# WP1-3: Hw and Sw technologies for HPC and their use in SyeC applications

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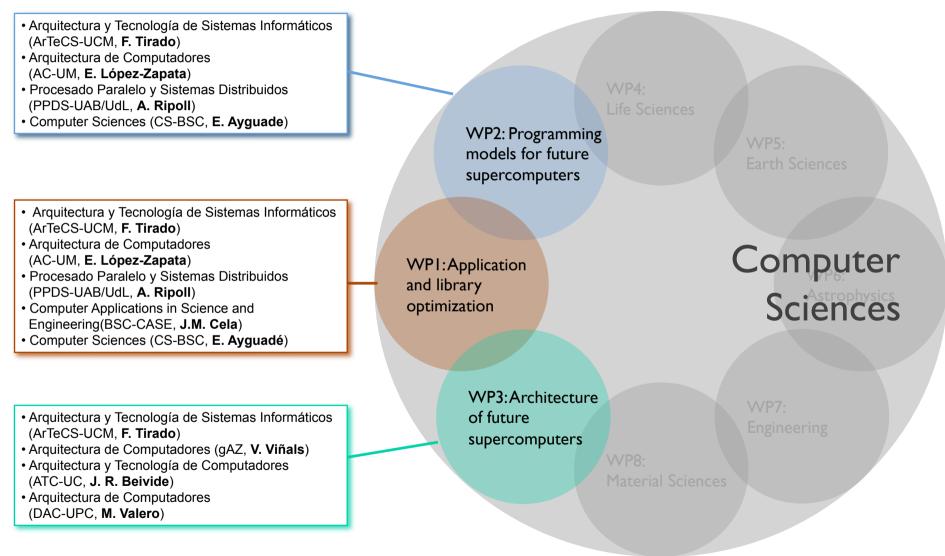
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#### WP1-3: project structure and participating groups (7)





#### WP1: activities and outcomes

#### From sequential optimization to parallelization

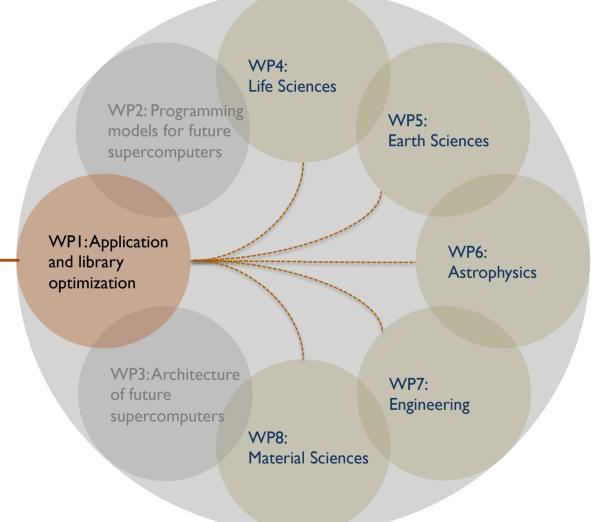
- Algorithmic improvements, load balancing
- Hybrid MPI/OpenMP parallelization
- Optimization for GPU and multicore

#### **Collaborations:**

- Gromacs: UB + CS-BSC
- T-coffee: CRG + PPDS-UAB/UdL
- GEM: CRG + gAZ
- NCAR-WACCM: UCM + ArTeCS-UCM
- Atmospheric: ES-BSC + CASE-BSC
- SPEV: UV + AC-UM
- MRGENESIS: UV + CS-BSC
- RATPENAT: UV + CASE-BSC
- pDEVA: UAM + CASE-BSC
- BLKTRI: CIEMAT + ArTeCS-UCM
- SIESTA: CSIC-ICN2 + CASE-BSC

#### Other applications

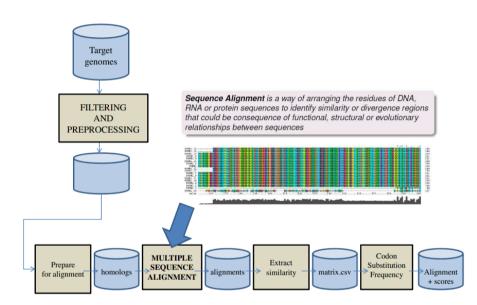
- Histogram, k-means, biomedicine, 3D Fast Wavelet, Generalized Hough Transform: AC-UM
- Image segmentation (gAZ)





#### Multiple sequence alignment and T-coffee

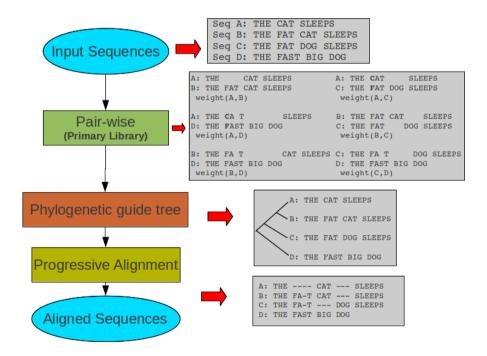
( MSA common in many biocomputing workflows



( Limitations: memory requirements and execution time (quadratric with number of sequences and length)

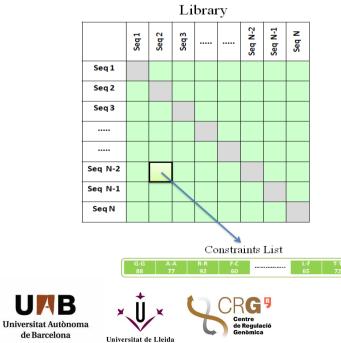


#### ( T-coffee: reference MSA algorithm of Life Science community

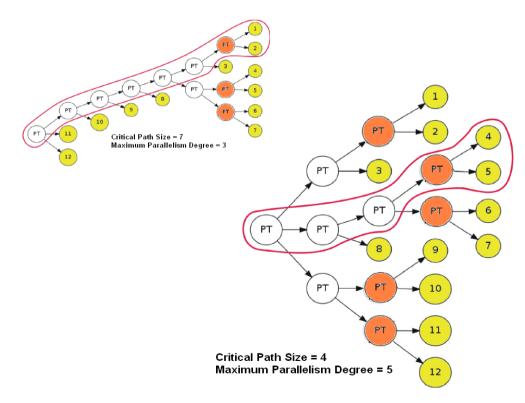


#### T-coffee: two main optimizations

- ( Improved primary library generation, discarding redundant/less representative information
  - Reduce memory requirements
  - Better execution time and increased scalability
  - Keep alignment quality

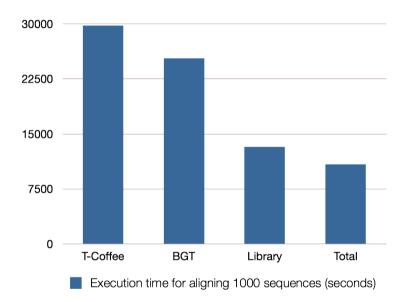


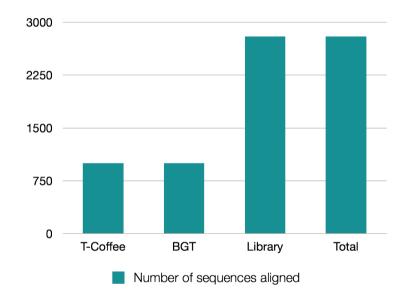
- ( Balanced Guide Tree (BGT)
  - Better parallelism degree and load balancing in the progressive alignment phase
  - Keep alignment quality



#### T-coffee: performance improvement

#### ( Execution time and number of sequences aligned





#### ( Biological accuracy of alignment

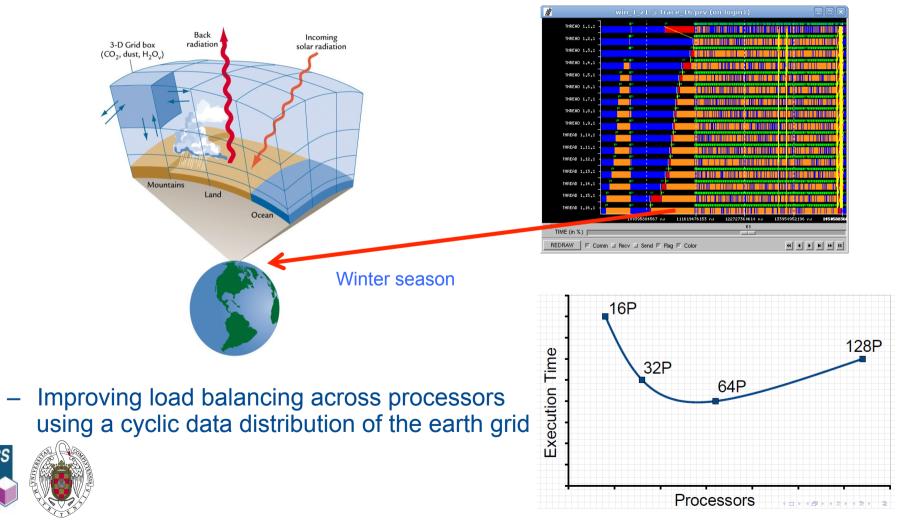
Similarity	standard T-Coffee quality	optimized T-Coffee quality
0-15	0,421	0,379
15-35	0,721	0,695
25-35	0,876	0,865
35-100	0,951	0,956
average	0,709	0,687

PREFAB MSA benchmark results



# Whole-Atmosphere Community Climate Model (WACAMM)

- ( Main bottleneck in radiation code
  - Computational cost depends on seasonal cycles: temporal variations and spatial variations



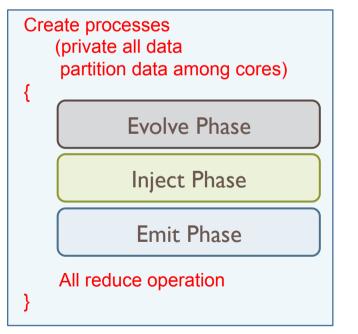
### SPectral EVolution simulation (SPEV)

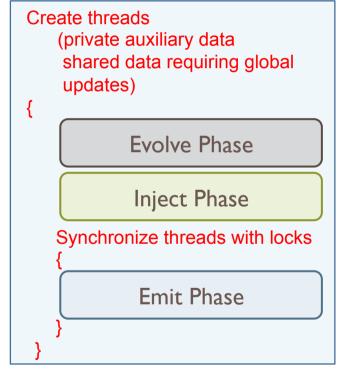
- ( Radio emissions from extragalactic ( Shared data approach jets → observed on Earth
- ( Data-sharing on multi-socket and multicore architectures
- ( Private data approach

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- ( Hybrid approach
  - Private data across chips (sockets)
  - Shared data inside chips

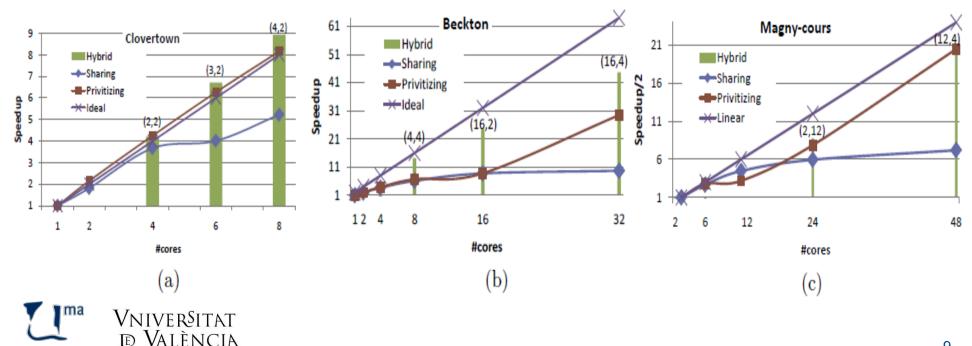
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### SPEV: performance results

( Platforms

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- 2-socket 4-core Intel Clovertown
- 4-socket 8-core Intel Beckton
- 4-socket 12-core AMD Magny-Cours
- ( Programming models
  - OpenMP (threads and synchronization)
  - MPI (processes and reduction)

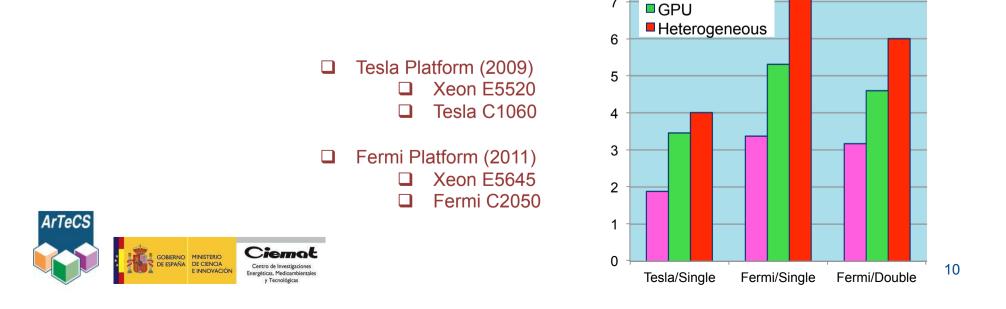


### **BLKTRI solver from FISH-PACK library**

- ( Numerical simulation of incompressible fluid flows
  - The most time consuming part of incompressible unsteady Navier Stokes solver is related to the solution of a pressure Poisson equation
- Target architecture: heterogeneous multicore/GPU
  - Coarse-grain loop parallelism (OpenMP) in tri-diagonal problems for multicore
  - Fine-grain parallelism (CUDA) for GPU, with major algorithm changes
- ( Substantial speedup over the standard BLKTRI solver using a combination of the two strategies (overlapping) 8

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#### WP2: activities and outcomes

Definition and implementation of programming models:

- COMPSs for the specification of workflows, support for web services
- OmpSs for asynchronous/dataflow programming in OpenMP, heterogeneity support (CUDA, OpenCL)
- **Chapel** (from Cray): implementation of the communication layer handling the transference of array regions in clusters
- Extensions to Intel **TBB** for the specification of wavefront computations
- Source-to-source compilation (PPCG), from sequential to parallel code for modern GPU (collaboration with IMEC/INRIA)

Scheduling techniques for homogeneous and heterogeneous multicore architectures (accelerators, asymmetric multicores, ...)

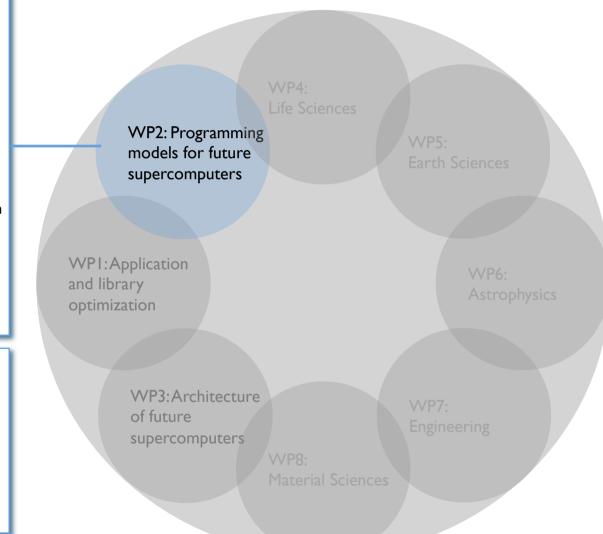
New data distribution policies and scheduling policies for **Hadoop/MapReduce**:

#### **Optimization of application workflows**

COMPSs and MapReduce

#### Collaborations

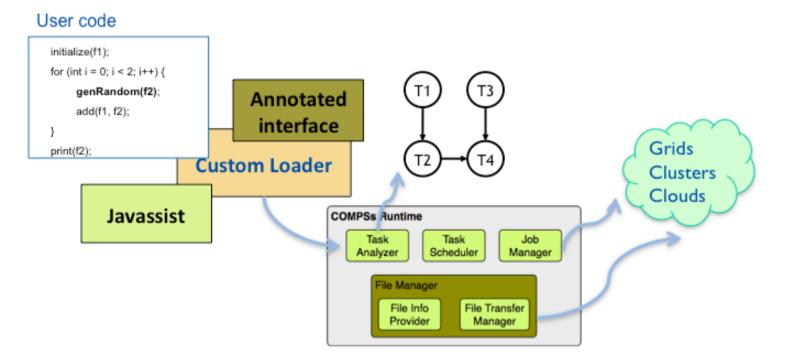
- Milky Way origin (GASS, GOG): UB + CS-BSC
- Protein dynamics, gene detection: IRB-UB
  + CS-BSC
- LNC-RNA: CRG + PPDS-UAB/UdL
- Ocean-atmosphere circulation model: ES-BSC + CS-BSC





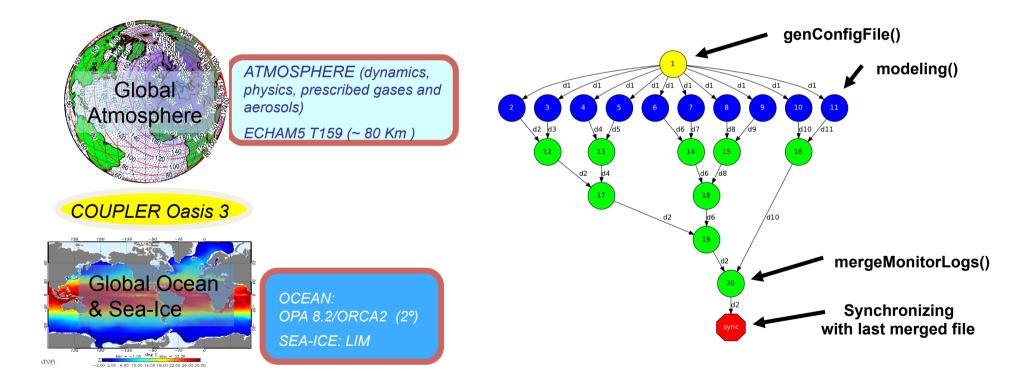
## COMPSs programming model

- ( Easy programming, execution in any platform (cluster, Grid and Cloud)
- ( Task based programming model (automatic generation of data-dependent task graph)
  - Coarse grain tasks: methods and web services
  - Whole application can be exposed as a new web-service
- (Cloud interoperability: commercial solutions (e.g. Azure and Amazon), open source (e.g. OpenNebula, EMOTIVE Cloud, OpenStack)



### COMPSs: coupled models in Earth Sciences models

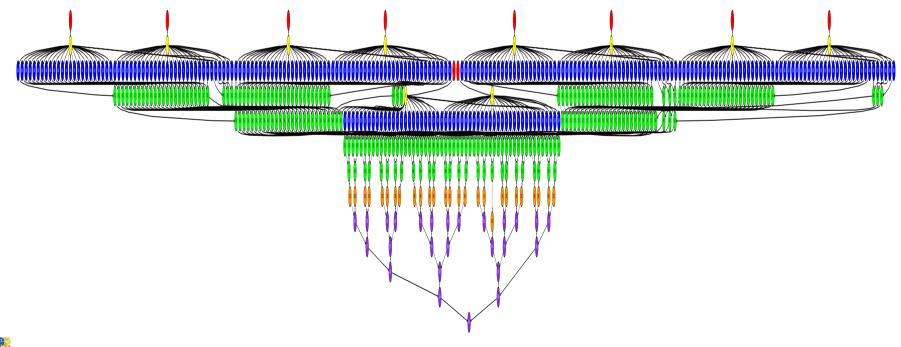
- ( Global coupled ocean-atmosphere general circulation model
  - Communication between atmospheric and ocean models done through the CMCC parallel version of OASIS3 coupler





### COMPSs: protein dynamics in Life Sciences workflow

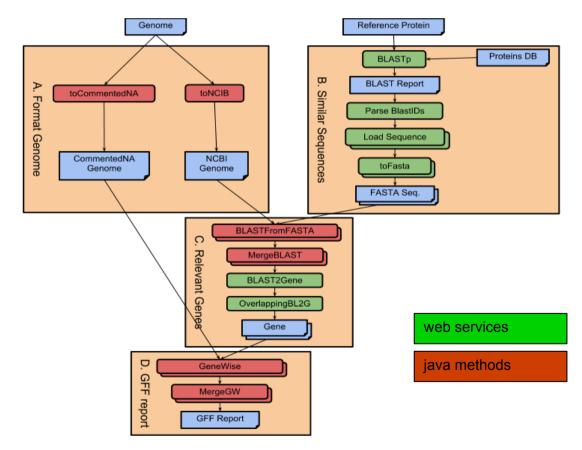
- ( DISCRETE is a package devised to simulate the dynamics of proteins using the Discrete Molecular Dynamics (DMD) method
  - From a set of protein structures, perform a series of simulations to optimize 3 parameters: FDVW, FSOLV, EPS
  - For each structure,  $N_{fdvw} \cdot M_{fsolv} \cdot L_{eps}$  simulations are done





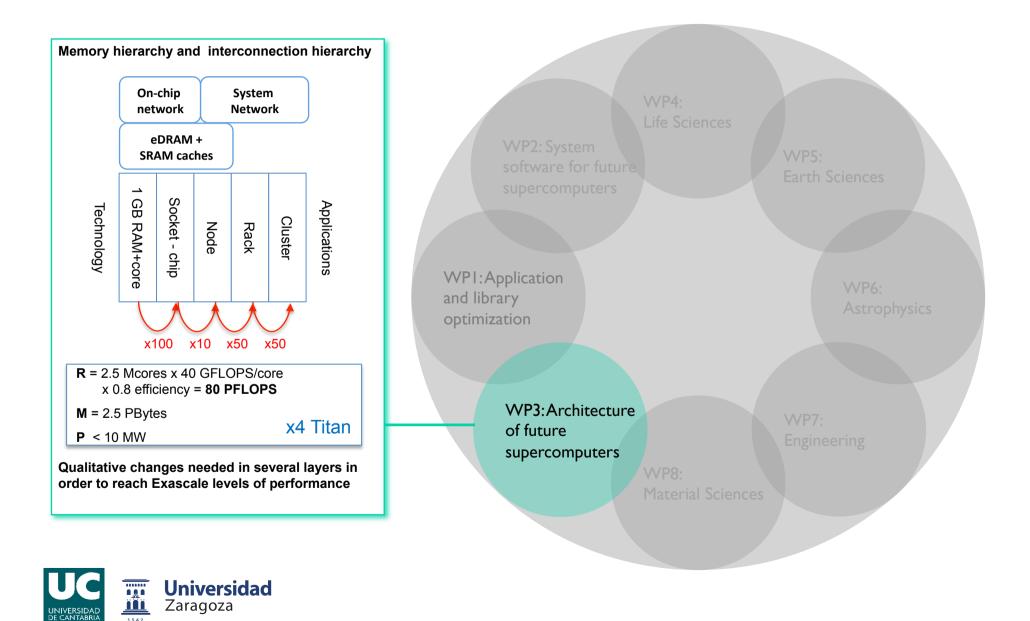
#### COMPSs: gene detection in Life Sciences workflow

- Automatic identification of genes which causes a disease
- Combine web services with computations
- Implemented as a new composite service

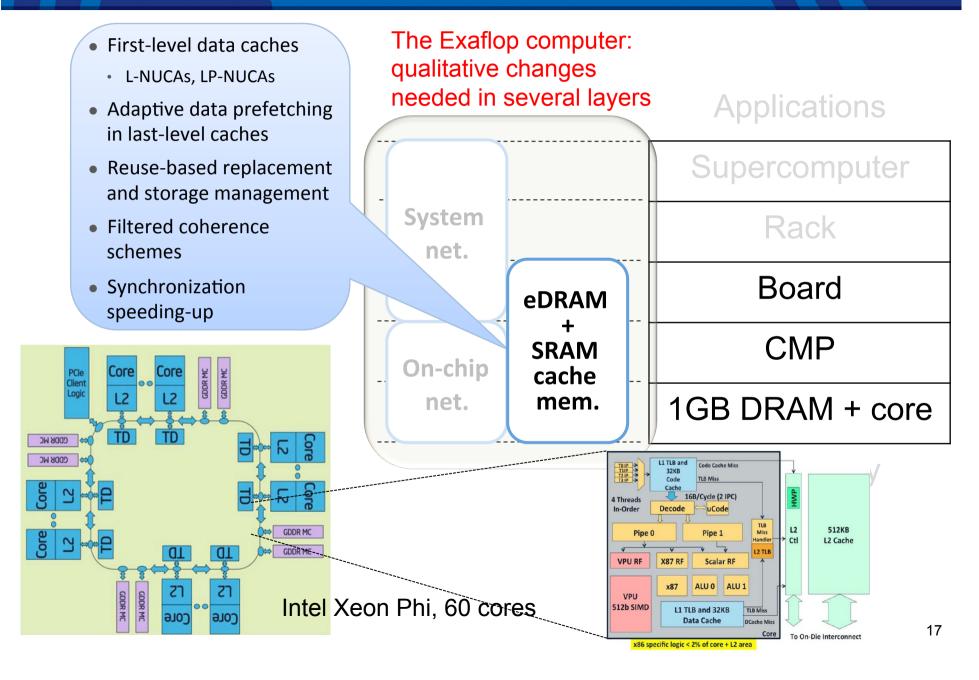




#### WP3: activities and motivation



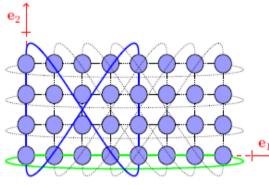
#### Cache hierarchy

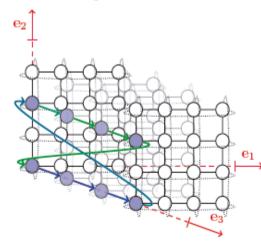


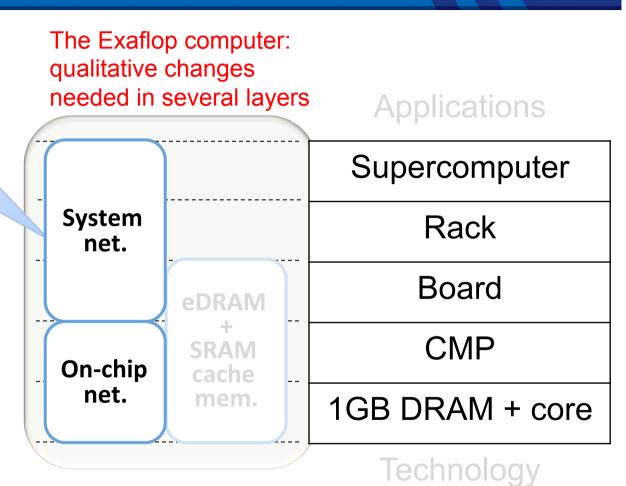
#### **Network foundations**

# Theory: applying topological results to network design

- low-degree networks
- cayley graphs
- gaussian integers
- error-correcting codes

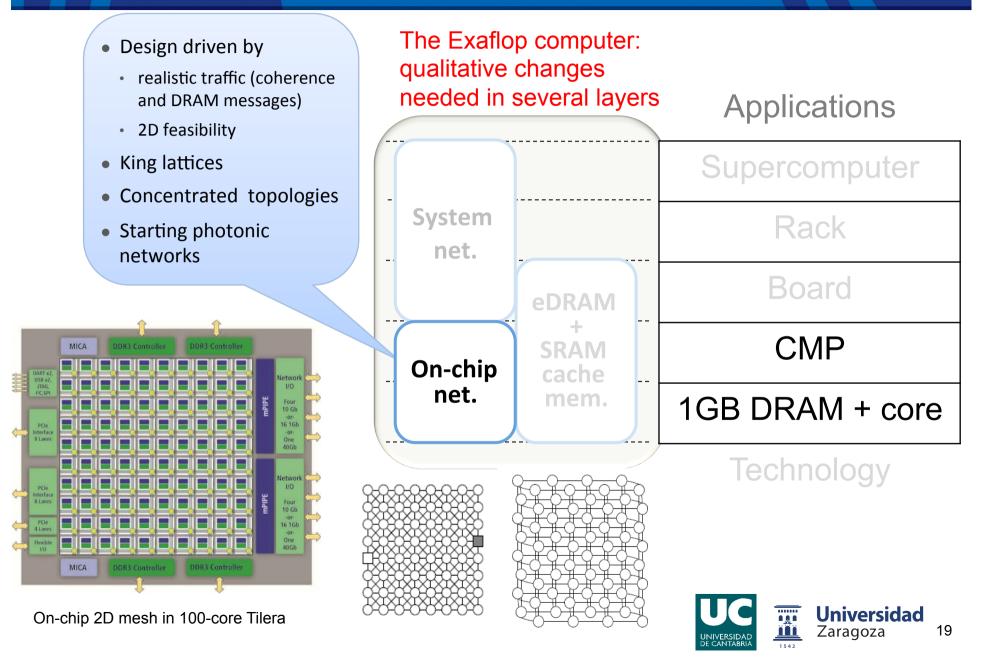




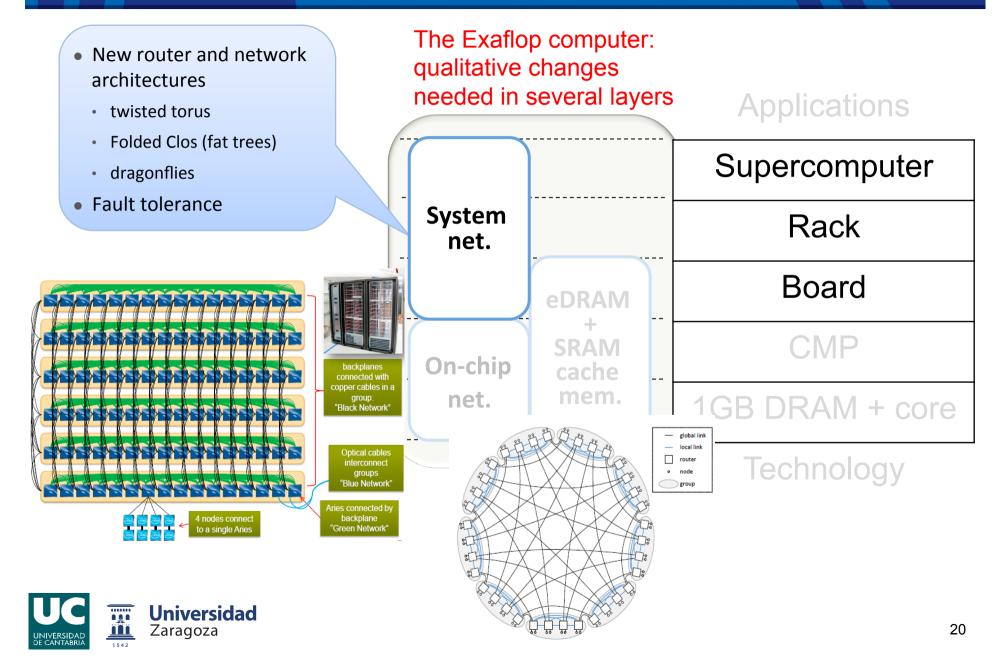




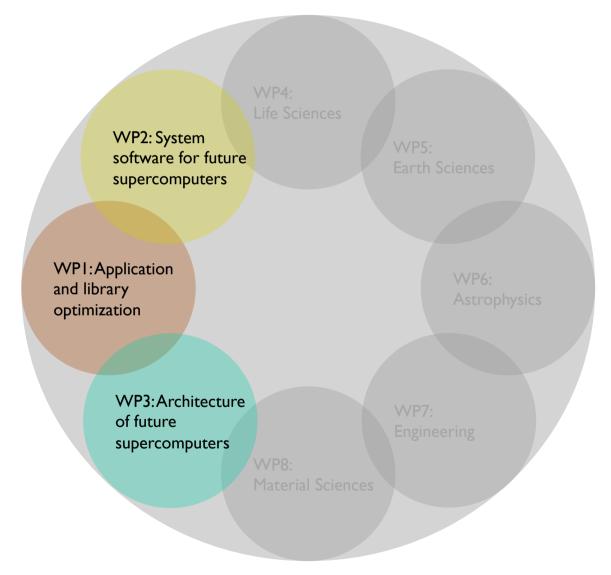
#### **Networks on Chip**



#### System networks



#### WP3: conclusions





# Conclusions

- Collaboration between Computer Science and Application groups resulted in important improvements
  - Needed to adapt to novel hw technologies, sometimes difficult to predict their impact in actual applications
- ( Impact in the evolution of programming models
  - Novel programming models: OmpSs
  - Evolution of current ones: TBB (Thread Building Blocks)
  - Efficient implementation of current ones: data movement for Chapel, automatic compilation for CUDA
  - We missed the possibility of using some of them in applications from other groups in the Consolider project (only COMPSs for workflows)
- ( Architectural design of future supercomputers
  - Intra-chip and intra-node interconnect and memory hierarchy
  - Inter-node interconnection network
  - We also missed the opportunity of driving architectural simulations using real applications from the Consolider project