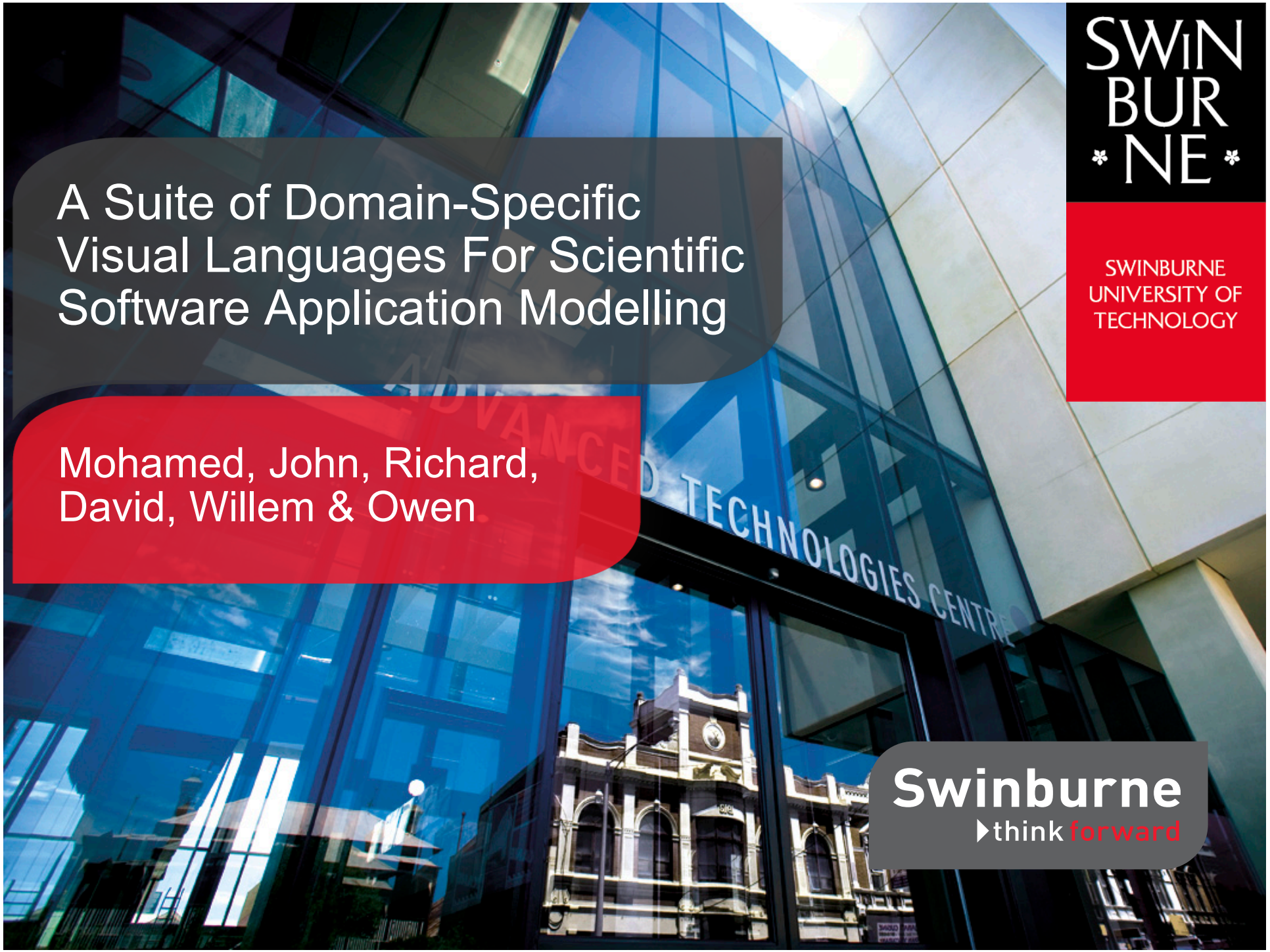

A Suite of Domain-Specific Visual Languages For Scientific Software Application Modelling

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Outline

- Problem / Motivation
- Approach
- Example Usage
- Future work

Problem

- Scientists are increasingly developing complex software for data analysis
- Most are not trained programmers
- Many are using complex software platforms and techniques e.g. distributed & parallel programming, GPUs, etc - that are hard even for experienced CS grads to use
- Approaches to address range from packaged software (Lack flexibility), DSLs (Also Flexibiity/Domain issues), programming patterns and toolkits (still complex)
- Still lack high-level, human-centric support; still really hard to develop high quality software for scientific apps

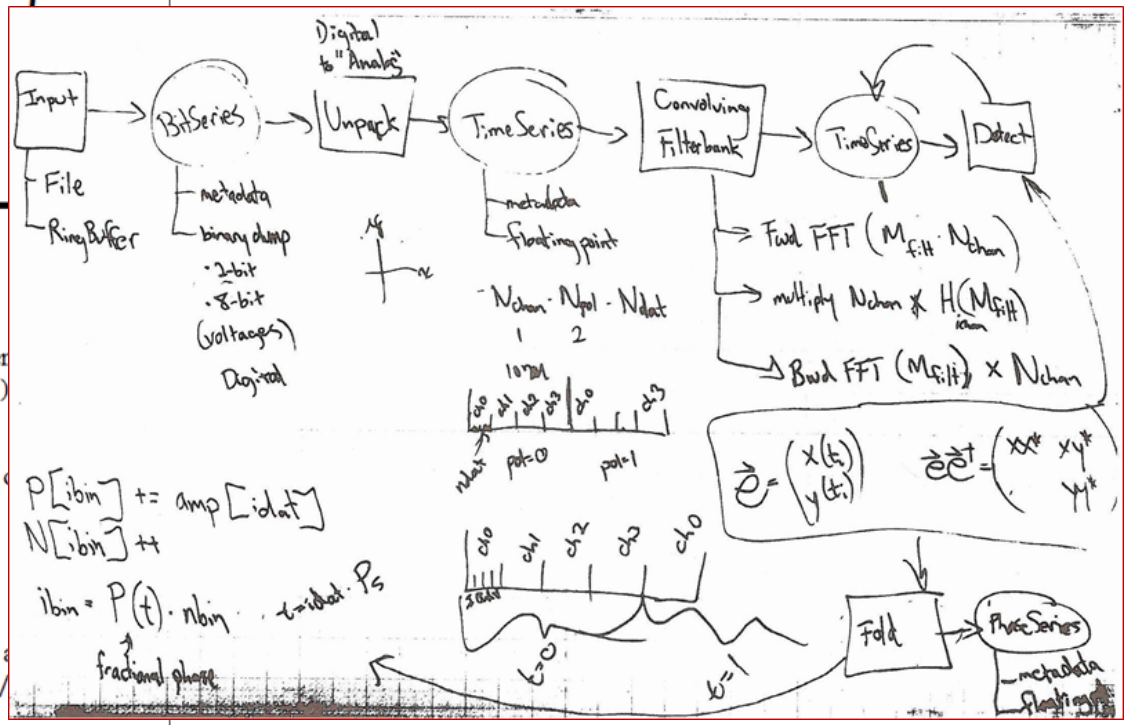
Motivation – how scientists design their applications…

$$\mathbf{F}_i = \sum_j^{N_m} \mathbf{F}_{ij}(\mathbf{r}_i, \mathbf{r}_j) - \nabla_i \left(-\frac{1}{2} \sum_j^{N_m} \mathbf{E}_j^0 \cdot \boldsymbol{\mu}_j^{\text{ind}} \right) + \sum_j^{N_m} \sum_k^{N_m} \mathbf{F}_{ijk}(\mathbf{r}_i, \mathbf{r}_j, \mathbf{r}_k),$$

Part 1 Initialization:
Part 1.1 Assign number of particles, energy, volume, etc.
Assign coordinates (r_i) at time (t) = 0.
Assign molecules an initial velocity (v_i).
Scale v_i consistent with the energy of the er
Assign any other $(\partial r^n/\partial t^n)_i$ values ($n > 2$)

Part 1.2 Initial force calculations:
Calculate forces (f_i) on each atom from all
Calculate acceleration (a_i) for each atom.

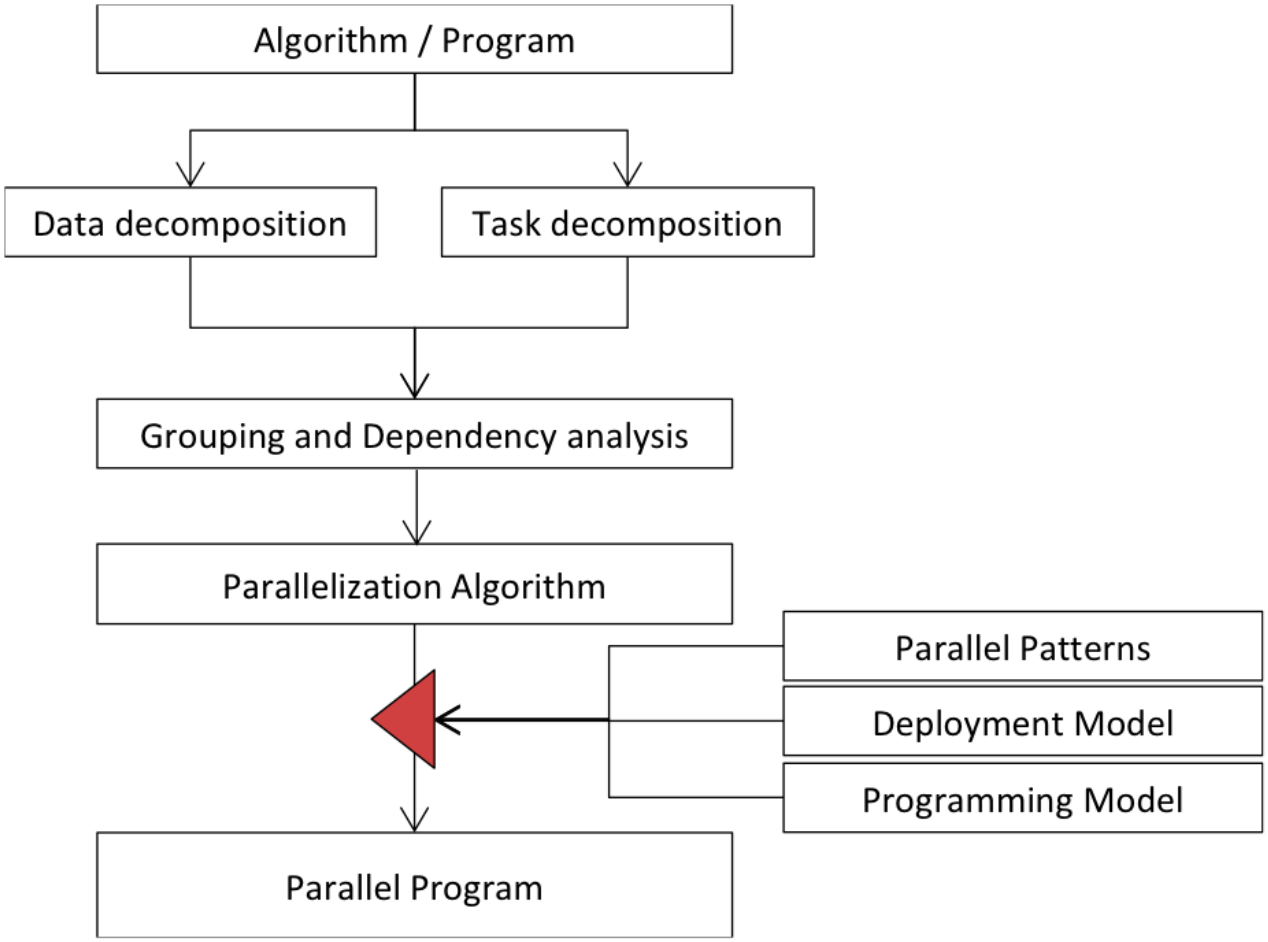
Part 2 Simulation Process:
loop
Part 2.1 Integrate equations of motion:
Part 2.1.1 Calculate f_i on each atom from all other j
Part 2.1.2 Apply integrator to update r_i, a_i, v_i, $(\partial r^n/$
Part 2.2 **if** ($t > tEquilibration$)
Accumulate averages.
else if (mod(m, *scalingInterval*) = 0)



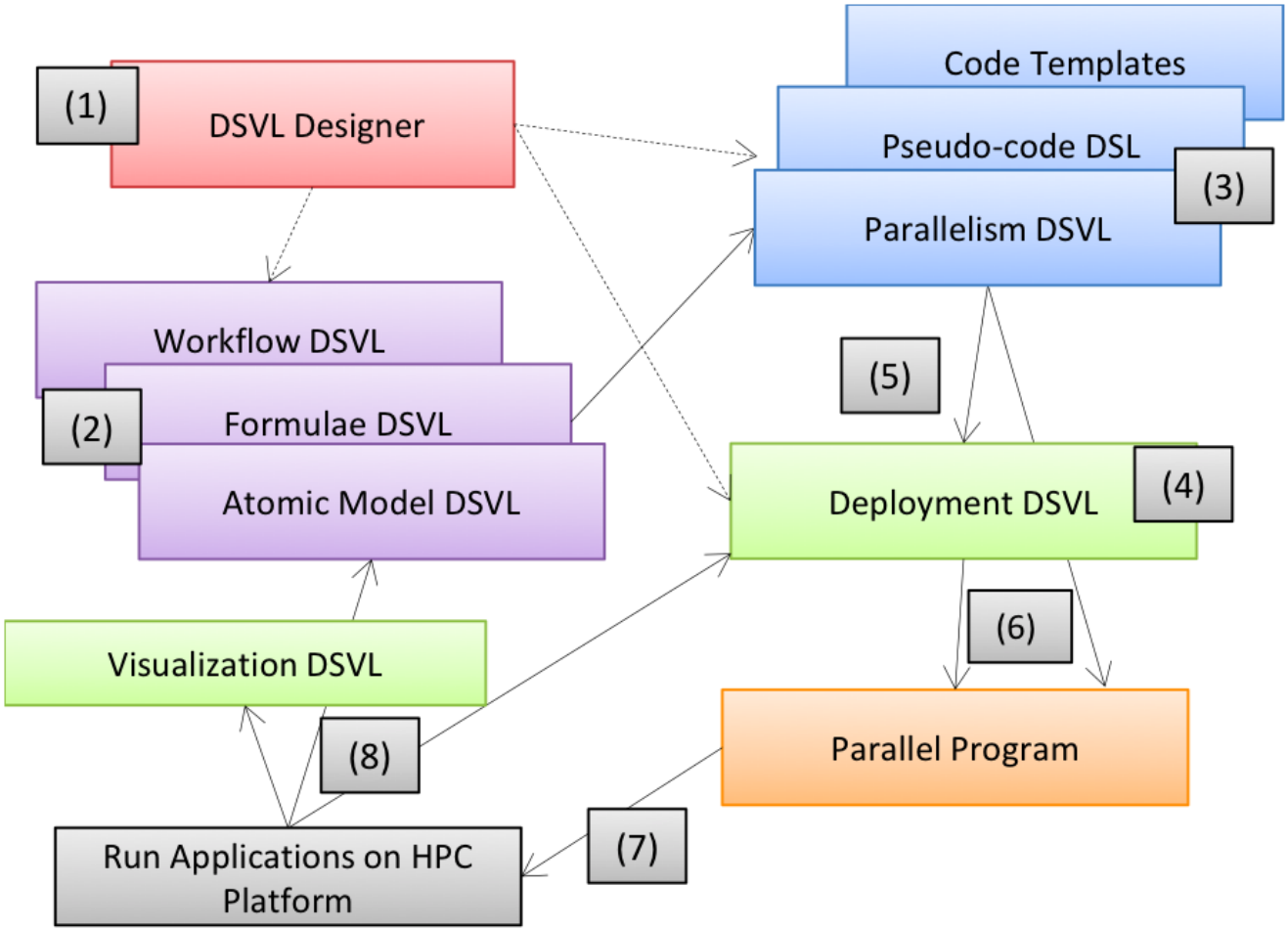
Approach

- Support scientists – and developers! – to model their applications at multiple levels of abstraction – domain right down to detailed C/C++/GPU kernel code
- Use set of user-defined and reusable DSVLs to model
- Provide web-based envrionment including DSVLs designer, coding, debugging, linking DSVL views
- Provide semi-automated support for generating lower-level models, generate code/code annotations, reverse-engineer higher-level models from (existing) code

Development approach to support

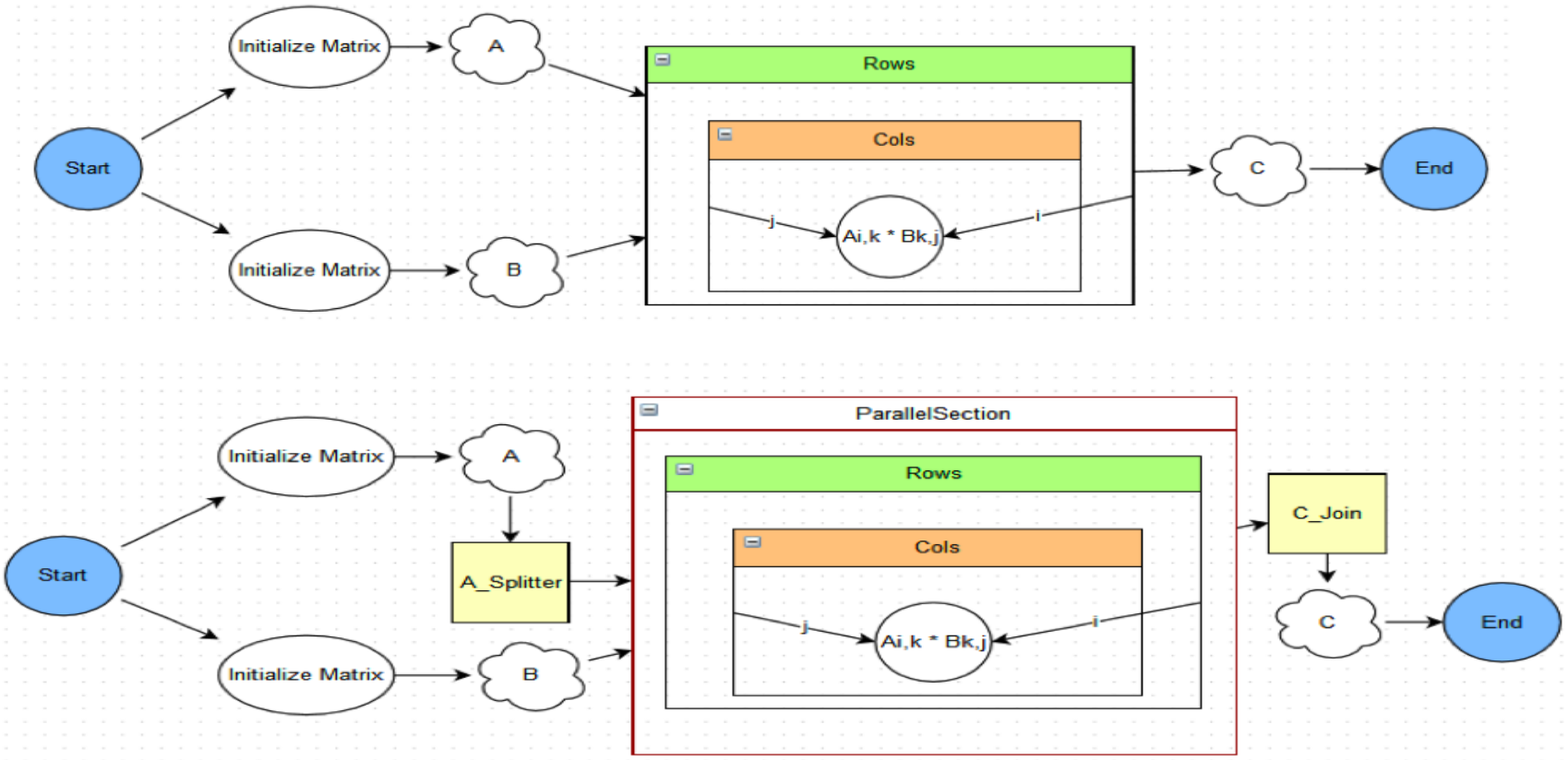


Our Approach

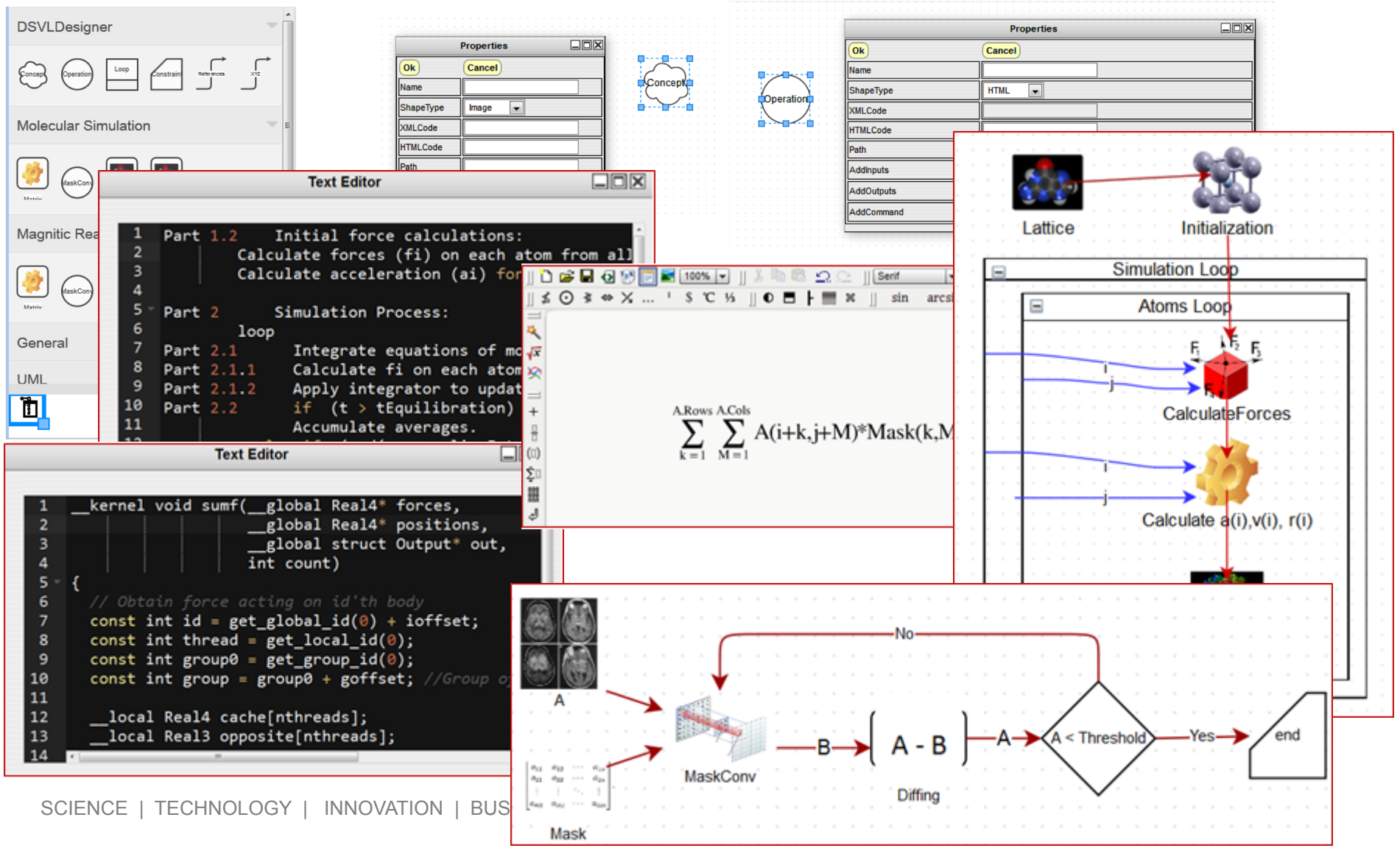


Example

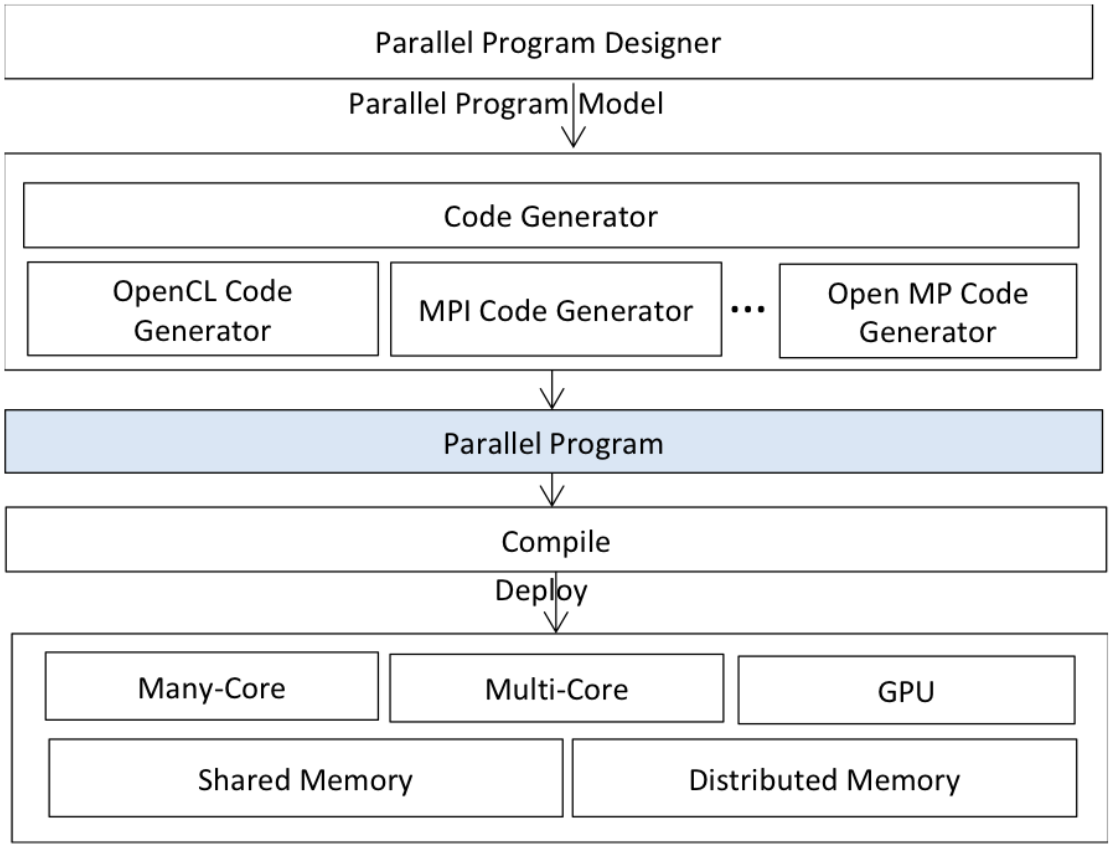
$C = A \times B$, $C_{i,j} = \text{Sum}(A_{i,k} * B_{k,j})$



Web-based development tool



Tool


Parallel Program Designer
Parallel Program Model
Code Generator
OpenCL Code Generator
MPI Code Generator
...
Open MP Code Generator
Parallel Program
Compile
Deploy
Many-Core
Multi-Core
GPU
Shared Memory
Distributed Memory

Evaluation

- Define set of DSVLs to model domain (mollecular similation and signal processing for radio telecopes), architecture (GPU and MPI-based CPU), parallelism, processing models (MapReduce etc)
- Model hand-implemented programs at multiple levels of abstraction including links between models
- Generate C and OpenCL code
- Reverse-engineer (parts of) models from C code
- 3 scientists validating approach iteratively

Summary & Future work

- Integrated web-based development environment for GPU-based (and other) scientific applications
- Flexible DSVL designer with pre-packaged DSVLs and user-defined DSVLs
- Semi-automated roundtrip engineering support: model-> code-> model
- Working on:
  - Improve generation/reverse engineering – C/C++, OpenCL
  - Patterns and critics to guide users, analyse models/code
  - Visualisation of running GPU code in models
  - “Liveness”…? ☺

Questions?

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