X Improvement
Performance/Cost	40X Improvement
Performance/Space	40X Improvement

Supercomputers = many P4-like processors
Why not FPGA processors instead?

The TM3 project has been made possible by support from Micronet and Xilinx.