



Accelerating Scientific Model Optimization with a Pipelined FPGA-Based Differential Evolution Engine

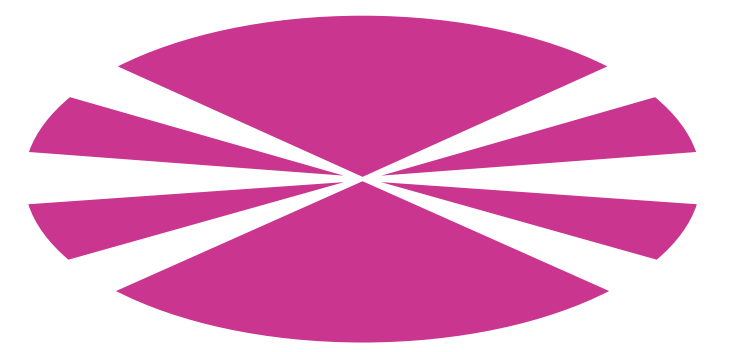
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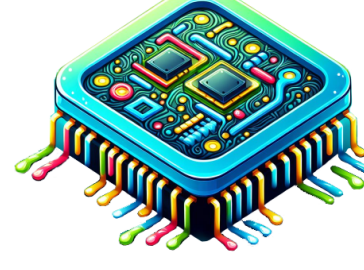
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INTRODUCTION

- Differential Evolution (DE) is a genetic algorithm that, when combined with Numerical Integration of scientific model on application-specific hardware:
 - Produces deep floating-point pipelines
 - Requires minimal external memory bandwidth
 - Benefits from large internal memory bandwidth
- Proposal: **Generic architecture for Differential Evolution with Numerical Integration methods**
 - Evaluated with 2 scientific models: Hodgking-Huxley (with non-adaptive 4th order Runge-Kutta) and Circadian (with adaptive Runge-Kutta-Fehlberg)

FPGA's strong points!



DIFFERENTIAL EVOLUTION

- Mutation:**
Alter each solution's parameters, using the others (with randomness)
$$m_{i,G+1} = s_{r1,G} + F \cdot (s_{r2,G} - s_{r3,G})$$
- Crossover:**
New candidate solution, hybrid from original solution and mutant
- Limiting:**
Ensure all parameter values are in valid range
- Selection:**
Compute cost function of new candidate; keep solution with lower cost

SCIENTIFIC MODELS USED FOR ASSESSMENT

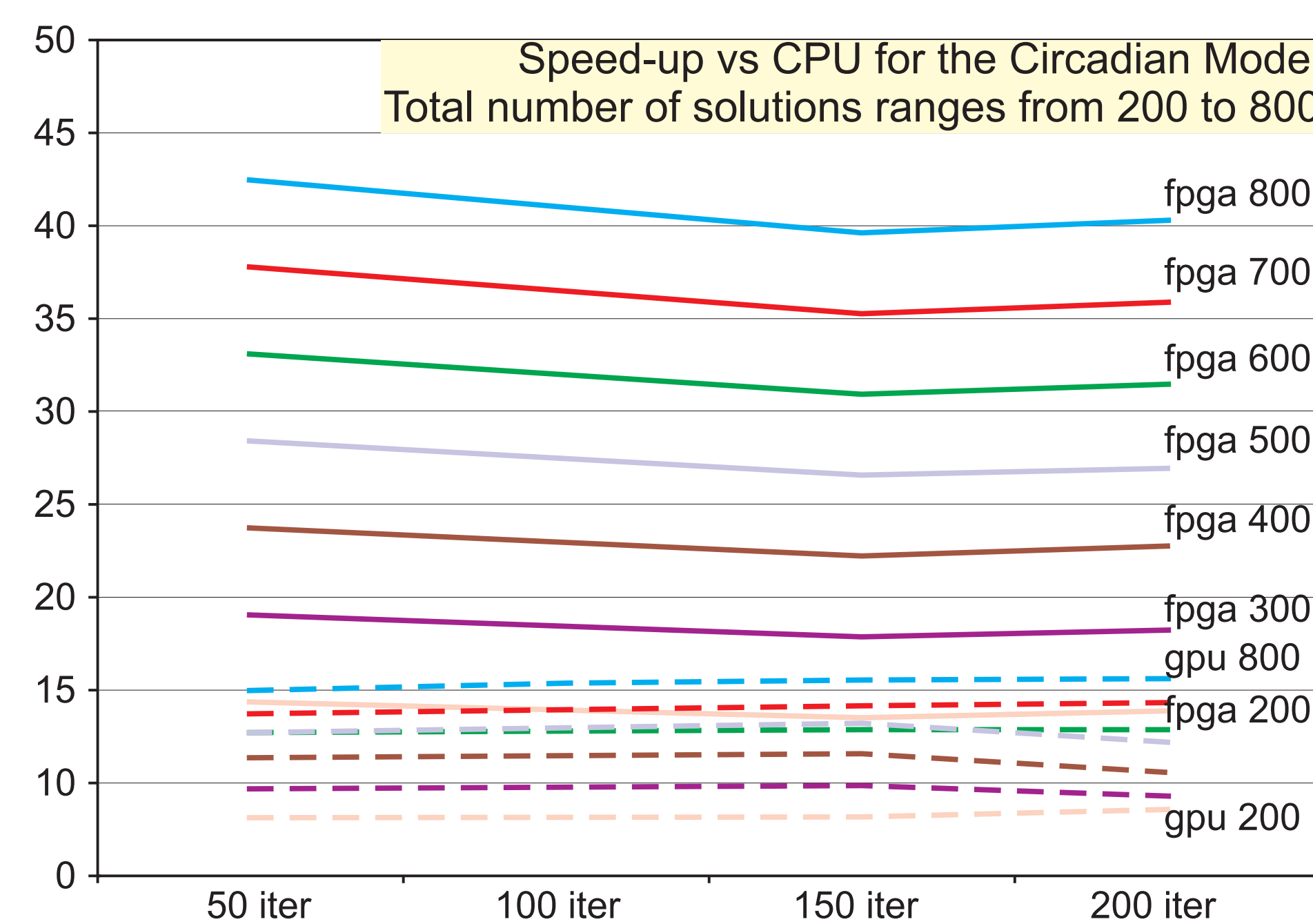
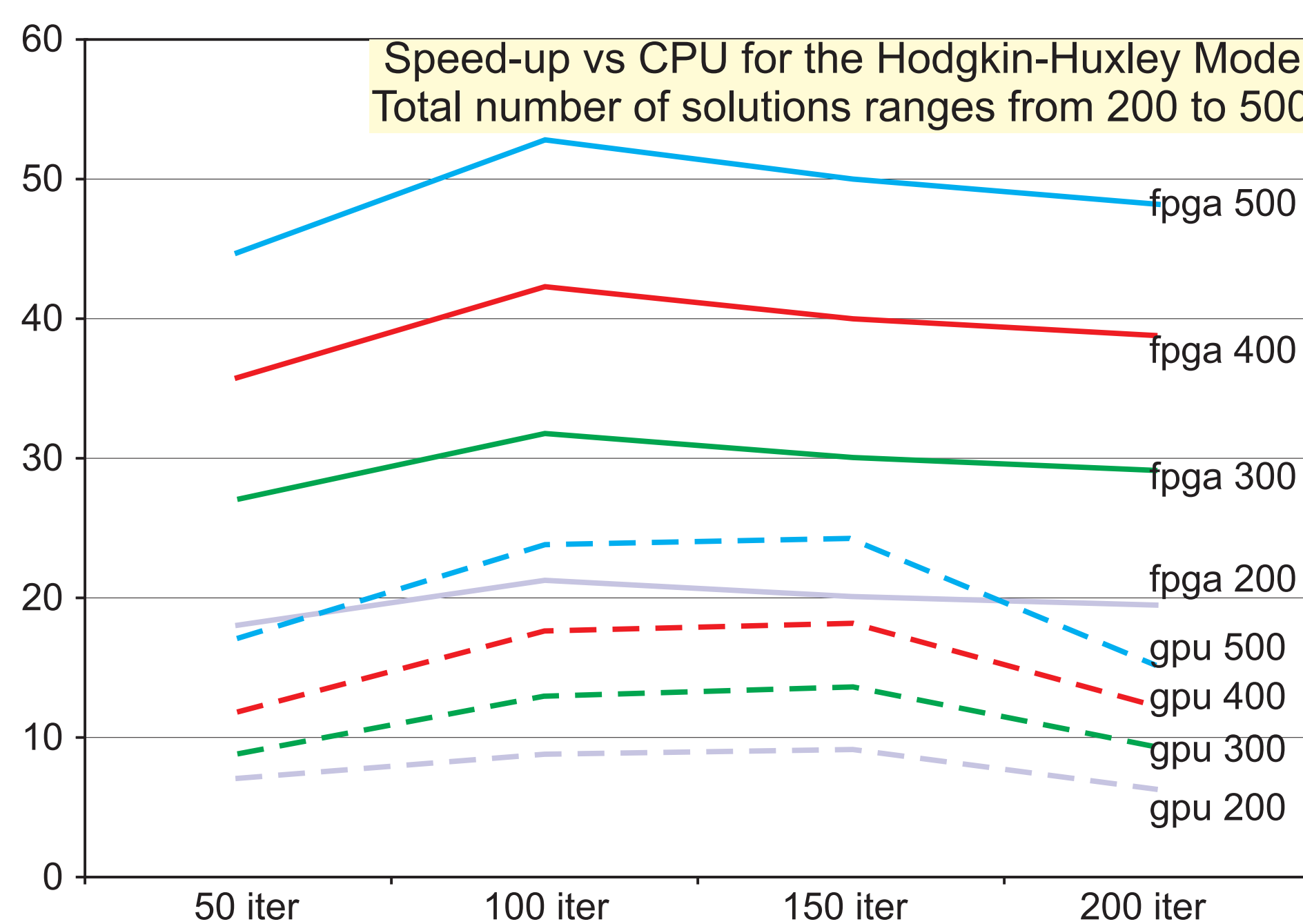
- Hodgking-Huxley:** action potentials in neuron axons
 - Multiple non-linear differential equations
 - Always converges with non-adaptive RK4
 - 11 unknown parameters per solution
 - Circadian:** biological clock of plants
 - Multiple coupled stiff differential equations
 - Requires adaptive numerical integration: RKF
 - 27 unknown parameters per solution
- × Complex, real-world models, not tackled previously in the literature in a fully-pipelined way

RESOURCE USAGE

- Synthesized for Virtex Ultrascale+ XCVU13P
- | | LUT | FF | BRAM | DSP |
|----------------|---------------|---------------|-----------|-------------|
| Hodgkin-Huxley | 220 123 (13%) | 230 438 (7%) | 114 (4%) | 2 272 (18%) |
| Circadian | 648 976 (38%) | 659 083 (12%) | 293 (11%) | 4250 (37%) |
- More than ×10 LUT, BRAM; ×100 FF, DSP than previous works

PERFORMANCE

- Tested: Multiple number of candidate solutions, multiple amount of numerical integration iterations
- Speedup with respect to single-threaded CPU implementation, executed in an AMD EPYC 7713 @ 3.67 GHz, compiled with *gcc* and *-O3* optimizations
- Comparison against **NVIDIA A100 GPU**, using a GPU-optimized high-performance implementation: One thread per candidate solution



- FPGA and GPU performance scale with the number of candidate solutions
 - FPGA scaling is more linear
- Average energy savings vs GPU:
 - Hodgkin-Huxley:**
 - ×9.0 more energy efficient (11.04% the energy consumption of the GPU)
 - Circadian:**
 - ×4.11 more energy efficient (24.33% the energy consumption of the GPU)

CONCLUSION AND FUTURE WORK

- Differential Evolution with Numerical Integration is well suited for FPGA-based CCMs: Parallelism, independent memory accesses, deep floating-point pipelines
- We present a generic, parameterized, modular architecture for Differential Evolution
- Our architecture presents superior performance to GPU-based high performance solutions
- Future work:** optimizing engine performance and resource usage, investigating new fitness functions, and comparing against multicore CPU implementations

Acknowledgements

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REFERENCES

