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COMPSs Tutorial

February 20th 2014, Barcelona

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Outline (Feb 20th 2014)

- Session 1 / 9am – 11am: Introduction to COMPSs
- Roundtable: presentation and background of participants
- Programming model
  - Overview
  - Steps
  - Properties
- COMPSs runtime system
  - Overview
  - Features
- Coffee break – 11:00 – 11:30
- Session 2 / 11:30am – 1pm: Application examples
  - Sample codes & Demos
    - Matmul
  - Graphical interface (IDE)
    - Gene Detection

Outline (Feb 20th 2014)

- Lunch Break 1pm to 2pm
- Session 3 / 2 pm- 3:30 pm: Hands-on I
  - Virtual Machine Setup
  - BLAST overview
  - Code modification
    - All-to-one and tree-reduction
  - Compilation and Execution
- Coffee break: 3:30 – 4:00
- Session 4 / 4 pm- 6 pm: Hands-on II
  - HMMER overview & code modification
  - Configuration, monitoring, debugging
  - Overview of tracing, trace analysis
  - IDE Hands-on
  - Final notes

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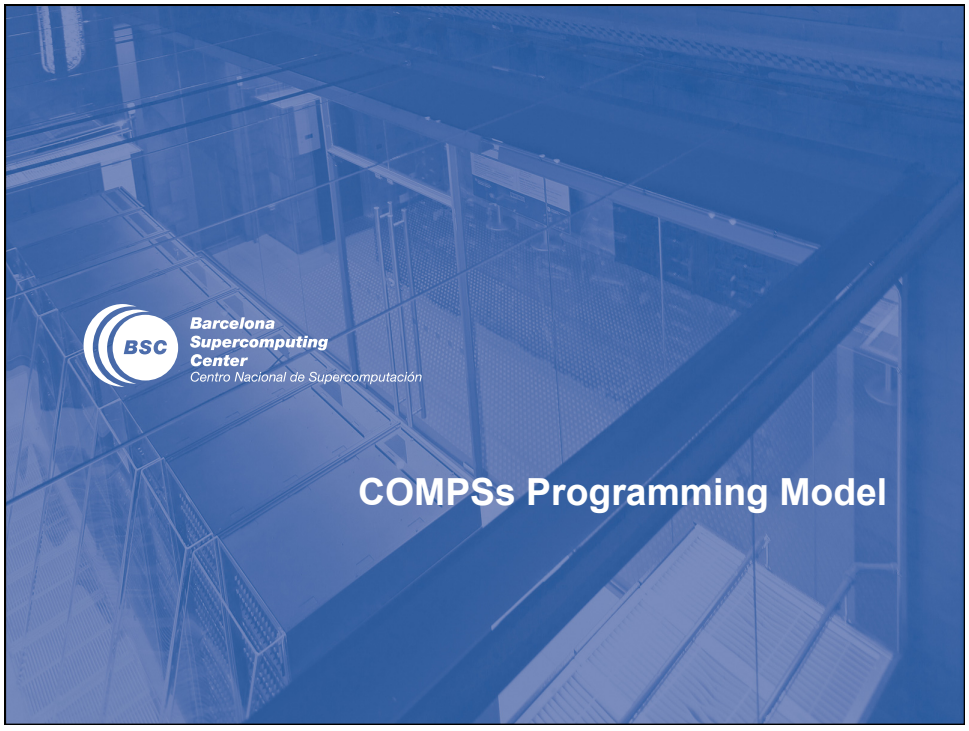
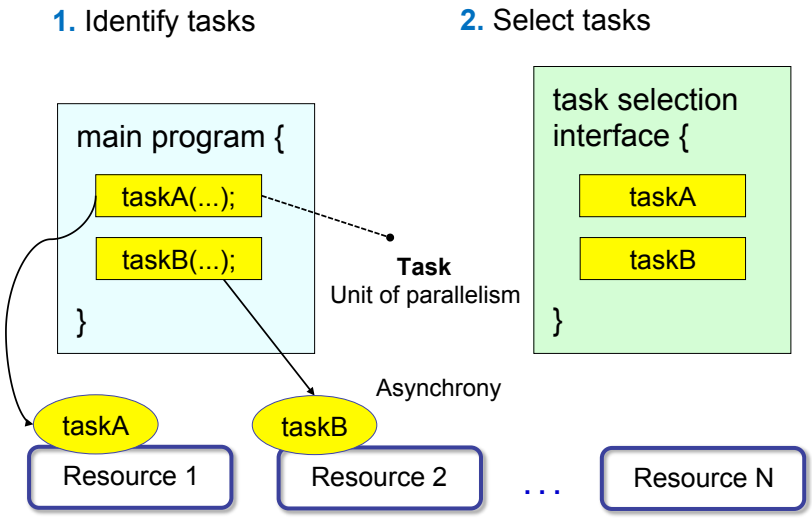


Overview: Objectives

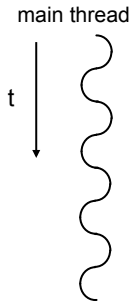
- Reduce the development complexity of Grid/Cluster/ Cloud applications to the minimum
  - Writing an application for a computational distributed infrastructure may be as easy as writing a sequential application
- Target applications: composed of tasks, most of them repetitive
  - Granularity of the tasks or programs
  - Data: files, objects, arrays and primitive types

Programming Model: Steps

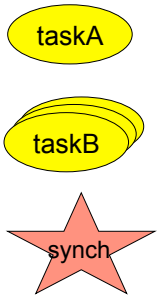


Programming Model: Properties (I)

- Based on pure-Java fully-sequential programming
  - No APIs, no threading, no messaging
  - No parallel constructs, no pragmas
  - Sequential consistency




```
Main Program {
  taskA(data1);

  for (int i=0; i< N; i++)
    taskB(data1, data2);

  if (condition)
    process(data2);
}
```

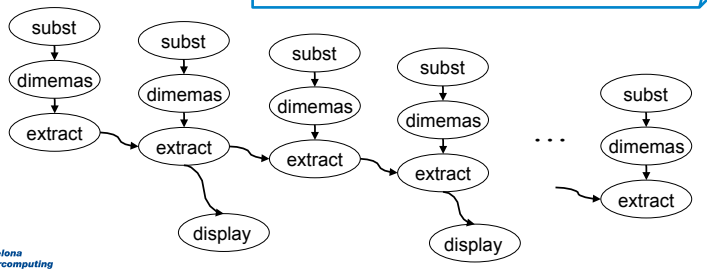


Programming Model: Dependency detection

- Automatic on-the-fly creation of a task dependency graph

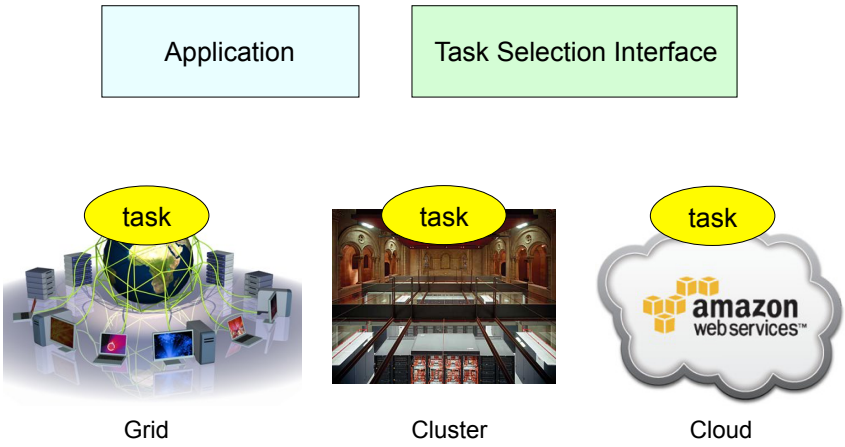
Main Program

```
for (int i = 0; i < N; i++) {
   newBWD = random();
   subst(refCFG, newBWD, newCFG);
   dimemas(newCFG, trace, dimOUT);
   extract(newBWD, dimOUT, finalOUT);
   if (i % 2 == 0) display(finalOUT);
}
```



Programming Model: Properties (II)

- Infrastructure unaware



Programming Model: Task selection interface

```
public interface SampleItf {
   @Constraints(processorCPUCount = 1, memoryPhysicalSize = 0.5f)
   @Method(declaringClass = "servicess.Example")
   void myMethod(
      @Parameter(direction = INOUT)
      Reply r
    );

   @Service(namespace = "http://servicess.es/example",
            name = "SampleService",
            port = "SamplePort")
   Reply myServiceOp(
      @Parameter(direction = IN)
      Query q
    );
}
```

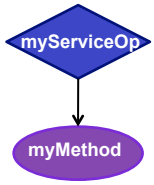
Programming Model: Regular Main program

```
public class App {

   public static void main(String[] args) {
       Query query = new Query(…);
       Reply reply = myServiceOp(query);

       myMethod(reply);

       reply.printToLog();
   }
}
```

Service task call

Method task call

Synchronization



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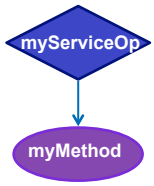
Programming Model: Service Operation

```
public class ServiceApp {
   @Orchestration
   public static void sampleComposite() {
       Query query = new Query(…);
       Reply reply = myServiceOp(query);

       myMethod(reply);

       reply.printToLog();
   }
}
```



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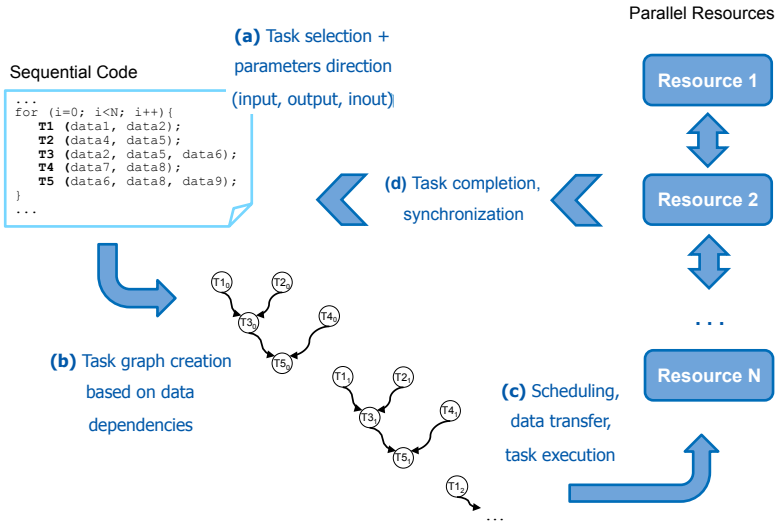
Programming Model: Summary



Programming Model: Sample Application

```
public static void main(String[] args) {
        String counter1 = args[0], counter2 = args[1],
                          counter3 = args[2];

        initializeCounters(counter1, counter2, counter3);

        for (i = 0; i < 3; i++) {
          increment(counter1);
          increment(counter2);
          increment(counter3);
        }
}
```

Main program

Subroutine

```
public static void increment(String counterFile) {
        int value = readCounter(counterFile);
        value++;
        writeCounter(counterFile, value);
}
```

Programming Model: Sample App (Interface)

Task selection interface

```
public interface SimpleItf {

        @Method(declaringClass = "SimpleImpl")
        void increment(
                @Parameter(type = FILE, direction = INOUT)
                String counterFile
        );

}
```

Implementation

Parameter metadata

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Programming Model: Final App Code

```
public static void main(String[] args) {
        String counter1 = args[0], counter2 = args[1],
                                   counter3 = args[2];

        initializeCounters(counter1, counter2, counter3);

        for (i = 0; i < 3; i++) {
          increment(counter1);
          increment(counter2);
          increment(counter3);
        }
}
```

Main program of the application NO CHANGES!

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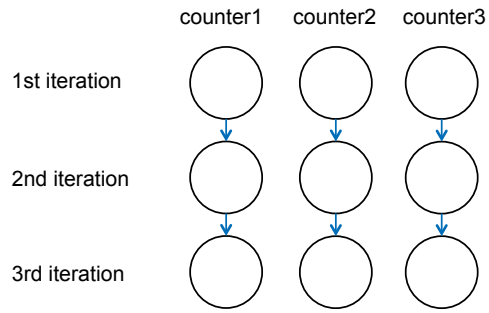
Programming Model: Task Graph

Main loop

```
for (i = 0; i < 3; i++) {
        increment(counter1);
        increment(counter2);
        increment(counter3);
}
```

Task graph



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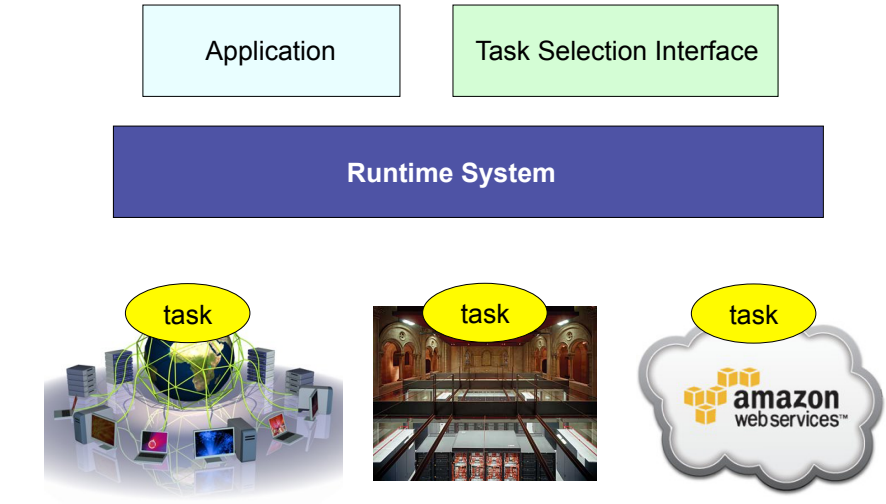
COMPSs Runtime System

Runtime System



Basic features

- Supported Features:
  - Data dependency analysis
  - Data renaming
  - Data transfer
  - Task scheduling
  - Resource management
  - Results collection
  - Fault tolerance
  - Shared disks management
- In Progress:
  - Checkpointing
  - Constraint matching
  - Task nesting

Interoperability

Grid/Cluster Configuration: Resources Specification

Resources.xml

```
<?xml version="1.0" encoding="UTF-8"?>
<ResourceList>
    <!--Description for any physical node-->
    <Resource Name="172.20.200.18">
        <Capabilities>
            <Host>
                <TaskCount>0</TaskCount>
                <Queue>short</Queue>
                <Queue/>
            </Host>
            <Processor>
                <Architecture>IA32</Architecture>
                <Speed>3.0</Speed>
                <CPUCount>1</CPUCount>
            </Processor>
            <OS>
                <OSType>Linux</OSType>
                <MaxProcessesPerUser>32</MaxProcessesPerUser>
            </OS>
            <StorageElement>
                <Size>30</Size>
            </StorageElement>
            …
            <Memory>
                <PhysicalSize>1</PhysicalSize>
                <VirtualSize>8</VirtualSize>
            </Memory>
            <ApplicationSoftware>
                <Software>Java</Software>
            </ApplicationSoftware>
            <Service/>
            <VO/>
            <Cluster/>
            <FileSystem/>
            <NetworkAdaptor/>
            <JobPolicy/>
            <AccessControlPolicy/>
        </Capabilities>
        <Requirements/>
    </Resource>
    <Resource Name="172.20.200.19">
        …
    </Resource>
</ResourceList>
```

Grid/Cluster Configuration: Project Specification

Project.xml

```
<?xml version="1.0" encoding="UTF-8"?>
<Project>
      <!--Description for any physical node-->
      <Worker Name="172.20.200.18">
            <InstallDir>/opt/COMPSs/Runtime/scripts/</InstallDir>
            <WorkingDir>/tmp/</WorkingDir>
            <User>user</User>
            <LimitOfTasks>1</LimitOfTasks>
      </Worker>

      <Worker Name="172.20.200.19">
        …
      </Worker>
      ….
</Project>
```

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Cloud Configuration: Resources Specification

Resources.xml

```
<ResourceList>
   <CloudProvider name="BSCCloud">
         <Server>https://bscgrid20.bsc.es:8443/DRP</Server>
         <Connector>
               integratedtoolkit.connectors.emotivecloud.DRPSecureClientConnector
         </Connector>
         <ImageList>
           <Image name="debianbase"/>
         </ImageList>
         <InstanceTypes>
               <Resource Name="bsc.small">
                     <Capabilities>
                           <Processor>
                                 <CPUCount>1</CPUCount>
                           </Processor>
                           <StorageElement>
                                 <Size>0.5</Size>
                           </StorageElement>
                           <Memory>
                                 <PhysicalSize>1</PhysicalSize>
                           </Memory>
                     </Capabilities>
               </Resource>
               <Resource Name="bsc.medium">
                 …
               </Resource>
         </InstanceTypes>
   </CloudProvider>
</ResourceList>
```

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Cloud Configuration: Project Specification

Project.xml

```
<Project>
 <Cloud>
        <InitialVMs>0</InitialVMs>
        <minVMCount>2</minVMCount>
        <maxVMCount>5</maxVMCount>

        <Provider name="BSCCloud">
            <LimitOfVMs>5</LimitOfVMs>
            <Property>
                <Name>Cert</Name>
                <Value>/home/.../cert.p12</Value>
            </Property>
            <Property>
                <Name>Owner</Name>
                <Value>userbsc</Value>
            </Property>
            <Property>
                <Name>JobNameTag</Name>
                <Value>Job</Value>
            </Property>
    ...
```

```
...
<ImageList>
        <Image name="debianbase">

            <InstallDir>/opt/COMPSs/Runtime/scripts</InstallDir>
                <WorkingDir>/tmp/</WorkingDir>
                <User>user</User>
                 <Package>
                  <Source>/home/.../AppName.tar.gz</Source>
                  <Target>/home/user</Target>
                 </Package>
        </Image>
    </ImageList>
    <InstanceTypes>
        <Resource name="bsc.small"/>
    </InstanceTypes>
</Provider>
</Cloud>
</Project>
```

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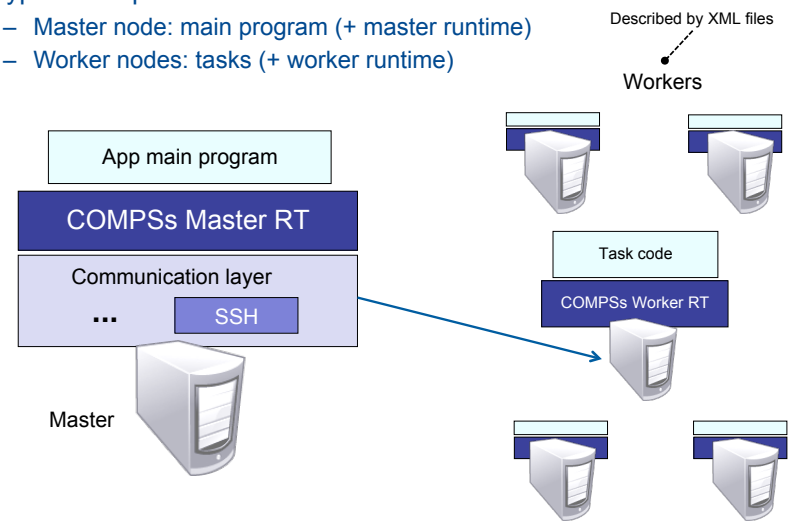
COMPSs in a Cluster (interactive)

- Typical setup:
  - Master node: main program (+ master runtime)
  - Worker nodes: tasks (+ worker runtime)



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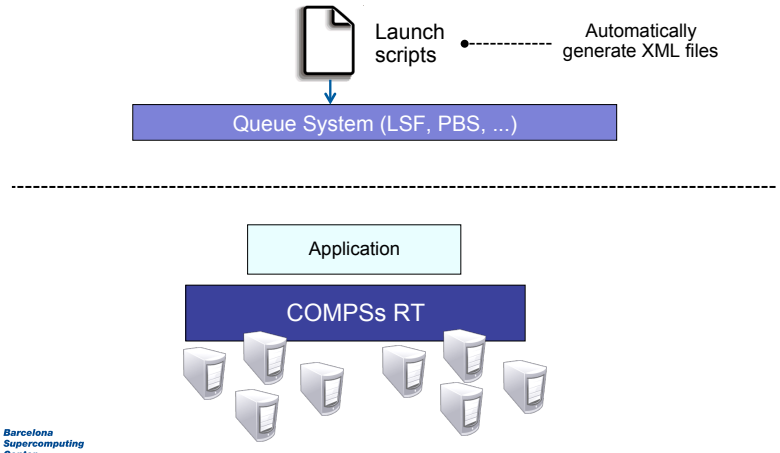
COMPSs in a Cluster (queue system)

- Execution divided in two phases
  - Launch scripts queue a whole COMPSs app execution
  - Actual execution starts when reservation is obtained



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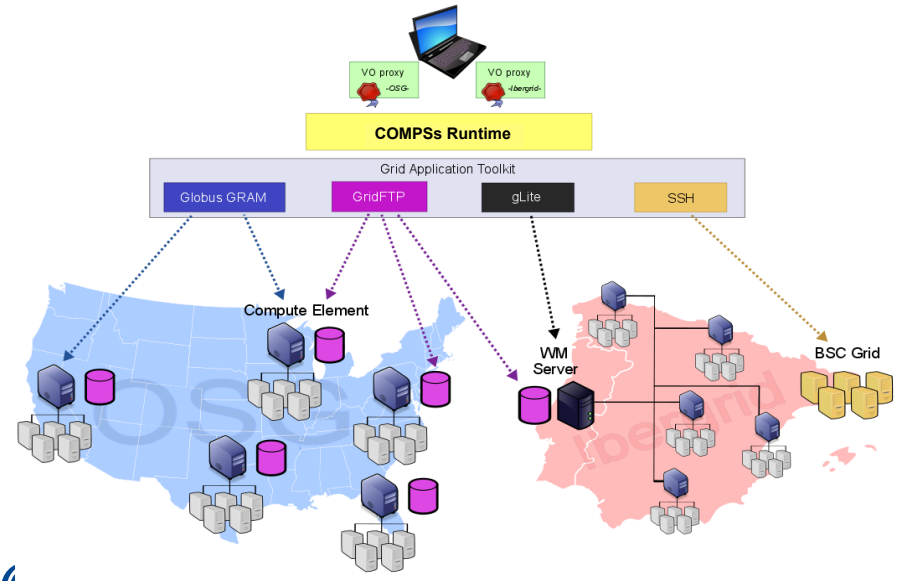
COMPSs in a Grid



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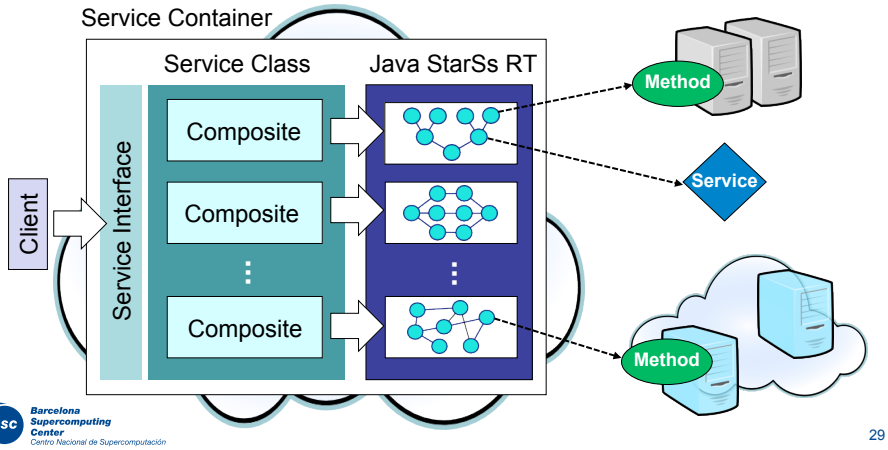
COMPSs in the Cloud

- Runtime integrated in a platform with:
  - Service orientation
  - Virtualization



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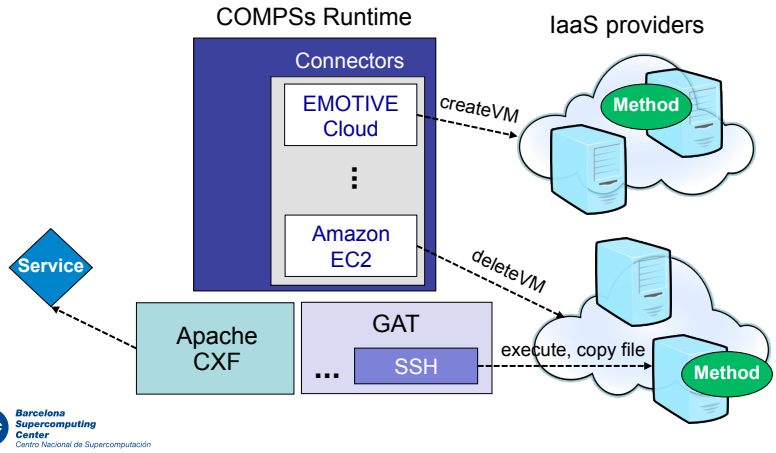
Cloud: Connector design

- Interaction with:
  - Cloud providers: connectors, SSH
  - Service providers: WS client



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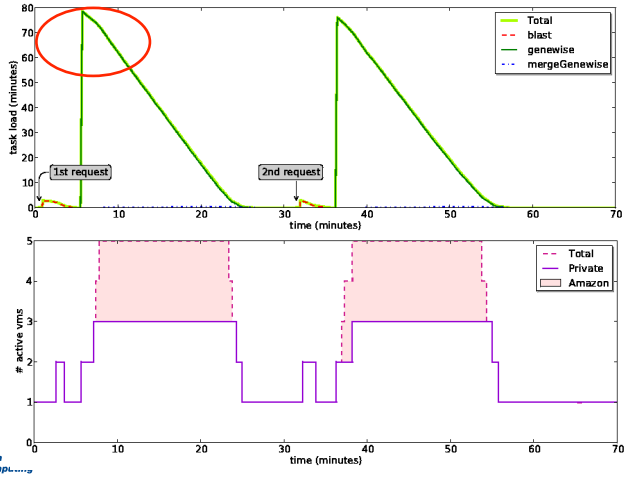
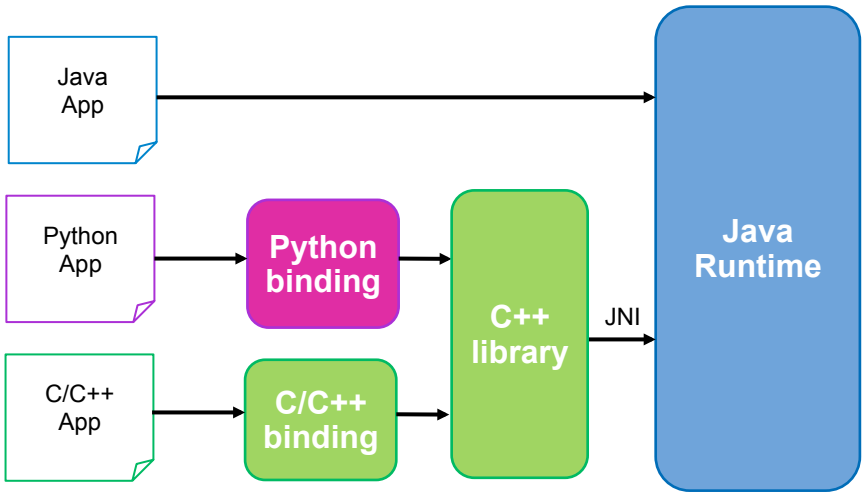
Cloud: Elasticity

- Increase/decrease number of VMs depending on task load
- Bursting to Amazon EC2 to face peak load



COMPSs Bindings



Matmul example

```
for (int i = 0; i < MSIZE; i++){
        for (int j = 0; j < MSIZE; j++)
                for (int k = 0; k < MSIZE; k++)
                {
                        long ini, fi;
                        ini = System.currentTimeMillis();
                        MatmulImpl.multiplyAccumulative( _C[i][j], _A[i][k], _B[k][j] );
                        fi = System.currentTimeMillis();
                        System.out.println("TASK: " + ((fi - ini) / 1000) + " seconds\n");
                }
}
```

```
public static void multiplyAccumulative( String f3, String f1, String f2 )
{
        Block a = new Block( f1 );
        Block b = new Block( f2 );
        Block c = new Block( f3 );

        c.multiplyAccum( a, b );
        try
                ...
}

public void multiplyAccum ( Block a, Block b )
{
        for( int i = 0; i < this.bRows; i++ )                   // rows
                for( int j = 0; j < this.bCols; j++ )           // cols
                        for ( int k = 0; k < this.bCols; k++ )  // cols
                                this.data[i][j] += a.data[i][k] * b.data[k][j];
}
```

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Matmul interface

```
package matmul;

import integratedtoolkit.types.annotations.Constraints;
import integratedtoolkit.types.annotations.Method;
import integratedtoolkit.types.annotations.Parameter;
import integratedtoolkit.types.annotations.Parameter.*;


public interface MatmulItf {
        @Constraints(processorCPUCount = 4, memoryPhysicalSize = 1.5f)
        @Method(declaringClass = "matmul.MatmulImpl")
        void multiplyAccumulative(
                @Parameter(type = Type.FILE, direction = Direction.INOUT)
                String file1,

                @Parameter(type = Type.FILE, direction = Direction.IN)
                String file2,

                @Parameter(type = Type.FILE, direction = Direction.IN)
                String file3
        );

}
```


Matmul: Compiling

- Compiling with command line:
  - cd workspace
  - javac matmul/src/matmul/*.java
  - cd matmul/src/
  - jar cf matmul.jar matmul

- From eclipse:
  - Package Explorer -> Project (matmul) -> Export…

Matmul: Deploying

- In this case, in the same machine
  - Copy to home directory
  - cd
  - cp ./matmul/src/matmul.jar .
- In remote machines
  - Code needs to be transfer to machine that will host main code

Matmul: Executing

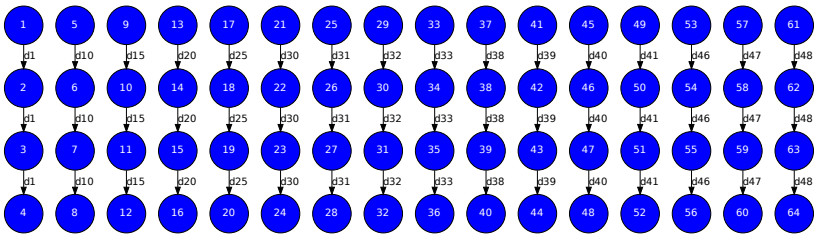
- Set CLASSPATH
  - export CLASSPATH=$CLASSPATH:/home/user/matmul.jar
  - runcompss matmul.Matmul 4

Matmul: Monitoring execution

- Browse
  - http://localhost:8080/compss-monitor

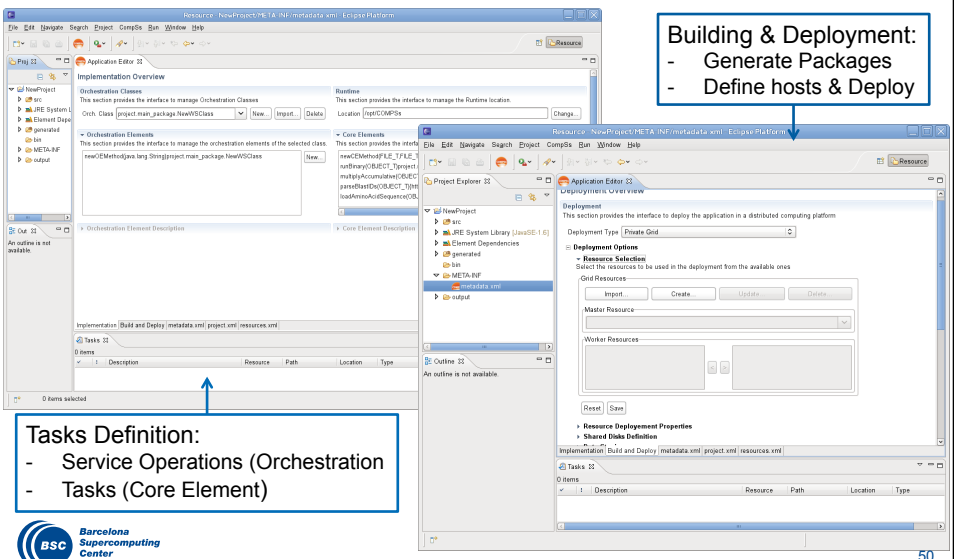
Demos: Matmul

- Blocks matrixes multiplication



IDE COMPSs applications as a Service

- IDE for implementing and deploying applications



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Demos: Gene Detection Application

- Gene Detection algorithm designed by the BSC Life Sciences department
  - Automatic Homology-based gene detection and analysis
- Combine services with computations
  - Example that shows different capabilities of COMPSs
  - Implicit Synchronization points
  - Different method and service call types
  - Objects and files as parameters

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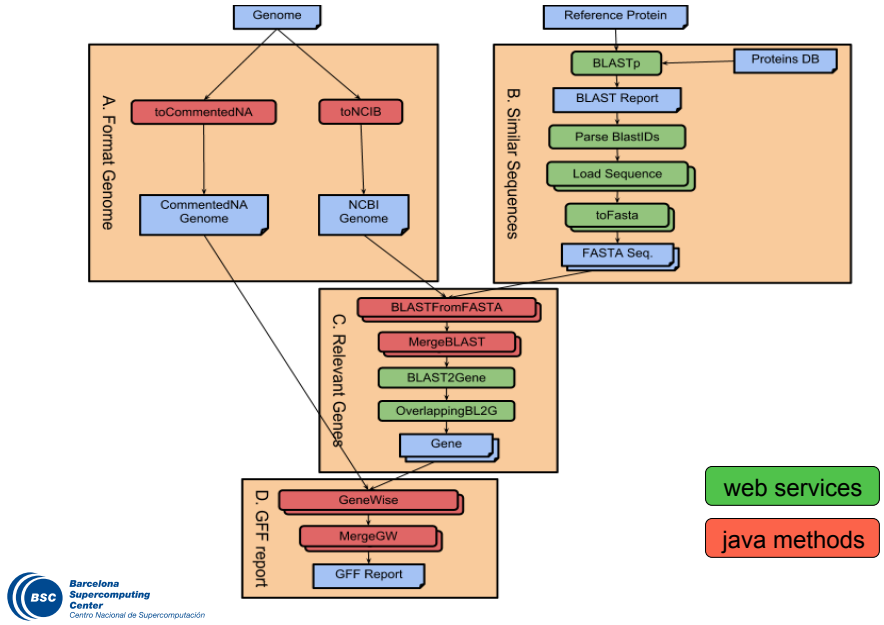
Gene Detection



Hands-On: Overview

- COMPSs Virtual Machine setup
- Applications Overview (BLAST, HMMER, …)
  - Code modification
  - Configuration, compilation & execution
  - Monitoring, debugging
  - Overview of tracing and trace performance analysis
  - IDE

COMPSs development VM Installation

- **COMPSs Development & Test VM (64-bit) OVA**
  - **Available from USB**
- **Import the virtual appliance in VirtualBox**

BLAST: Hands-on

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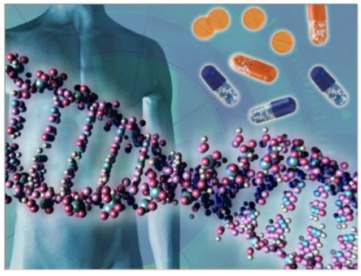
BLAST: Hands-on

Bioinformatics Scenario

- BLAST (Basic Local Alignment Search Tool) Suite:
  - BLAST: An algorithm for comparing primary biological sequence information, such as the amino-acid sequences of different proteins or nucleotides of DNA sequences.

BLAST enables a researcher to compare a query sequence with a library or database of sequences, and identify sequences that resemble the query sequence above a certain threshold.



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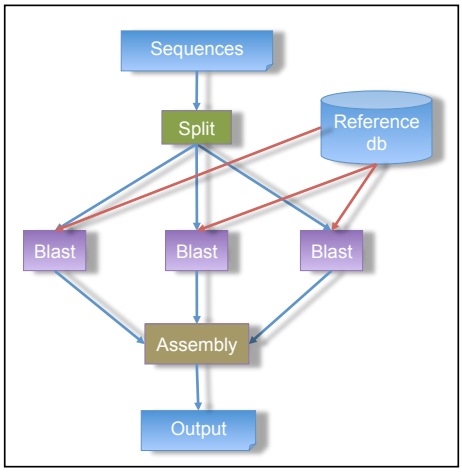
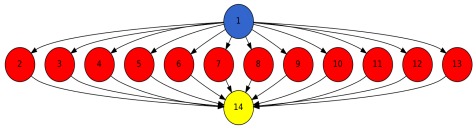
BLAST: Hands-on

- **BLAST**

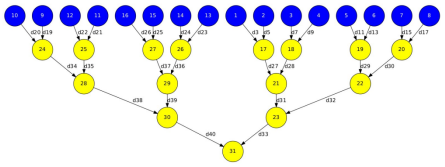


All-to-One Reduction:



OR

Tree-based Reduction:



BLAST: All-to-One reduction

- Main Application (All-to-One):

```
public static void main(String args[]) throws Exception {

  sequences[] = splitSequences(inputFile, nFrags);

  for (partition: sequences)
  {
    BlastImpl.align(database, partition, partitionOutput, blastBinary, commandArgs);
    partitionOutputs.add(partitionOutput);
  }

  assemblyPartitions(partialOutputs, outputFileName, tempDir, nFrags);
}
```

BLAST: All-to-One reduction

- Remote task implementation:

```
public class BlastImpl{

   public void align(String databasePath, String partitionFile,
                     String partitionOutput, String blastBinary, String commandArgs)
   {
     String cmd = blastBinary+ " " +"-p blastx -d " + databasePath + " -i " +partitionFile+ " -o "+
            partitionOutput+ " " +commandArgs;

     Process simProc = Runtime.getRuntime().exec(cmd);
       .......
   }
}
```


BLAST: All-to-One reduction

Creation of the annotated interface for the selection of remote tasks

```
public interface BlastItf {

    @Method(declaringClass = "blast.BlastImpl")
    @Constraints(processorCPUCount = 4, memoryPhysicalSize = 4.0f)
    void align(
        @Parameter(type = Type.STRING, direction = Direction.IN)
        String databasePath,

        @Parameter(type = Type.FILE, direction = Direction.IN)
        String partitionFile,

        @Parameter(type = Type.FILE, direction = Direction.OUT)
        String partitionOutput,

        @Parameter(type = Type.STRING, direction = Direction.IN)
        String blastBinary,

        @Parameter(type = Type.STRING, direction = Direction.IN)
        String commandArgs);
}
```

BLAST: Compilation and execution

- Compilation (Eclipse IDE)
  - ***Package Explorer -> Project (blastallone) -> Export…***
- Usage
  - ***runcompss blast.Blast <debug> <binary> <database> <sequences> <#fragments> <tmpdir> <output>***
- Execution
  - ***cp ~/workspace/blastallone/jar/blast.jar ~***
  - ***export CLASSPATH=$CLASSPATH:/home/user/blast.jar***
  - ***runcompss blast.Blast true /home/user/workspace/blast/binary/blastall / sharedDisk/Blast/databases/swissprot/swissprot /sharedDisk/Blast/ sequences/sargasso_test.fasta 4 /tmp/ /home/user/out.txt_***

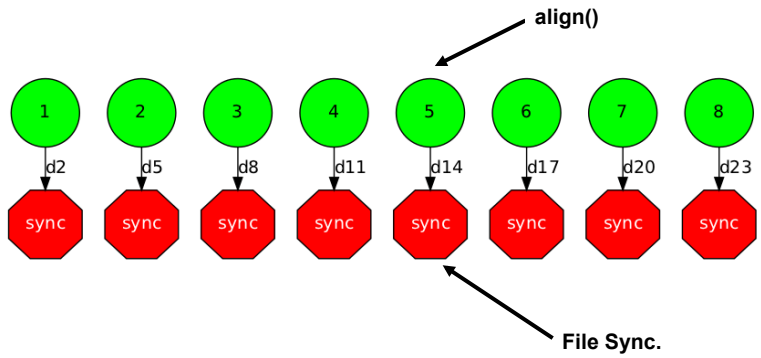
BLAST: Compilation and execution

```
---------- Executing blast.Blast in IT mode total----------

…
BLAST Sequence Alignment Tool:

Parameters:
- Debug Enabled
- Blast binary: /home/user/workspace/blastAllOne/binary/blastall
- Number of expected fragments: 8
- Database Name with Path: /sharedDisk/Blast/databases/swissprot/swissprot
- Database Name: swissprot
- Input Sequences File: /sharedDisk/Blast/sequences/sargasso_test.fasta
- Temporary Directory: /tmp/
- Output File: /home/user/IT/blast.Blast/out.txt
- Command Line Arguments:

- The total number of sequences is: 20
- The total number of sequences of a fragment is: 3

- Splitting sequences among fragment files...
[  API]  -  Opening file /tmp/seqFile1b495168-e913-430a-a347-9894015911e1.sqf in mode APPEND
…

Aligning Sequences:

 - Number of fragments to assemble -> 8

[  API]  -  Opening file /home/user/IT/blast.Blast/out.txt in mode WRITE
 - Assembling partial output -> /tmp/resFile1b495168-e913-430a-a347-9894015911e1.result.txt to final output file -> /home/user/IT/blast.Blast/out.txt
 …
 - Assembling partial output -> /tmp/resFile270855af-307b-4a1e-bc42-0e0cf22256ae.result.txt to final output file -> /home/user/IT/blast.Blast/out.txt
-Sequences assembled in 184 seconds
 …
------------------------------------------------------------
```

BLAST: All-to-One (work)

- Generate the final graph
- Launch BLAST (All-to-One)

BLAST: All-to-One (Graph)



BLAST: Tree-based reduction (work)

- Code the final reduction and its interface.

BLAST: Tree-based reduction

- Main Application (Tree-based):

```
public static void main(String args[]) throws Exception {

  sequences[] = splitSequences(inputFile, nFrags);

  for (partition: sequences)
  {
     BlastImpl.align(database, partition, partitionOutput, blastBinary, commandArgs);
     partitionOutputs.add(partitionOutput);
  }

  //Final Assembly process -> Merge 2 by 2
    int neighbour=1;
    while (neighbour<partialOutputs.size()){
       for (int result=0; result<partialOutputs.size(); result+=2*neighbour){
          if (result+neighbour < partialOutputs.size()){
               BlastImpl.assemblyPartitions(partialOutputs.get(result),partialOutputs.get(result+neighbor));
               lastMerge = partialOutputs.get(result);
          }
       }
      neighbor*=2;
    }
}
```

BLAST: Tree-based reduction

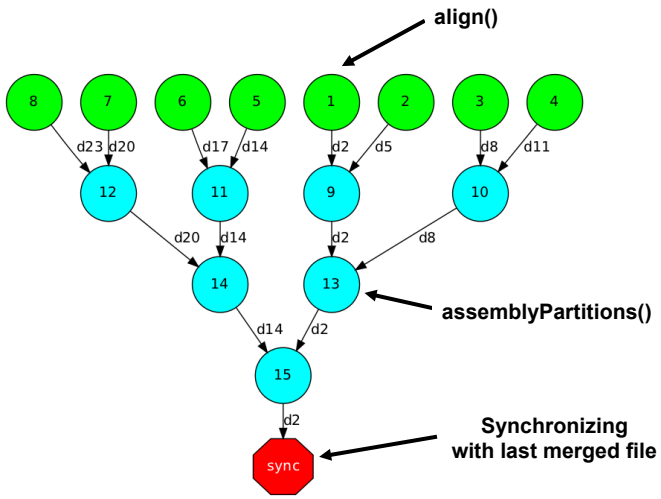
Creation the annotated interface for the selection of the remote tasks

```
public interface BlastItf {

    @Method(declaringClass = "blast.BlastImpl")
    @Constraints(processorCPUCount = 4, memoryPhysicalSize = 4.0f)
    void align(
        @Parameter(type = Type.STRING, direction = Direction.IN)
        String databasePath,

        @Parameter(type = Type.FILE, direction = Direction.IN)
        String partitionFile,

        ....

        @Parameter(type = Type.STRING, direction = Direction.IN)
        String commandArgs);

    @Method(declaringClass = "blast.BlastImpl")
    @Constraints(processorCPUCount = 2, memoryPhysicalSize = 2.0f)
    void assemblyPartitions(
        @Parameter(type = Type.FILE, direction = Direction.INOUT)
        String partialFileA,

        @Parameter(type = Type.FILE, direction = Direction.IN)
        String partialFileB);
}
```


BLAST: Tree-based execution

```
---------- Executing blast.Blast in IT mode total----------
…

BLAST Sequence Alignment Tool:

Parameters:
- Debug Enabled
- Blast binary: /home/user/workspace/blastAllOne/binary/blastall
- Number of expected fragments: 8
- Database Name with Path: /sharedDisk/Blast/databases/swissprot/swissprot
- Database Name: swissprot
- Input Sequences File: /sharedDisk/Blast/sequences/sargasso_test.fasta
- Temporary Directory: /tmp/
- Output File: /home/user/IT/blast.Blast/out.txt
- Command Line Arguments:

- The total number of sequences is: 20
- The total number of sequences of a fragment is: 3

- Splitting sequences among fragment files...
[  API]  -  Opening file /tmp/seqFileb0fa2b12-d0f6-42c1-b499-1e207e30ad84.sqf in mode APPEND
…

Aligning Sequences:

 - Number of fragments to assemble -> 8
 - Merging files -> /tmp/resFileb0fa2b12-d0f6-42c1-b499-1e207e30ad84.result.txt and /tmp/resFile815b4ff6-a077-422b-bc9b-9c6e10d8a417.result.txt
…
- Merging files -> /tmp/resFileb0fa2b12-d0f6-42c1-b499-1e207e30ad84.result.txt and /tmp/resFile81605bf8-b0f4-46bc-a521-9f289d219ef3.result.txt

Moving last merged file: /tmp/resFileb0fa2b12-d0f6-42c1-b499-1e207e30ad84.result.txt to /home/user/IT/blast.Blast/out.txt

[  API]  -  Opening file /home/user/IT/blast.Blast/out.txt in mode WRITE
- /sharedDisk/Blast/sequences/sargasso_test.fasta sequences aligned successfully in 193 seconds
…
------------------------------------------------------------
```

BLAST: Tree-based (Graph)



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HMMER: Hands-on

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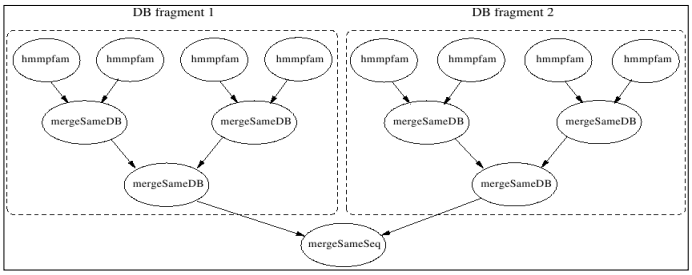
HMMER Hands On

Application: HMMER suite (hmmpfam)

- hmmpfam is part of the HMMER suite: set of tools for protein sequence analysis
  - Reads a sequences file and compares each sequence in it against a database of HMMs
  - HMM (Hidden Markov Model): statistical figure that represents a protein family
- Goal: create an hmmpfam efficient service
  - Starting point: sequential version of the hmmpfam tool
- With the COMPSs: hmmpfam becomes parallel
  - Phase 1: Split both input sequences and database
  - Phase 2: Process them in parallel (speed up execution)
  - Phase 3: Reduction of results



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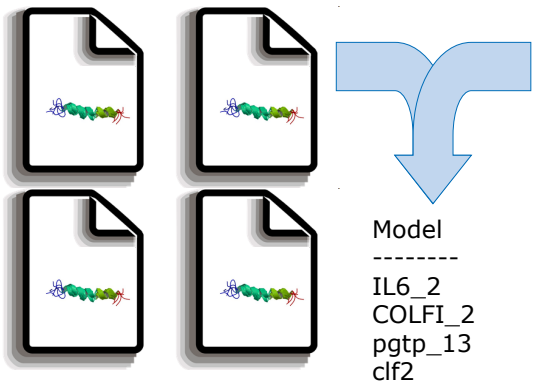
HMMER example

HMMER

Protein Database



Aminoacid Sequence

IQKKSGKWHTLTDLRA
VNAVIQPMGPLQPGLP
SPAMIPKDWPLIIIDLK
DCFFTIPLAEQDCEKFA
FTIPAINNKEPATRF

Model	Score	E-value	N
IL6_2	-78.5	0.13	1
COLFI_2	-164.5	0.35	1
pgtp_13	-36.3	0.48	1
clf2	-15.6	3.6	1
PKD_9	-24.0	5	1

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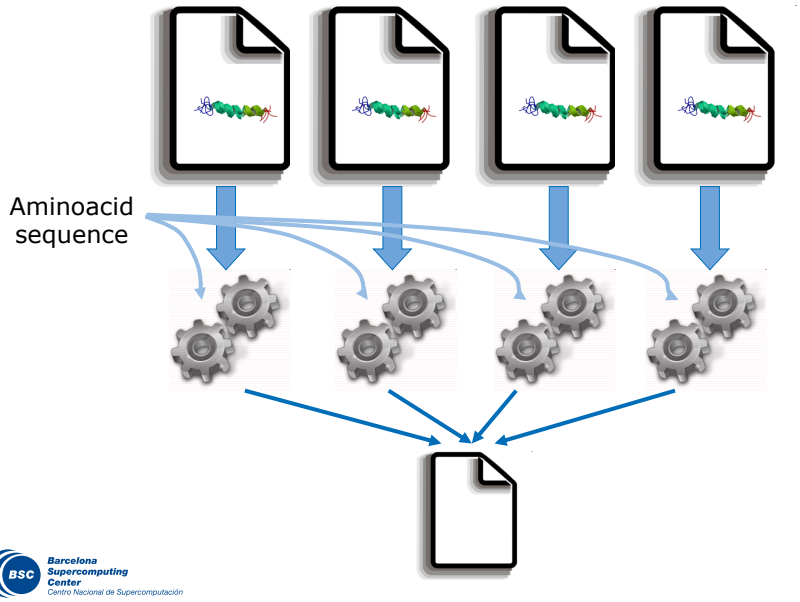
HMMER example



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HMMER example (code)

```
String[] outputs = new String[numDBFrags];

//Process
for (String dbFrag : dbFrags) {
     outputs[dbNum]= HMMPfamImpl.hmmpfam(sequence, dbFrag);
}


//Merge all DB outputs of the same DB fragment
int neighbour = 1;
while (neighbour < numDBFrags) {
  for (int db = 0; db < numDBFrags; db += 2 * neighbour) {
     if (db + neighbour < numDBFrags) {
        HMMPfamImpl.mergeSameDB(outputs[db], outputs[db + neighbour]);
     }
  }
  neighbour *= 2;
}
```

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HMMER example (code)

```
public interface HMMPfamItf {

    @Method(declaringClass = "worker.hmmerobj.HMMPfamImpl")
    @Constraints(storageElemSize = 1.5f)
    String hmmpfam(
            @Parameter(type = Type.FILE, direction = Direction.IN)
            String seqFile,
            @Parameter(type = Type.FILE, direction = Direction.IN)
            String dbFile,
            …
    );

    @Method(declaringClass = "worker.hmmerobj.HMMPfamImpl")
    void mergeSameDB(
            @Parameter(type = Type.OBJECT, direction = Direction.IN)
            String resultFile1,
            @Parameter(type = Type.OBJECT, direction = Direction.IN)
            String resultFile2
    );
    …
}
```

Implementation

Task constraints

Parameter metadata

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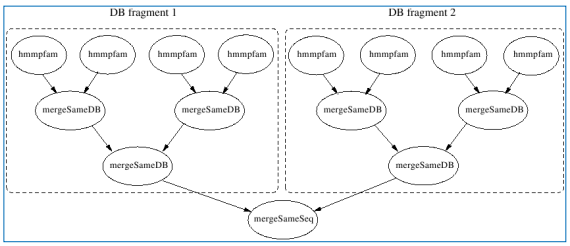
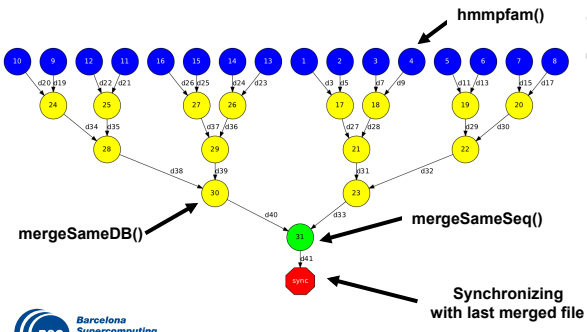
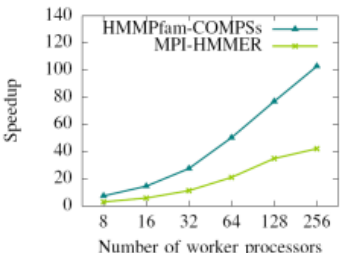
HMMER example (workflow)







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HMMER: Task Selection (work)

- Complete the hmmpfam & mergeSameSeq method interfaces.

HMMER: Configuration, compilation and execution

- Project.xml: /opt/COMPSs/Runtime/xml/projects/project.xml

```
<?xml version="1.0" encoding="UTF-8"?>
<Project>
    <!--Description for any physical node-->
    <Worker Name="localhost">
        <InstallDir>/opt/COMPSs/Runtime/scripts/</InstallDir>
        <WorkingDir>/tmp/</WorkingDir>
        <User>user</User>
        <LimitOfTasks>2</LimitOfTasks>
    </Worker>

</Project>
```

HMMER: Configuration, compilation and execution

- Configuration: /opt/COMPSs/Runtime/xml/resources/resources.xml

```
<?xml version="1.0" encoding="UTF-8"?>
<ResourceList>
    <!--Description for any physical node-->
    <Resource Name="localhost">
        <Capabilities>
            <Host>
                <TaskCount>0</TaskCount>
                <Queue>short</Queue>
                <Queue/>
            </Host>
            <Processor>
                <Architecture>AMD64</Architecture>
                <Speed>3.0</Speed>
                <CPUCount>2</CPUCount>
            </Processor>
            <OS>
                <OSType>Linux</OSType>
                <MaxProcessesPerUser>32</MaxProcess
            </OS>
            <StorageElement>
                <Size>30</Size>
            </StorageElement>
            …
            …
            <Memory>
                <PhysicalSize>2</PhysicalSize>
                <VirtualSize>8</VirtualSize>
            </Memory>
            <ApplicationSoftware>
                <Software>Java</Software>
            </ApplicationSoftware>
            <Service/>
            <VO/>
            <Cluster/>
            <FileSystem/>
            <NetworkAdaptor/>
            <JobPolicy/>
            <AccessControlPolicy/>
        </Capabilities>
        <Requirements/>
    </Resource>

<ResourceList>
```


HMMER: Configuration, compilation and execution

- Compilation (Eclipse IDE)
  - ***Package Explorer -> Project (hmmerobjblanks) -> Export… (Hands-on)***
  - ***Package Explorer -> Project (hmmerobj) -> Export… (Solution)***
- Usage
  - ***runcompss hmmerobj.HMMPfam <database> <sequences> <output> <params>***
- Execution
  - ***cp ~/workspace/hmmerobj/jar/hmmerobj.jar ~***
  - ***export CLASSPATH=$CLASSPATH:/home/user/hmmerobj.jar***
  - ***runcompss hmmerobj.HMMPfam /sharedDisk/Hmmer/smart.HMMs.bin / sharedDisk/Hmmer/256seq /home/user/out.txt 2 8 -A 222***

HMMER: Configuration, compilation and execution

```
user@bsccompss:~$ runcompss hmmerobj.HMMPfam /sharedDisk/Hmmer/smart.HMMs.bin /sharedDisk/Hmmer/
      256seq /home/user/out.txt 2 8 -A 222
-e
---------- Executing hmmerobj.HMMPfam in IT mode total----------

[ API] - Deploying the Integrated Toolkit
[ API] - Starting the Integrated Toolkit
[ API] - Initializing components
[ API] - Ready to process tasks
[ API] - Opening file /tmp/hmmer_frags/seqF0_1 in mode WRITE
[ API] - Opening file /tmp/hmmer_frags/seqF1_1 in mode WRITE
[ API] - Opening file /tmp/hmmer_frags/seqF2_1 in mode WRITE
[ API] - Opening file /tmp/hmmer_frags/seqF3_1 in mode WRITE
[ API] - Opening file /tmp/hmmer_frags/seqF4_1 in mode WRITE
[ API] - Opening file /tmp/hmmer_frags/seqF5_1 in mode WRITE
[ API] - Opening file /tmp/hmmer_frags/seqF6_1 in mode WRITE
[ API] - Opening file /tmp/hmmer_frags/seqF7_1 in mode WRITE
[ API] - Opening file /tmp/hmmer_frags/dbF0_1 in mode WRITE
[ API] - Opening file /tmp/hmmer_frags/dbF1 in mode WRITE
[ API] - Opening file /home/user/out.txt in mode WRITE
[ API] - No more tasks for app 1
[ API] - Stopping IT
[ API] - Cleaning
[ API] - Integrated Toolkit stopped

----------------------------------------------------------------
```

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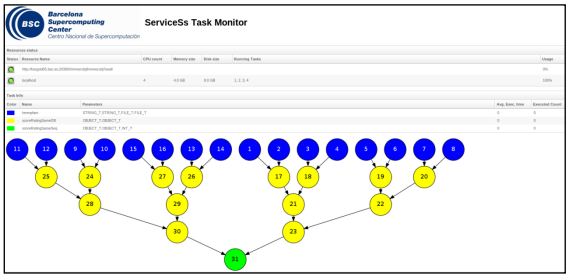
HMMER: Monitoring

- The runtime of COMPSs provides some information at execution time so the user can follow the progress of the application:
  - Real-time monitoring information (***http://localhost:8080/compss-monitor/***)
    - # tasks
    - Resources usage information
    - Execution time per task
    - Real-time execution graph
    - Etc.



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HMMER: Debugging

- COMPSs can be run in debug mode showing more information about the execution allowing to detect possible problems
  - Log level configurable at: **/opt/COMPSs/Runtime/log/it-log4j**
- The user can check the execution of its application by reading:
  - The output/errors of the main application (console stdout)
  - The output/error of a task # N
    - **~/IT/[*APP_NAME*]/jobs/jobN.[out|err]**
  - Messages from the runtime COMPSs
    - **~/it.log**
  - Task to resources allocation:
    - **~/resources.log**
- The user can verify the correct structure of the parallel application generating a complete post-mortem application graph
  - **gengraph $HOME/APP_NAME.dot**

Tracing: Overview

- **COMPSs can generate post-execution traces of the distributed execution of the application**
  - **Useful for performance analysis and diagnosis**
- **How it works?**
  - **Task execution and file transfers are application events**
  - **An XML file is created at workers to keep track of these events**
  - **At the end of the execution all the XML files are merged to get the final trace file**
  - **Instrumentation and Visualization tools from BSC are needed.**

Tracing: Instrumentation

- COMPSs uses Extrae tool to dynamically instrument the application
  - In a worker:
    - Extrae keeps track of the events in an intermediate file
  - In the master:
    - Extrae merges the intermediate files to get the final trace file
  - For more information about Extrae visit:
    - http://www.bsc.es/computer-sciences/extrae

Tracing: Instrumentation

```
---------- Executing hmmerobj.HMMPfam -------------------------

[ API] - Deploying the Integrated Toolkit
[ API] - Starting the Integrated Toolkit
[ API] - Initializing components

Welcome to Extrae 2.4.3rc4 (revision 311 based on framework/trunk/files/
extrae)
Extrae: Generating intermediate files for Paraver traces.
Extrae: Intermediate files will be stored in /home/user/IT/
hmmerobj.HMMPfam
Extrae: Tracing buffer can hold 500000 events
Extrae: Tracing mode is set to: Detail.
Extrae: Successfully initiated with 1 tasks

[ API] - Ready to process tasks

…
…
…
```

COMPSs runtime starts

Extrae starts before the user application execution

Extrae keeps tracing events in background

Tracing: Instrumentation

```
[ API] - No more tasks for app 1
[ API] - Stopping IT
[ API] - Cleaning
Extrae: Application has ended. Tracing has been terminated.
...

merger: Output trace format is: Paraver
merger: Extrae 2.4.3rc4 (revision 311 based on framework/trunk/files/extrae)
…

[ API] - Integrated Toolkit stopped
…

mpi2prv: Selected output trace format is Paraver

mpi2prv: Parsing intermediate files

mpi2prv: Generating tracefile (intermediate buffers of 1342156 events)

mpi2prv: Congratulations! hmmerobj.HMMPfam_compss_trace_1392736225.prv has been generated.

------------------------------------------------------------
```

The application finishes and the tracing process ends

The merge process starts

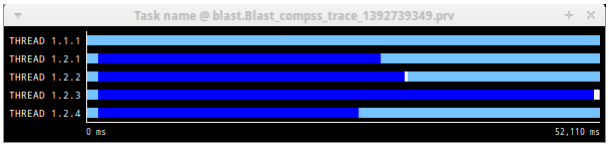
COMPSs runtime ends

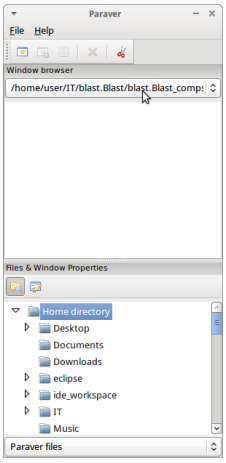
Intermediate trace files are processed

The final trace file is generated

Tracing: Visualization

- **Paraver is the BSC tool for trace visualization**
  - **Trace events are encoding in Paraver (.prv) format by Extrae**
  - **Paraver is a powerful tool for trace visualization.**
  - **An experimented user could obtain many different views of the trace events.**
  - **For more information about Paraver visit:**
    - **http://www.bsc.es/computer-sciences/performance-tools/paraver**



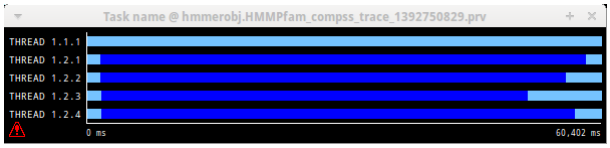


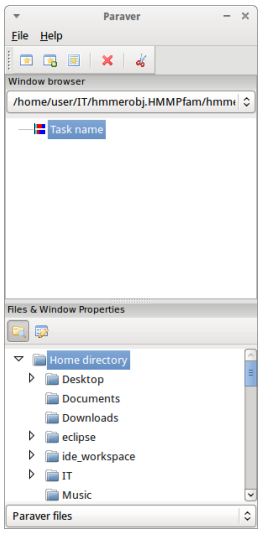
Tracing: Hands-on

- Compilation (Eclipse IDE)
  - ***Package Explorer -> Project (hmmerobj) -> Export...***
- Execution
  - ***cp ~/workspace/hmmerobj/jar/hmmerobj.jar ~***
  - ***export CLASSPATH=$CLASSPATH:/home/user/hmmerobj.jar***
  - ***runcompssext --app=hmmerobj.HMMPfam --tracing=true --cline_args="/sharedDisk/Hmmer/smart.HMMs.bin /sharedDisk/Hmmer/256seq /home/user/out.txt 2 8 -A 222"***
  - ***wxparaver /home/user/IT/hmmerobj.HMMPfam/*.prv***

Tracing: Hands-on

- COMPSs provides a configuration file to automatically obtain the view of the trace
  - ***File / Load Configuration...***
  - ***/opt/COMPSs/paraver/cfgs/tasks.cfg***

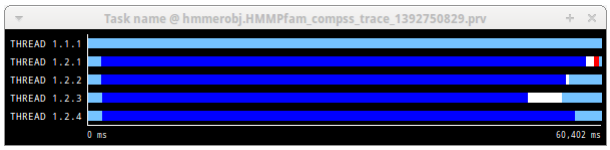




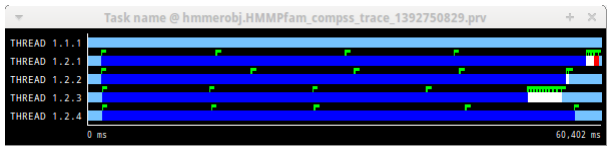
- Some small adjustments must be done in order to view the trace correctly

Tracing: Hands-on

- Fit window
  - ***Right click on the trace window***
  - ***Fit Semantic Scale / Fit Both***



- View Event Flags
  - ***Right click on the trace window***
  - ***View / Event Flags***



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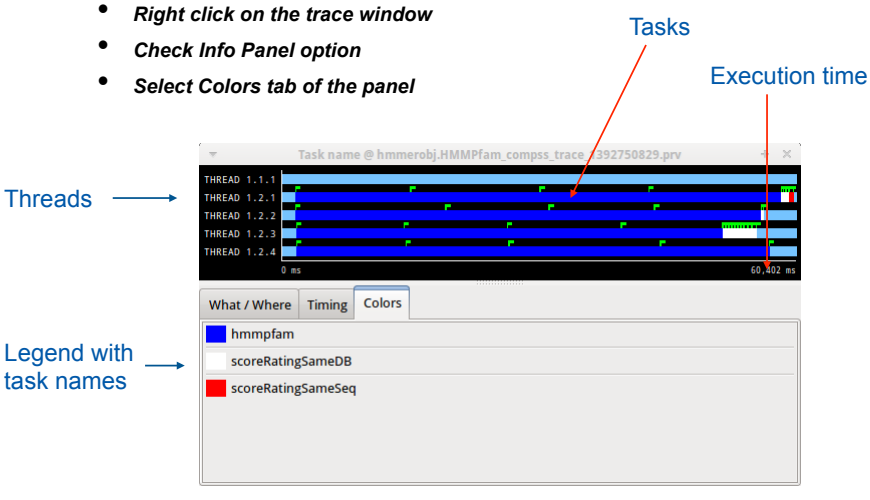
Tracing: Hands-on

- Show Info Panel
  - ***Right click on the trace window***
  - ***Check Info Panel option***
  - ***Select Colors tab of the panel***



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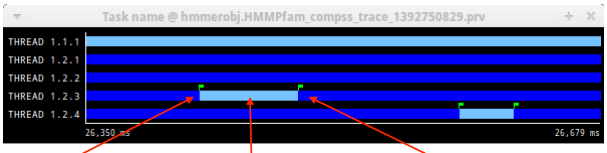
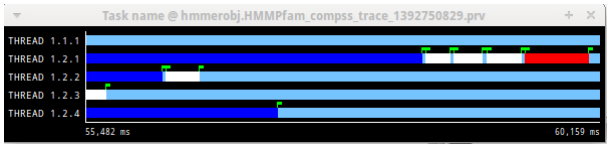
Tracing: Hands-on

- Zoom to see details
  - ***Select a region in the trace window to see in detail***
  - ***And repeat the process til the needed zoom level***
  - ***The undo zoom option is in the right click panel***



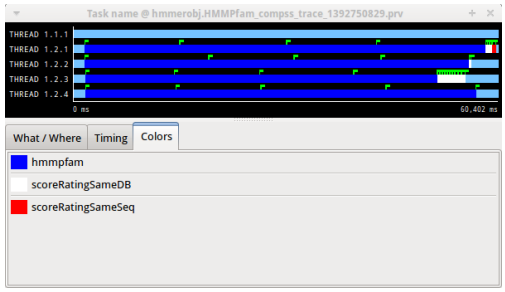
Previous task ends Processor is idle New task starts



Tracing: Hands-on

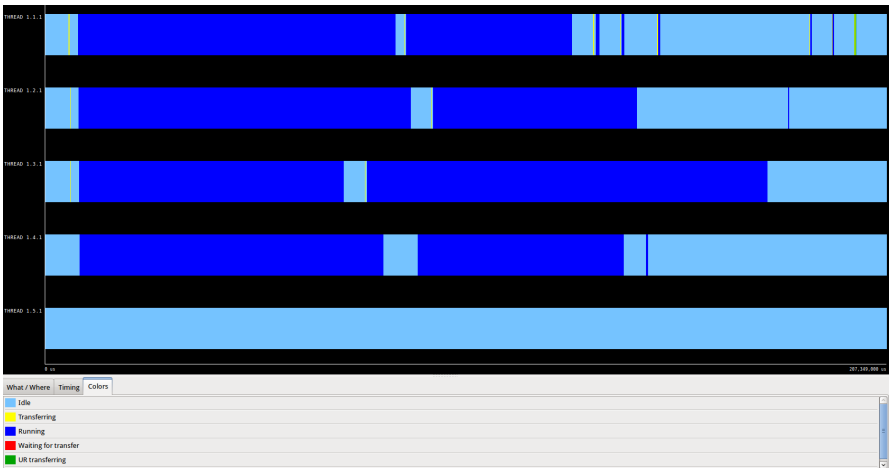
Summarizing:

- Lines in the trace:
  - One line for the master
  - N lines for the workers



- Meaning of the colours:
  - Light blue: idle
  - Other colors: task running, see the color legend
- Flags (events):
  - Start / end of task

Tracing: BLAST, Tree-based



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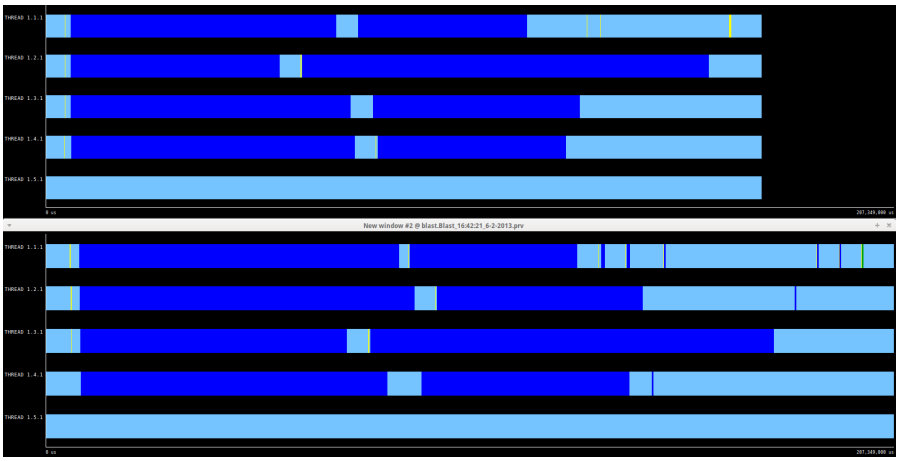
Tracing: BLAST, All-to-One vs Tree-based (Local)



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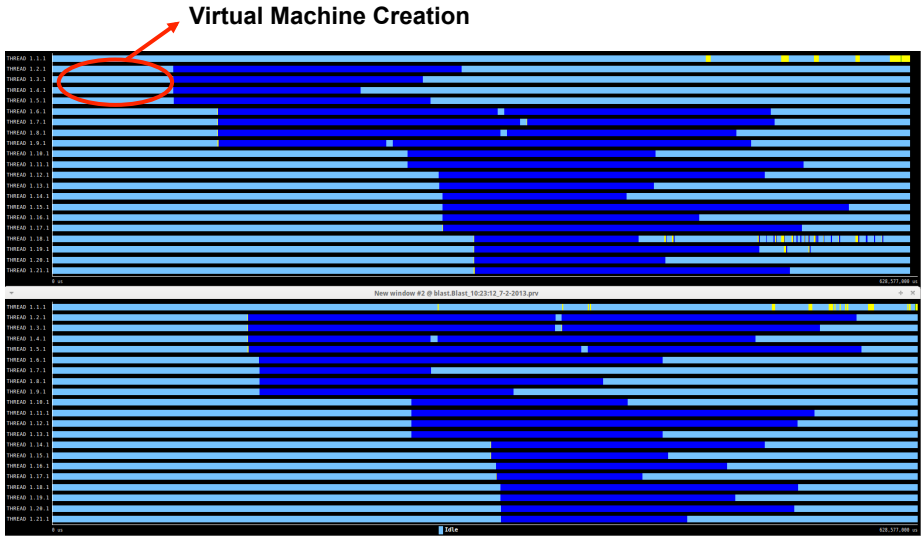
Tracing: BLAST, All-to-One vs Tree-based (Cloud)



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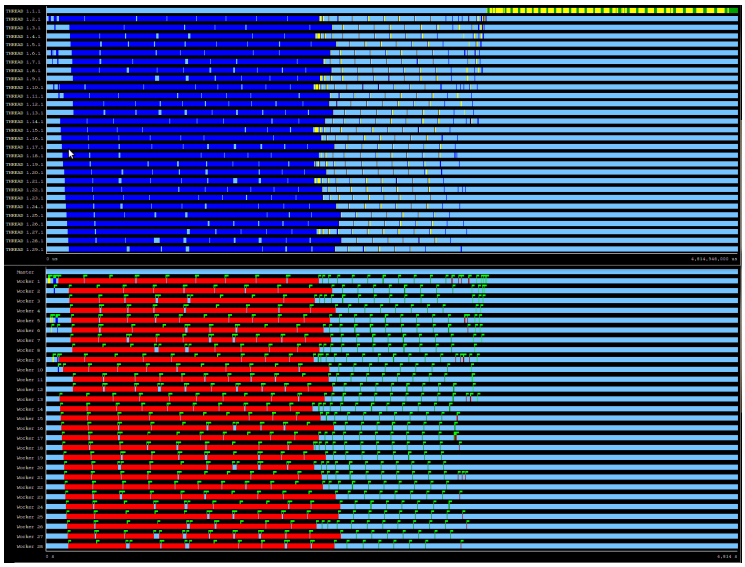
Tracing: Other examples



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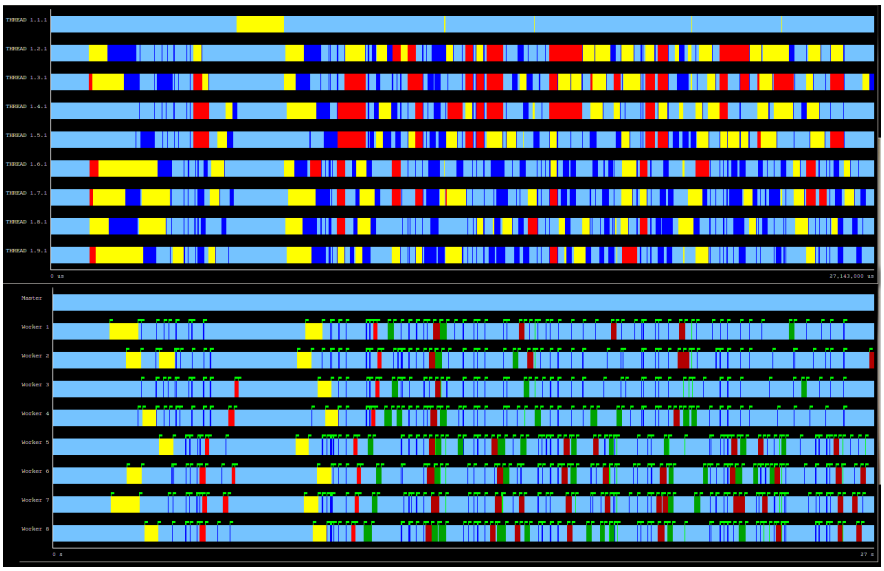
Tracing: Other examples



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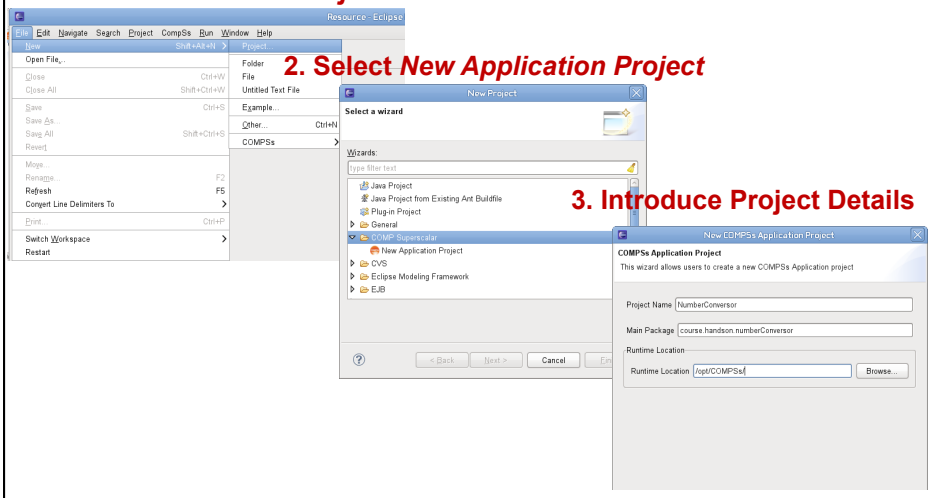
Integrated Development Environment
HANDS-ON

IDE Hands On – Create a COMPSs Project

1. Menu *File->New -> Project…*



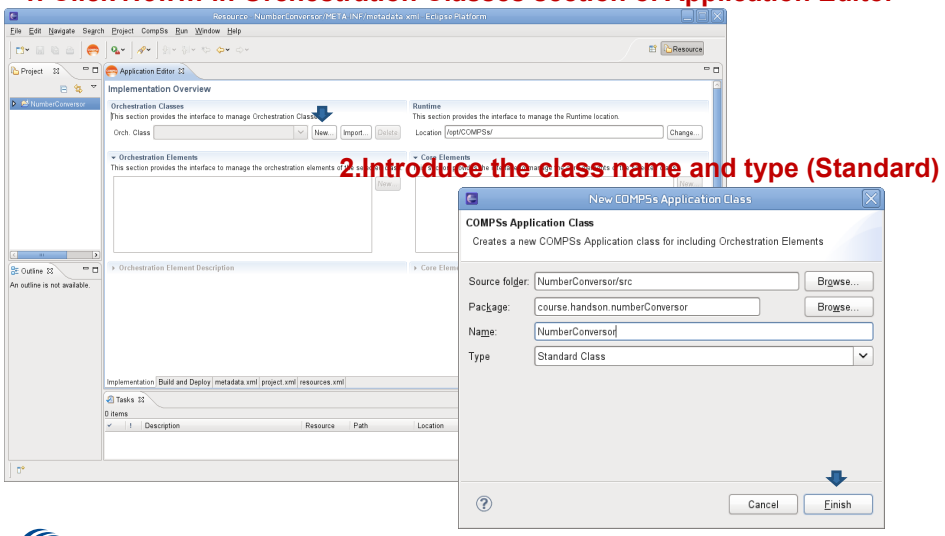
2. Select *New Application Project*

3. Introduce Project Details

(Also available *CompSs –>Implementation->Create Application Project*)

IDE Hands On – Create an Orchestration Class

1. Click *New…* in Orchestration Classes section of Application Editor

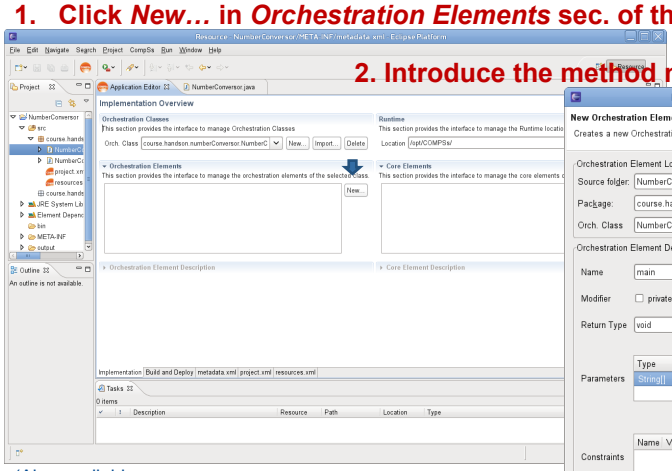


2.Introduce the class name and type (Standard)

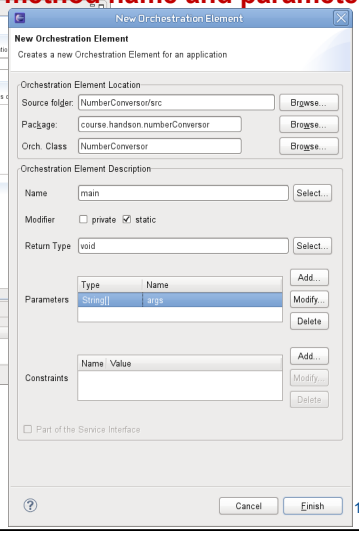
(Also available *CompSs –>Implementation->Create Orchestration Class*)

IDE Hands On – Create an Orchestration Element

1. **Click *New…* in *Orchestration Elements* sec. of the *Application Editor***



2. Introduce the method name and parameters



(Also available:

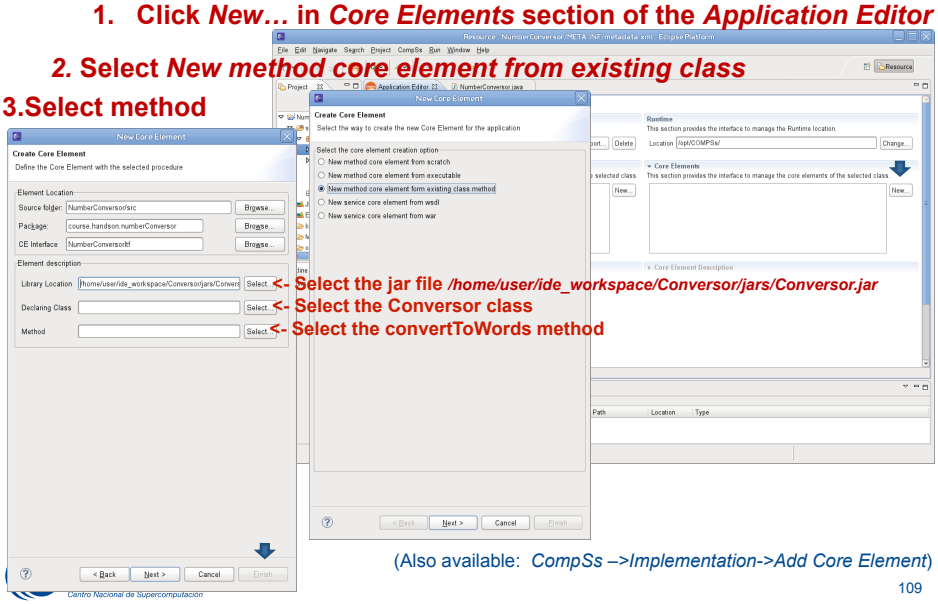
CompSs –>Implementation->Add Orchestration Element)

IDE Hands On – Add an Core Element from JAR

1. **Click *New…* in *Core Elements* section of the *Application Editor***

2. Select *New method core element from existing class*

3.Select method



<- Select the jar file */home/user/ide_workspace/Conversor/jars/Conversor.jar*

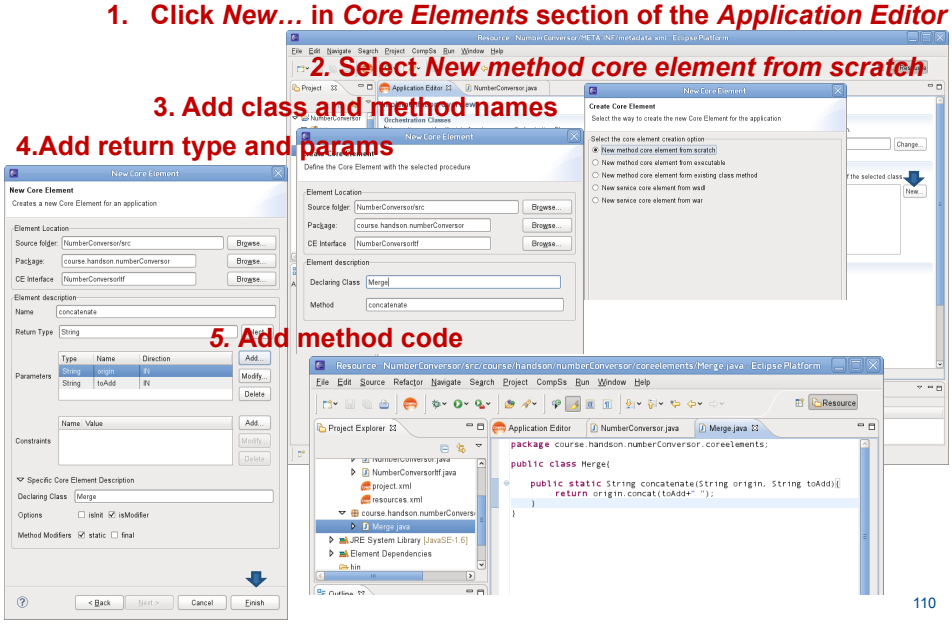
<- Select the Conversor class

<- Select the convertToWords method

(Also available: *CompSs –>Implementation->Add Core Element*)

IDE Hands On – Add an Core Element from scratch

1. Click *New…* in *Core Elements* section of the *Application Editor*
2. Select *New method core element from scratch*
3. Add class and method names
4. Add return type and params
5. Add method code



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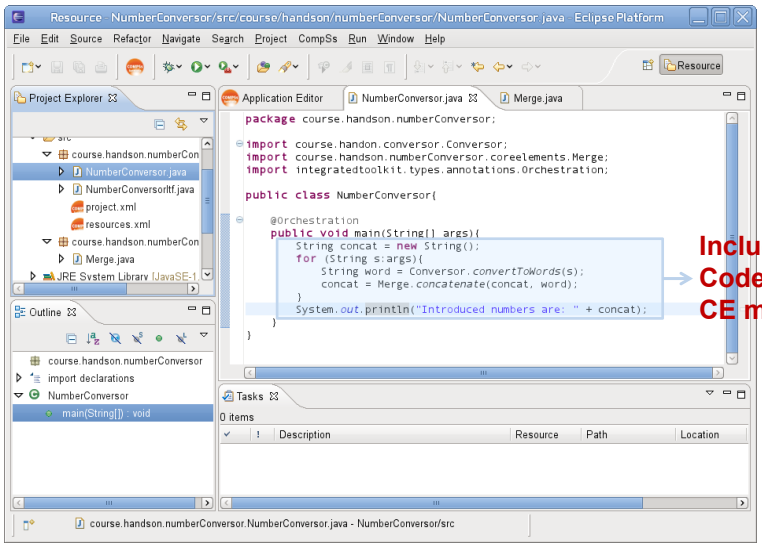
IDE Hands On – Introduce the OE code



Include the OE Code to call the CE methods

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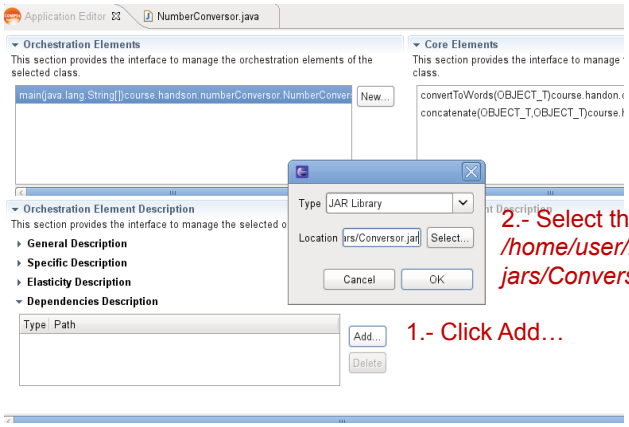
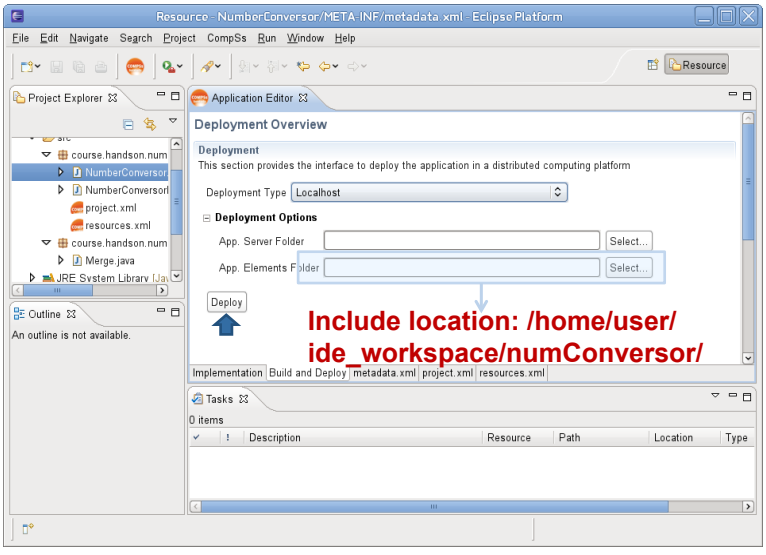
IDE Hands On – Add conversor dependency to OE



2.- Select the jar library: */home/user/ide_workspace/Conversor/jars/Conversor.jar*

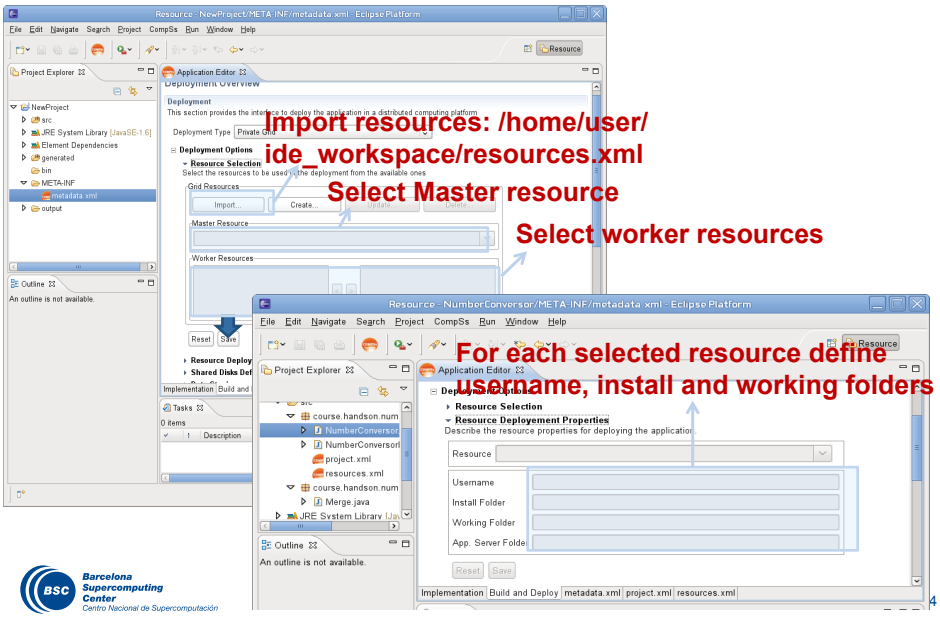
1.- Click Add…

IDE Hands On – Deploy Locally



Include location: /home/user/ide_workspace/numConversor/

IDE Hands On – Deploy Grid



Final Notes

- Sequential programming approach
- Parallelization at task level
- Transparent data management and remote execution
- Can operate on different infrastructures:
  - Cluster
  - Grid
  - Cloud (Public/Private)
    - PaaS
    - IaaS
  - Web services

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Final notes

- **Project page: http://www.bsc.es/compss**
- **Direct downloads page:**

  http://www.bsc.es/computer-sciences/grid-computing/comp-superscalar/download

  - *Sample applications & development virtual appliances*
  - *Tutorials*
  - *Red-Hat & Debian based installation packages*
  - *…*

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Thank you!

For further information please contact

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