

Scientific Applications Development: Why Code When You Can Draw?

Mohamed, John, Richard,
David, Willem & Owen

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▶ think **forward**



Objective

```

const int cRows = 300;
const int cCols = 300;
int matrix_C[cRows][cCols];
//@End
//@Task:init_matrixA
for(int iRows = 0 ; iRows < aRows; iRows++) {
    for(int j = 0 ; j < aCols; j++) {
        matrix_A[iRows][j] = rand();
    }
}
//@End Task:init_matrixA

//@Task:init_matrixB
for(int i = 0 ; i < bRows; i++) {
    for(int j = 0 ; j < bCols; j++) {
        matrix_B[i][j] = rand();
    }
}
//@End Task:init_matrixB

//@Loop:Rows
for(int i = 0; i < aRows ; i++)
{
    for(int j = 0; j < aCols ; j++) {
        for(int k = 0; k < aCols ; k++) {
            //@Task:element_value
            matrix_C[i][j] += matrix_A[i][k] * matrix_B[k][j];
            //@End Task:element_value
        }
    }
}
//@End Loop:Loop

//@Task:print_matrix
for(int i = 0 ; i < cRows; i++) {
    for(int j = 0 ; j < cCols; j++) {
        cout<<matrix_C[i][j]<<endl;
    }
    cout<<"\n";
}
    
```

Sequential Matrix Multiplication

```

kernel void
matrixMul( __global float* C, __global float* A, __global float* B,
           __local float* As, __local float* Bs,
           int uiWA, int uiWB, int trueLocalSize)
{
    for (i=1; i<=
        source = i;
        MPI_Recv(&C
        MPI_Recv(&A
        count = rows
        MPI_Recv(&B

        // Block index
        int bx = get_group_id(0);
        int by = get_group_id(1);

        // Thread index
        int tx = get_local_id(0);
        int ty = get_local_id(1);

        // Index of the first sub-matrix of A processed by the block
        int aBegin = uiWA * BLOCK_SIZE * by;

        // Index of the last sub-matrix of A processed by the block
        int aEnd = aBegin + uiWA - 1;

        // Step size used to iterate through the sub-matrices of A
        int aStep = BLOCK_SIZE;

        // Index of the first sub-matrix of B processed by the block
        int bBegin = BLOCK_SIZE * bx;

        // Step size used to iterate through the sub-matrices of B
        int bStep = BLOCK_SIZE * uiWB;

        // Loop over all the sub-matrices of A and B
        // required to compute the block sub-matrix
        for (int a = aBegin, b = bBegin;
             a <= aEnd;
             a += aStep, b += bStep) {
            // Load the matrices from device memory
            // to shared memory; each thread loads
            // one element of each matrix
            AS(ty, tx) = A[a + uiWA * ty + tx];
            BS(ty, tx) = B[b + uiWB * ty + tx];

            // Synchronize to make sure the matrices are loaded
            barrier(CLK_LOCAL_MEM_FENCE);

            // Multiply the two matrices together;
            // each thread computes one element
            // of the block sub-matrix
            #pragma unroll
            for (int k = 0; k < BLOCK_SIZE; ++k)
                C[a + uiWA * ty + tx] += BS(ty, tx) * A[b + uiWB * ty + tx];
        }

        //MPI_Send(&C
        MPI_Send(&off
        //MPI_Send(&A
        MPI_Send(&rows
        //MPI_Send(&C
        MPI_Send(&C,
    }
}
    
```

Matrix Multiplication - CUDA



Outline

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

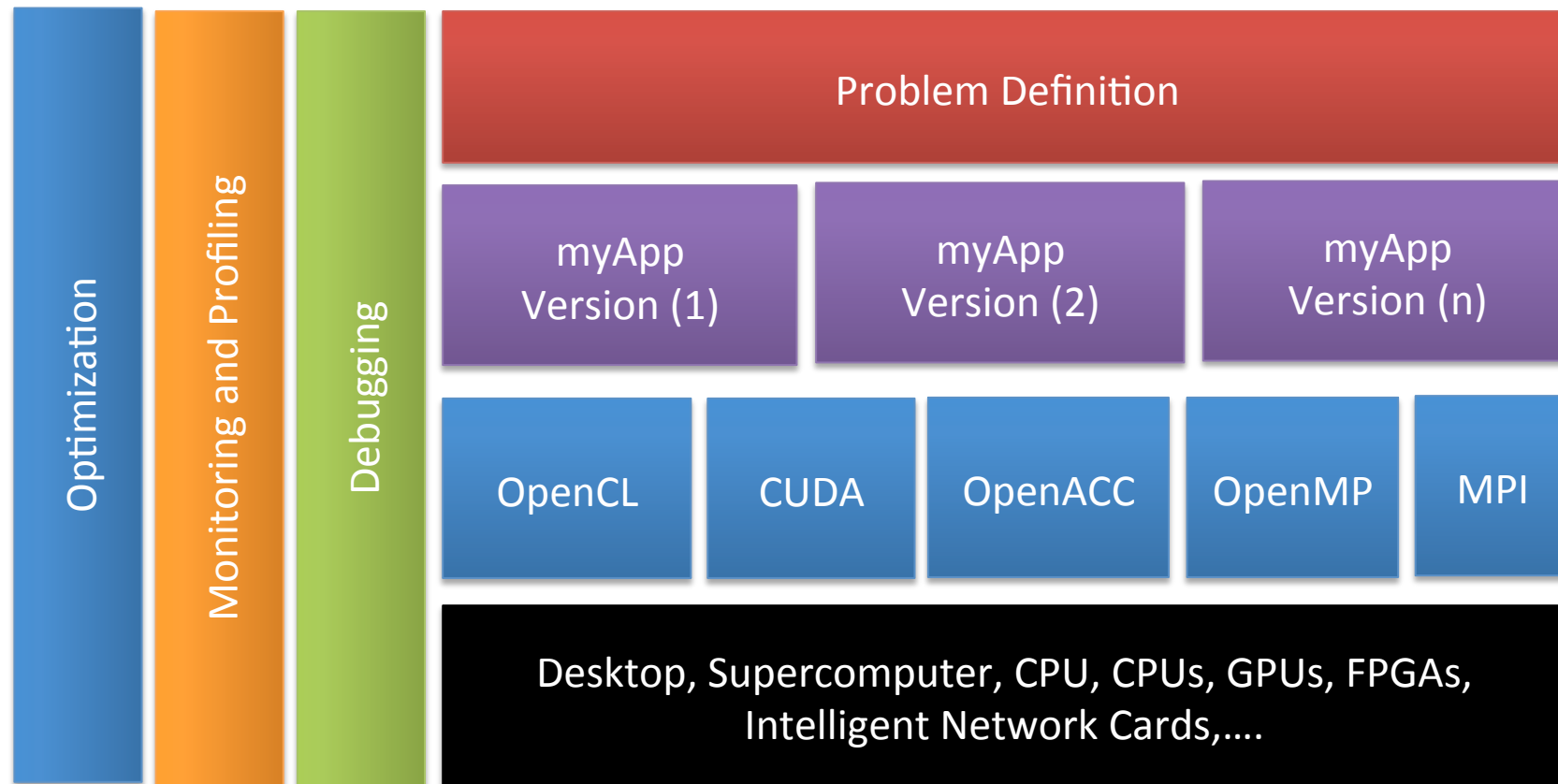


Problem

- Scientists are increasingly developing complex software for data analysis
- *“It’s all about the software...”*
- Most are not trained programmers
- Many are using complex software platforms and techniques e.g. distributed & parallel programming, GPUs, etc - that are hard for experienced CS grads to do
- Approaches to address range from packaged software (Lack flexibility), DSLs (Also Flexibility/Domain issues), programming patterns and toolkits (still complex)



More Problems...



Lack of high-level, human-centric approach to help in developing high quality scientific apps



Motivation – How scientists design their applications...

A

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

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 ens
 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 o
 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 at

Part 2.1.2 Apply integrator to update $r_i, a_i, v_i, (\partial r^n / \partial t^n)_i$.

Part 2.2 if ($t > t_{\text{Equilibration}}$)
 Accumulate averages.
 else if ($\text{mod}(m, \text{scalingInterval}) = 0$)

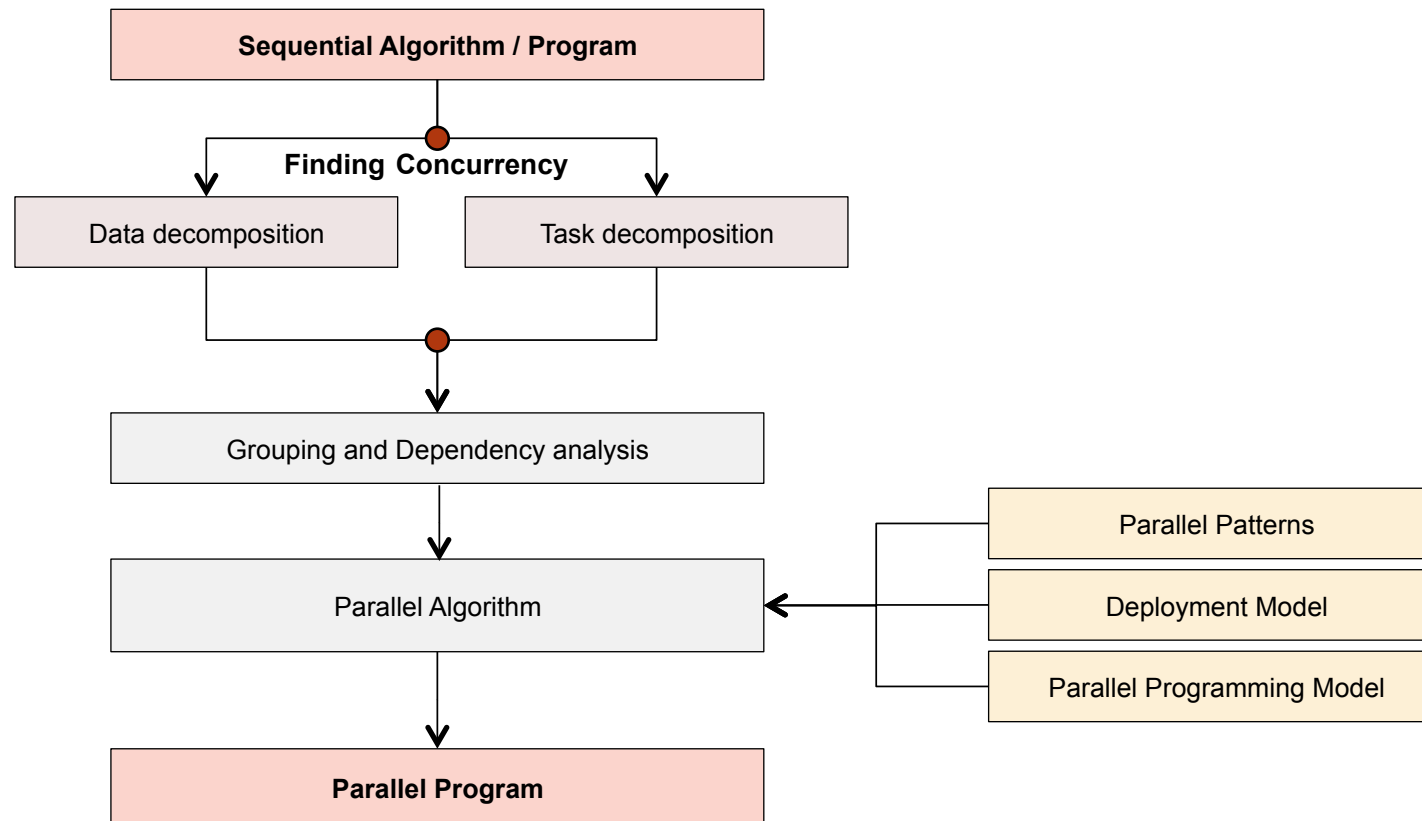
B

Handwritten notes and diagrams:

- $P[\text{ibin}] += \text{amp}[\text{idat}]$
- $N[\text{ibin}] ++$
- $\text{ibin} = P(t) \cdot \text{nbins} \cdot t - \text{idat} \cdot P_s$
- fractional phase
- $k=0$ and $k=1$ labels on a waveform diagram.
- Equation: $\vec{e} = \begin{pmatrix} x(t) \\ y(t) \end{pmatrix}$ and $\vec{e}^* = \begin{pmatrix} x^* & y^* \\ Y^* \end{pmatrix}$



Parallel Program Development Steps



* Timothy. G. Mattson et al., 2004, Patterns for Parallel Programming, Addison Wesley Software Patterns Series.



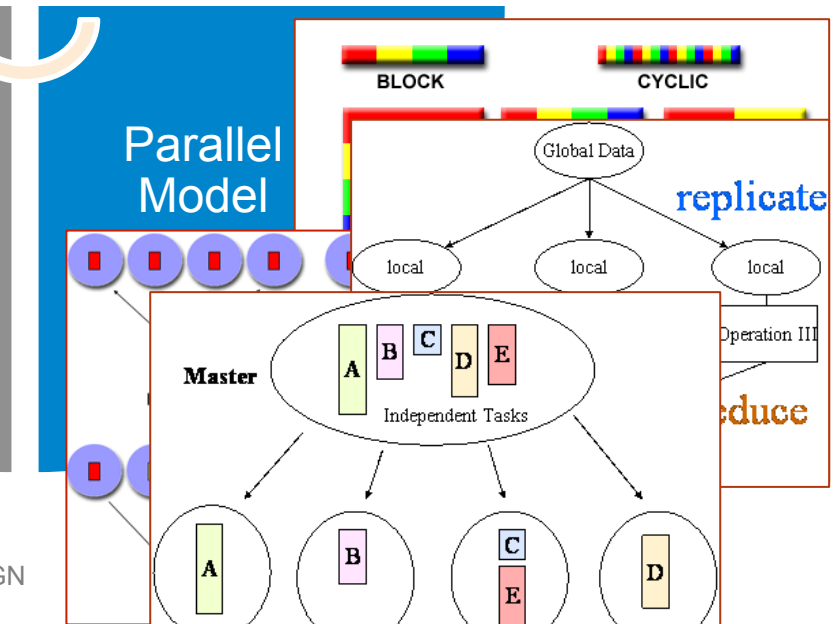
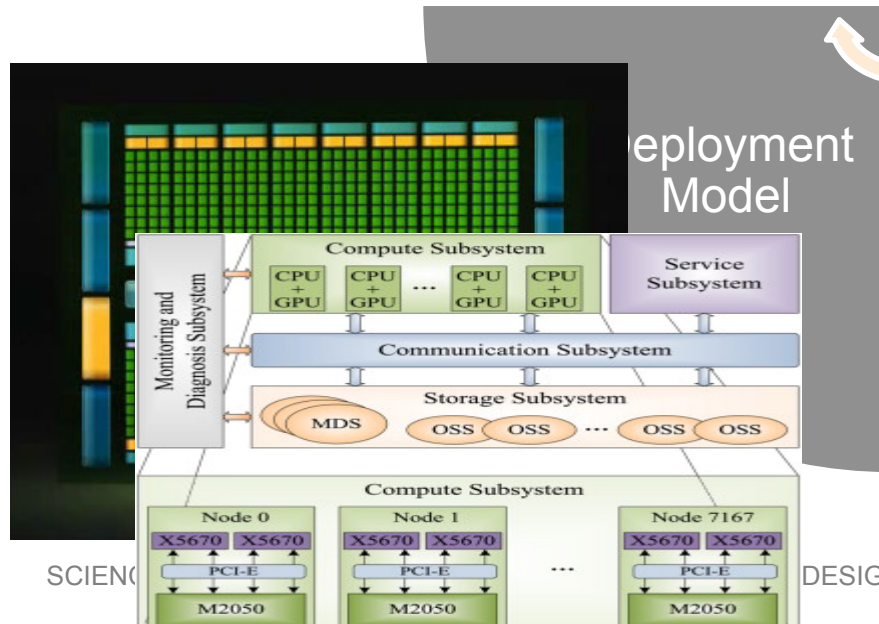
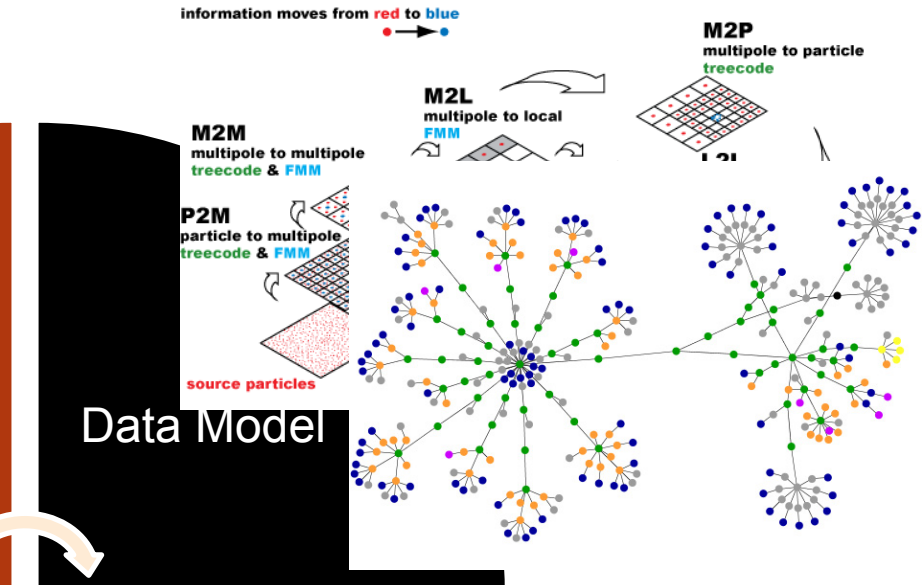
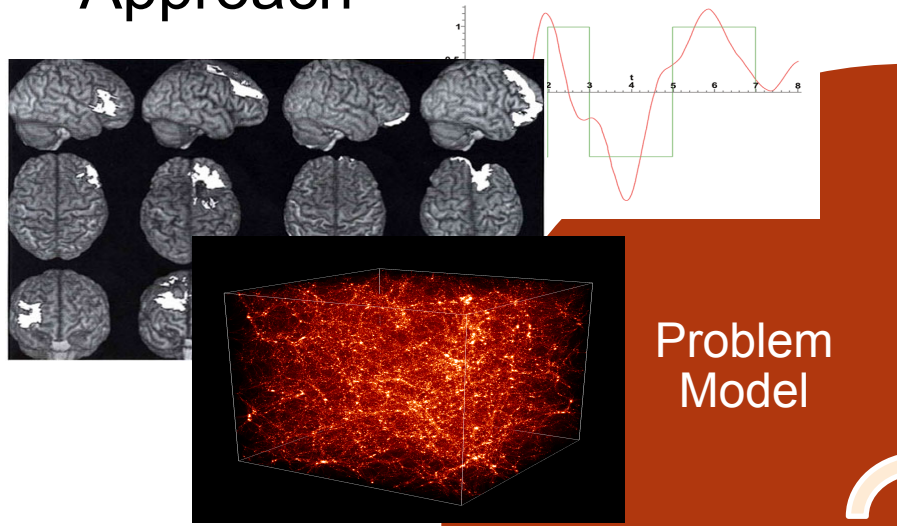
Program Organization vs Parallel Patterns vs Parallel Programming Models

		OpenMP	MPI	CUDA
SPMD	SPMD	☺ ☺ ☺	☺ ☺ ☺ ☺	☺ ☺ ☺ ☺ ☺
Loop Parallel	Loop Parallel	☺ ☺ ☺ ☺	☺	
Master/Worker	Master/Slave	☺ ☺	☺ ☺ ☺	
Fork/Join	Fork/Join	☺ ☺ ☺		

* Timothy. G. Mattson et al., 2004, Patterns for Parallel Programming, Addison Wesley Software Patterns Series.



Approach



SCIENCE

DESIGN

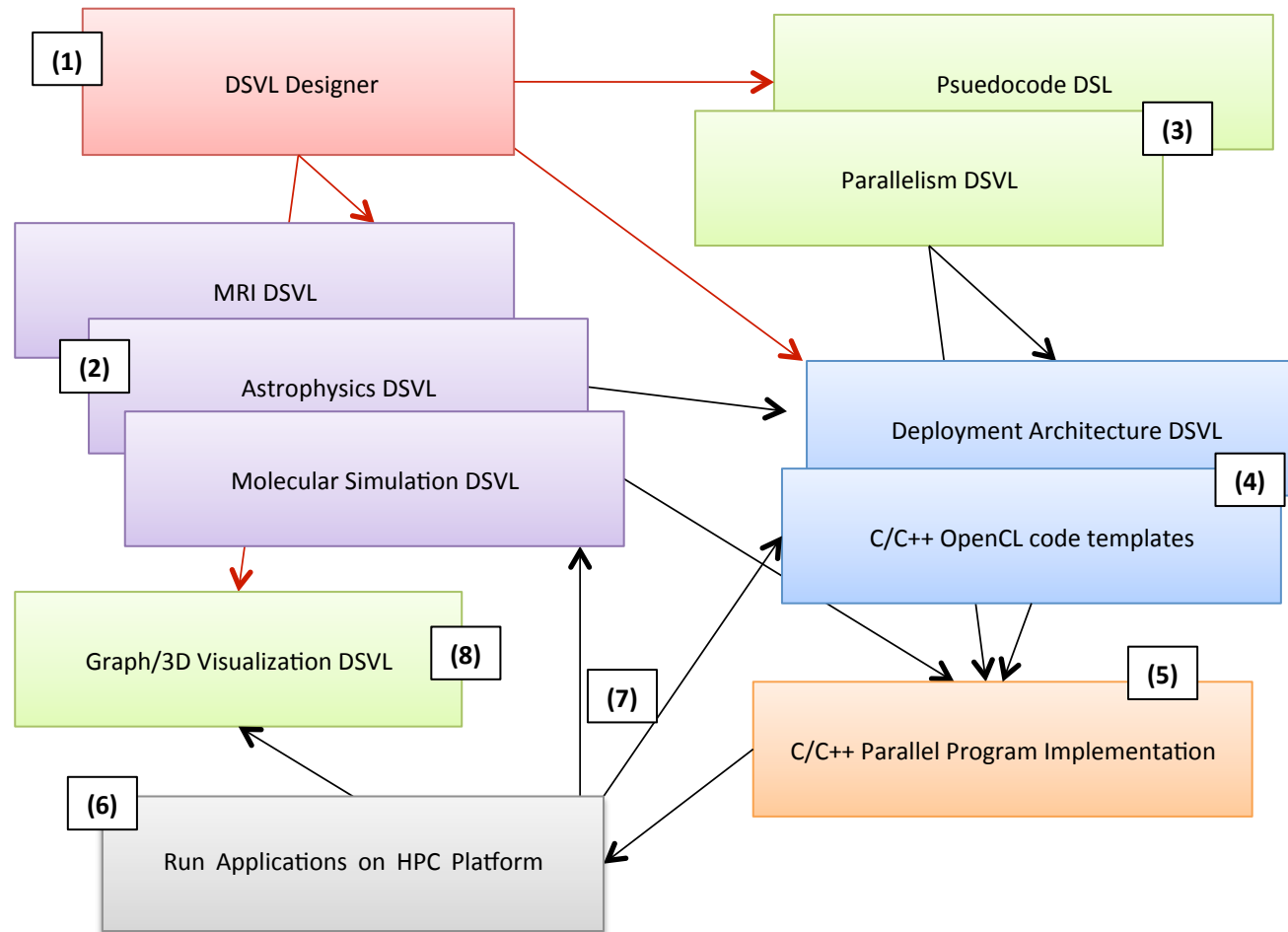


Approach

- 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 DSLs to model
- Provide web-based environment including DSLs designer, coding, debugging, linking DSL views
- Provide semi-automated support for generating lower-level models, generate code/code annotations, reverse-engineer higher-level models from (existing) code

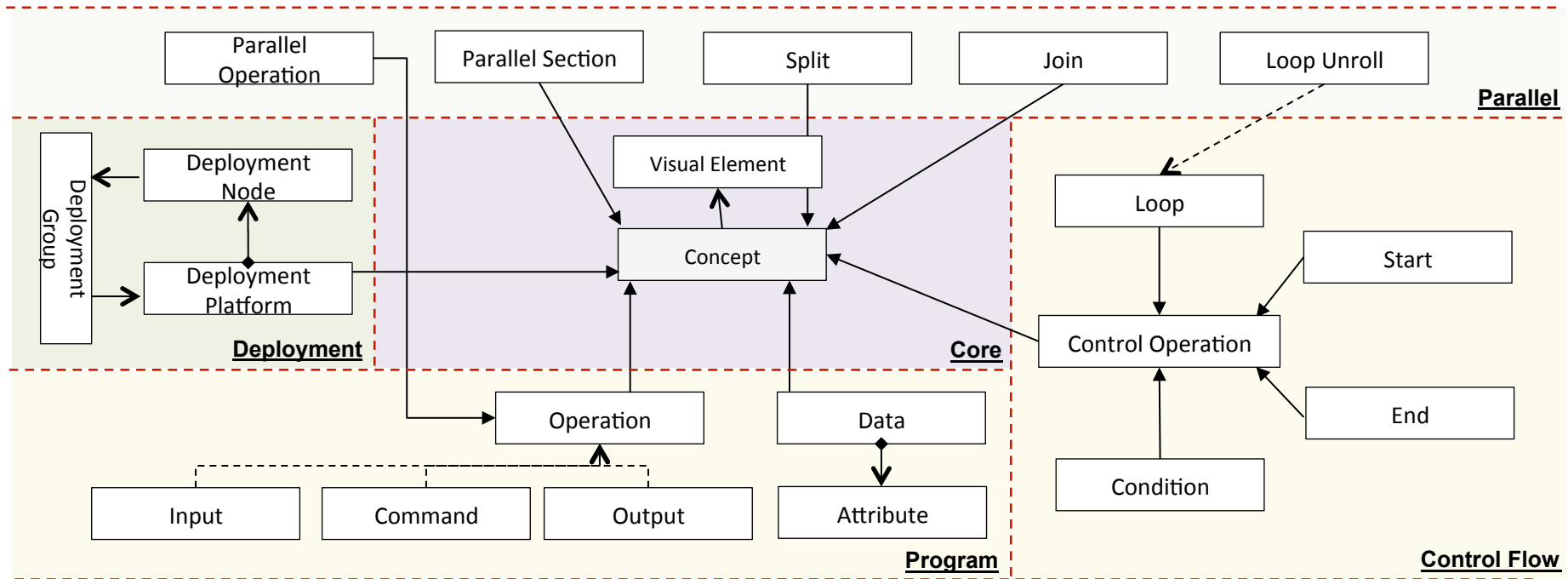


Approach

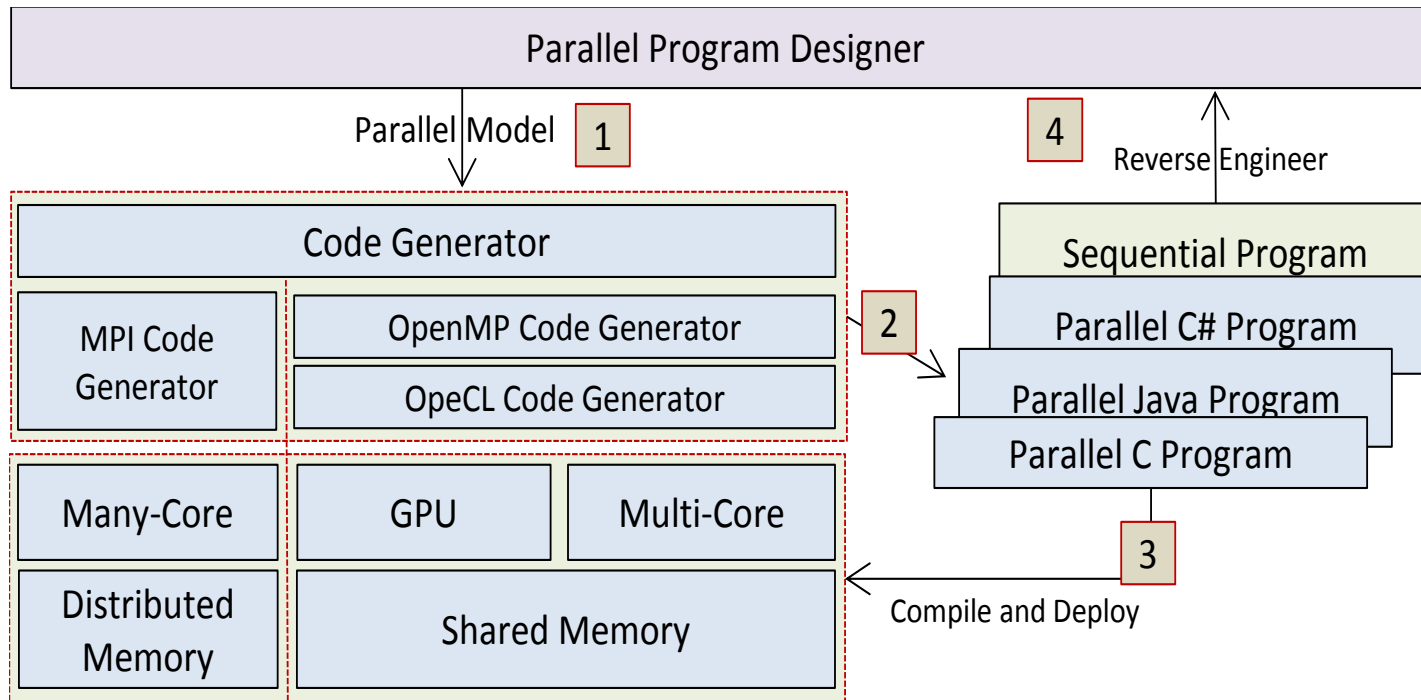




DSVL Designer Meta-model



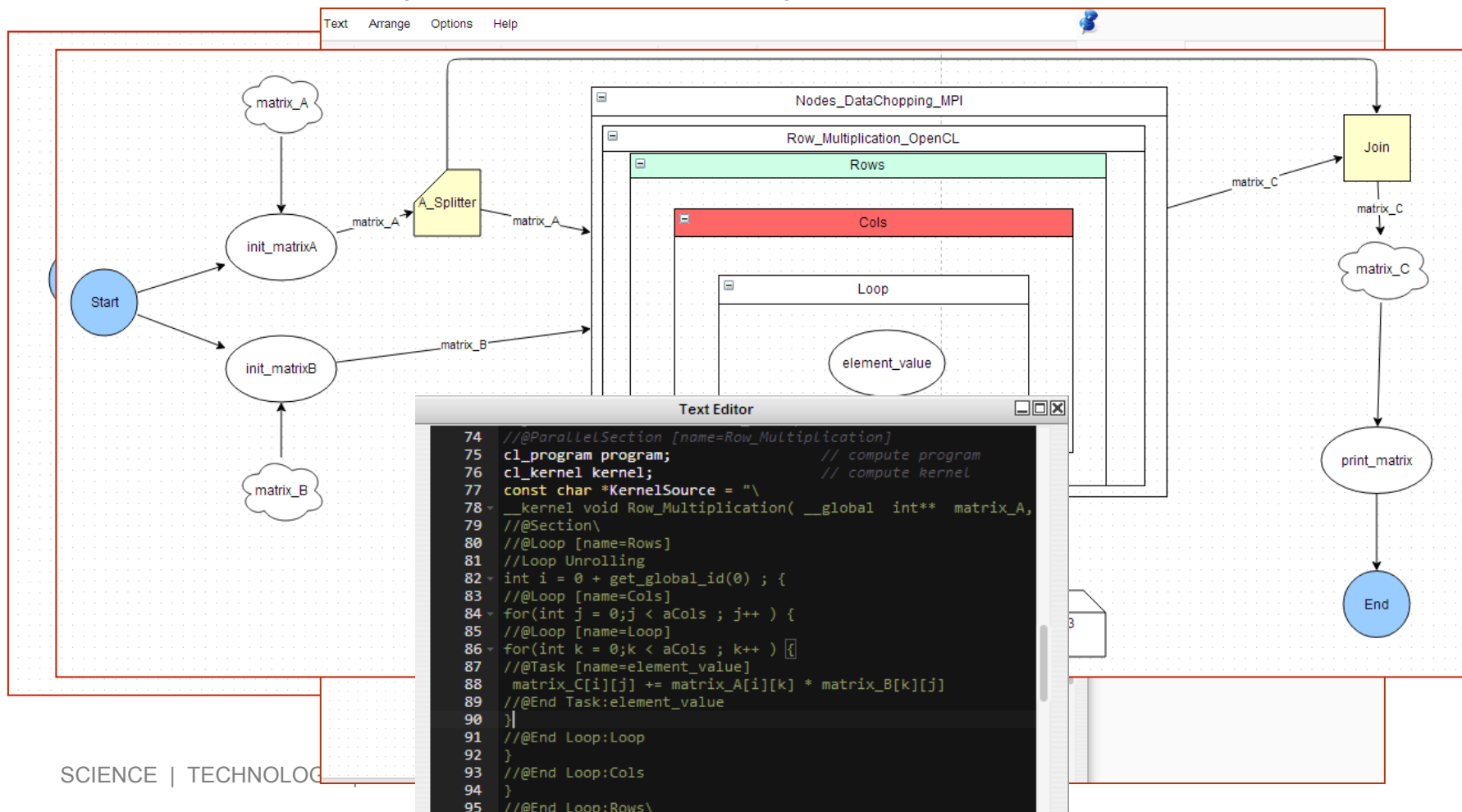
Platform





Example – Matrix Multiplication

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





Snapshots

```
46 //@@End Splitter :A_Splitter
47 //@@ParallelSection [name=matrixKernel]
48 #pragma acc parallel copyin(matrix_A,matrix_B) copyout(matrix_C)
49
50 {
51 //@Section
52 //@Loop [name=Rows]
53 for(int i = 0;i < aRows ; i++ ) {
54 //@Loop [name=Cols]
55 for(int j = 0;j < aCols ; j++ ) {
56 //@Loop [name=Loop]
57 for(int k = 0;k < aCols ; k++ ) {
58 //@Task [name=element_value]
59 matrix_C[i][j] += matrix_A[i][k] * matrix_B[k][j]
60 //@End Task:element_value
61 }
62 //@@End Loop:Loop
63 }
64 //@@End Loop:Cols
65 }
66 //@@End Loop:Rows
67 //@@End Section
68 }
69 //@@End Parallel Section:matrixKernel
70
71 //@Join [name=Join]
72 if(taskid !=0){
73 ierr = MPI Send( matrix_C, 1*aRows/ntasks * 1*aCols , MPI DOUBLE, 0 , 0 , MPI COMM WORLD);
```

Terminal output:

```
compute program
compute kernel
...
ata->nAtom / ntas
...
) {\
```



Summary & Future work

- Integrated web-based development environment for scientific applications
- Flexible DSL designer with pre-packaged DSLs (Parallel DSL, and Deployment DSL) and user-defined DSLs
- Semi-automated roundtrip engineering support: model->code-> model
- Working on:
 - Template-based DSL Designer
 - Patterns and critics to guide users, analyze models/code
 - Visualization of running parallel code onto models

Questions?

Mohamed Almorsy

malmorsy@swin.edu.au



Thanks to ARC Discovery Project scheme
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References

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