



Universitatea “Ștefan cel Mare” Suceava

Facultatea de Inginerie Electrică și Știința Calculatoarelor

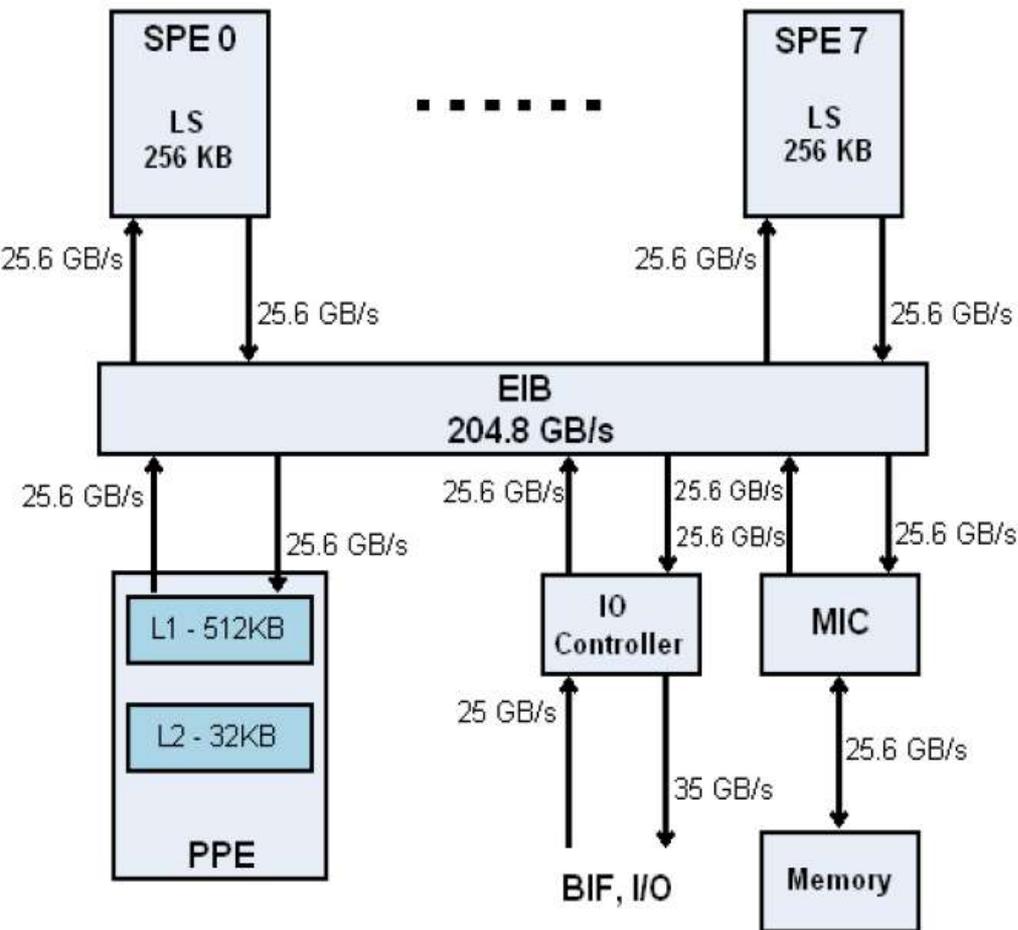
HIGH-PERFORMANCE COMPUTING ON A SUPERCOMPUTER BASED ON NEW-GENERATION PROCESSORS

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1. INTRODUCTION



- ▶ Cell Broadband Engine Architecture was developed by IBM for Cell B.E. processor from PlayStation 3 game console.
- ▶ This architecture was optimized for math intensive computing.

2. USV-ROADRUNNER SUPERCOMPUTER(1 / 3)

- ▶ In December 2009, the HPC lab was completed with an IBM Roadrunner-type cluster, with a peak performance of 9.98 TFlops.
- ▶ Based on Cell BE processors
- ▶ Energy efficient (~ 337 MFlops/watt)
- ▶ Architecture optimized for intensive arithmetic computations



2. USV-ROADRUNNER SUPERCOMPUTER(2/3)

- ▶ 48 processing QS22 dual-CPU blades with Cell BE processors:
 - 2 IBM PowerXCell 8i CPUs (3.2 GHz)
 - 8 GB RAM per blade
 - 8 GB modular SSD
 - Gigabit Ethernet interface for administration
 - Infiniband 4xDDR interface for data transfer
 - GPFS access on the storage system for concurrent parallel access of files



3. DETERMINATION OF A MODEL FOR A TYPE-3 CLASSIFIER (1 / 2)

- ▶ The CLASS algorithm is part of the supervised classification algorithms.
- ▶ This means that the algorithm needs a training set, which is used for classification of patterns from the input set.

3. DETERMINATION OF A MODEL FOR A TYPE-3 CLASSIFIER (2 / 2)

- ▶ **Algorithm 1. Sequential algorithm.**
- ▶ *) *Read the input data from a file.*
- ▶ *) *Allocate memory for the $A(m \times (p+1))$ matrix.*
- ▶ *) *Allocate memory for the $\text{miu}(m)$ vector.*
- ▶ *) *Initialize with 0 the elements of matrix A.*
- ▶ **for $i=1, n$ do**
- ▶ **for $j=1, p$ do**
- ▶ $A[\text{CLASA}[i]][j] += \text{ANTREN}[i][j]$
- ▶ **end for**
- ▶ $\text{miu}[\text{CLASA}[i]] += F[i]$
- ▶ **end for**
- ▶ **for $l=1, M$ do**
- ▶ **for $j=1, p$ do**
- ▶ $A[l][j] /= \text{miu}[l];$
- ▶ $A[l][p+1] += A[l][j] * A[l][j] * (-1/2)$
- ▶ **end for**
- ▶ **end for**
- ▶ **end.**

4. PARALLELIZATION OF THE ALGORITHM FOR USV-ROADRUNNER SUPERCOMPUTER

Algorithm 2. Parallelization of the algorithm for USV-RoadRunner supercomputer.

Read the input data from a file.

**) Allocate memory for the $A(m \times (p+1))$ matrix.*

**) Allocate memory for the $miu(m)$ vector.*

**) Initialize with 0 the elements of matrix A.*

**) Distribute the set of training patterns to the MPI process ($nLocal$ – the number of patterns assigned to the current MPI process).*

for $i=1, nLocal$ do

for $j=1, p$ do

$A[CLASA[i]][j] += ANTREN[i][j]$

end for

$miu[CLASA[i]] += F[i]$

end for

**) Use $MPI_AllReduce()$ function in order to update the matrix A and vector miu on all MPI processes from the system.*

for $l=1, M$ do

for $j=1, p$ do

$A[l][j] /= miu[l];$

$A[l][p+1] += A[l][j] * A[l][j] * (-1/2)$

end for

end for

**) Use $MPI_AllReduce()$ function in order to update the matrix A on all MPI processes from the system.*

end.

5. THE ACTIVATION OF THE SPE CORES

Algorithm 3. The activation of the SPE cores.

- *) Read the input data from a file.*
- *) Allocate memory for the $A(m \times (p+1))$ matrix.*
- *) Allocate memory for the $miu(m)$ vector.*
- *) Initialize with 0 the elements of matrix A .*
- *) Distribute the set of training patterns to the MPI process ($nLocal$ – the number of patterns assigned to the current MPI process).*

Distribute the local set of training patterns to the SPE cores

- *) Send a signal to the SPE cores in order to begin the compute of matrix A based on assigned patterns.*
- *) Wait for a mailbox from each SPE core, which signals the end of assigned jobs.*

```
for  $l=1,M$  do  
    for  $j=1,p$  do  
         $A[l][j] /= miu[l];$   
         $A[l][p+1] += A[l][j]*A[l][j]*(-1/2)$   
    end for  
end for  
*) Use MPI_AllReduce() function in order to update the matrix  $A$  and vector  $miu$  on all MPI processes from the system.  
end.
```

6. EXPERIMENTAL RESULTS (1 / 4)

- ▶ In order to test the performance of this algorithm, we have chosen the p53 mutants dataset.
- ▶ This dataset represents the models of the mutant p53 protein, models that can be used to predict the transcriptional activity of p53 protein (cancer diagnosis).
- ▶ This data set has 16,722 patterns, each pattern with 5409 features.

6. EXPERIMENTAL RESULTS (2/4)

- ▶ In first step, the performances were measured by execution of the algorithm on one processor PowerXCell 8i, without use of the SPE cores.
- ▶ Next, the number of PowerXCell 8i processor was doubled until the maximum number (96) (see Fig. 2).

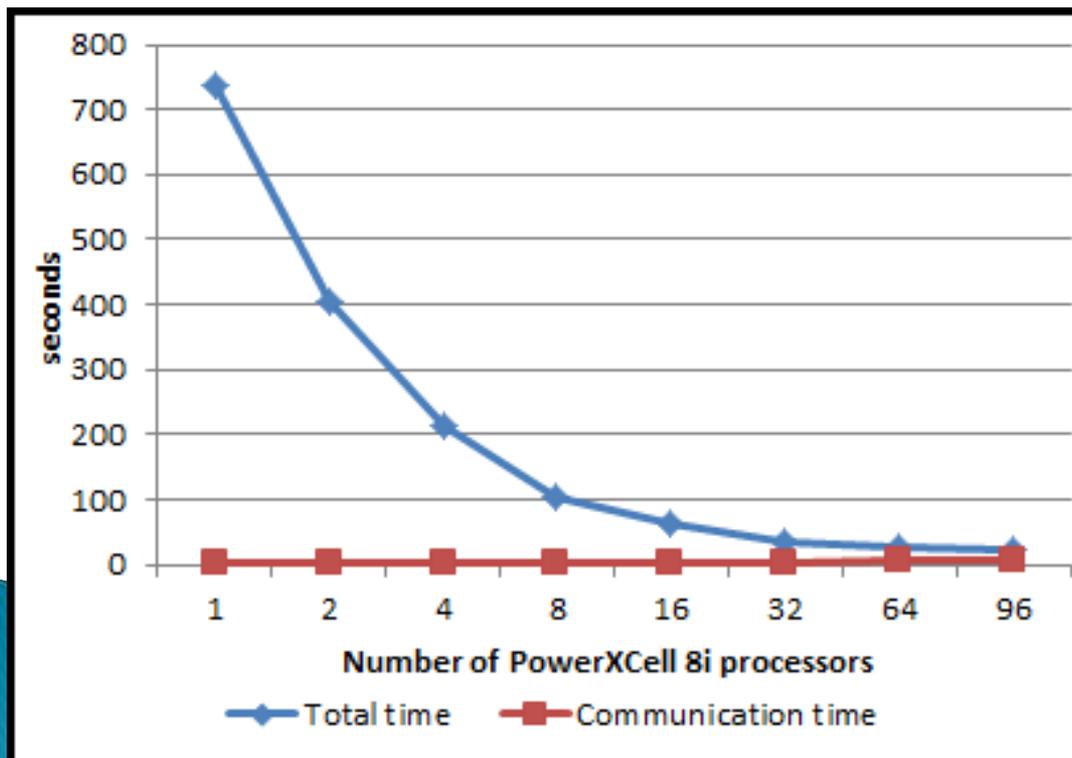


Figure 2. The performances obtained without use of SPE cores.

6. EXPERIMENTAL RESULTS (3/4)

- ▶ Fig 3 presents the total execution time and communication time for the case in which all 8 SPE cores on each PowerXCell 8i processor are used.

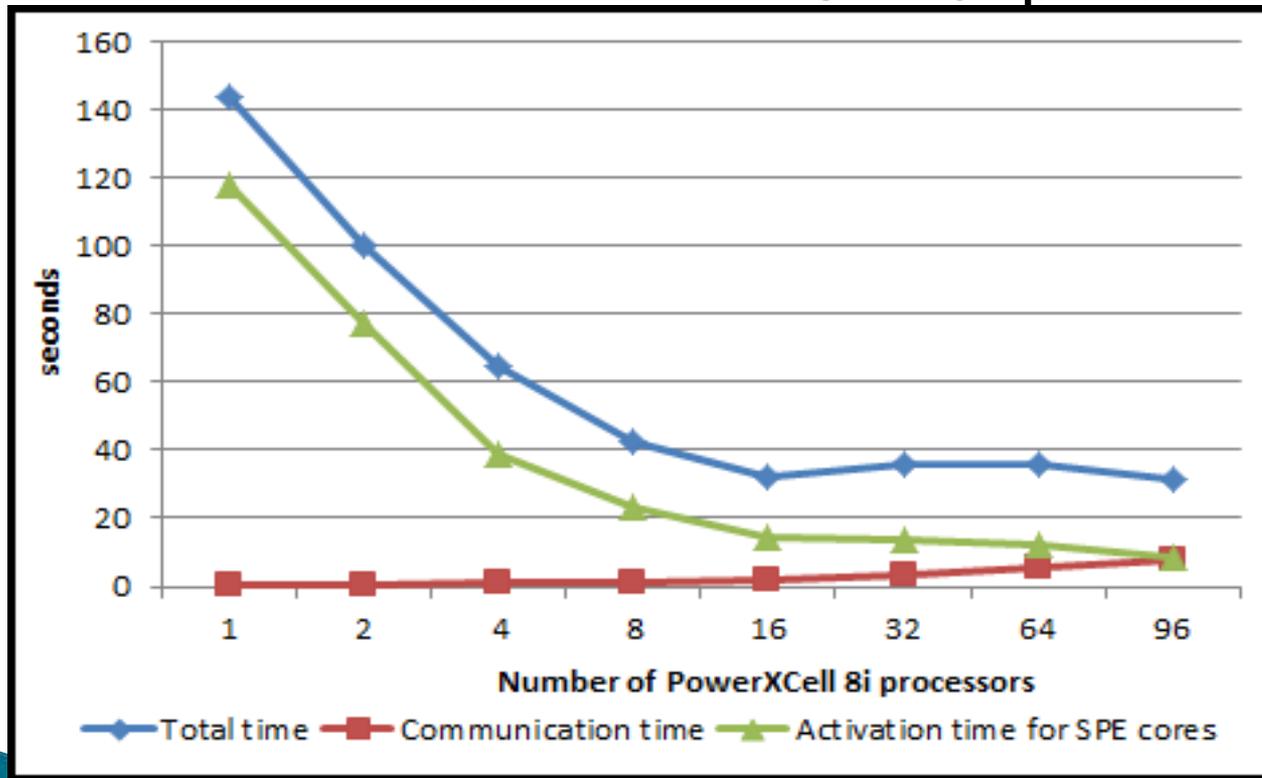


Figure 3. The performances obtained with use of SPE cores.

6. EXPERIMENTAL RESULTS (4/4)

- ▶ Fig 4 presents the speeds-up obtained by increasing the number of PowerXCell 8i processors with and without use of the SPE cores.
- ▶ Speeds-ups are reported to the algorithm execution on a PowerXCell 8i processor without use of the SPE cores.

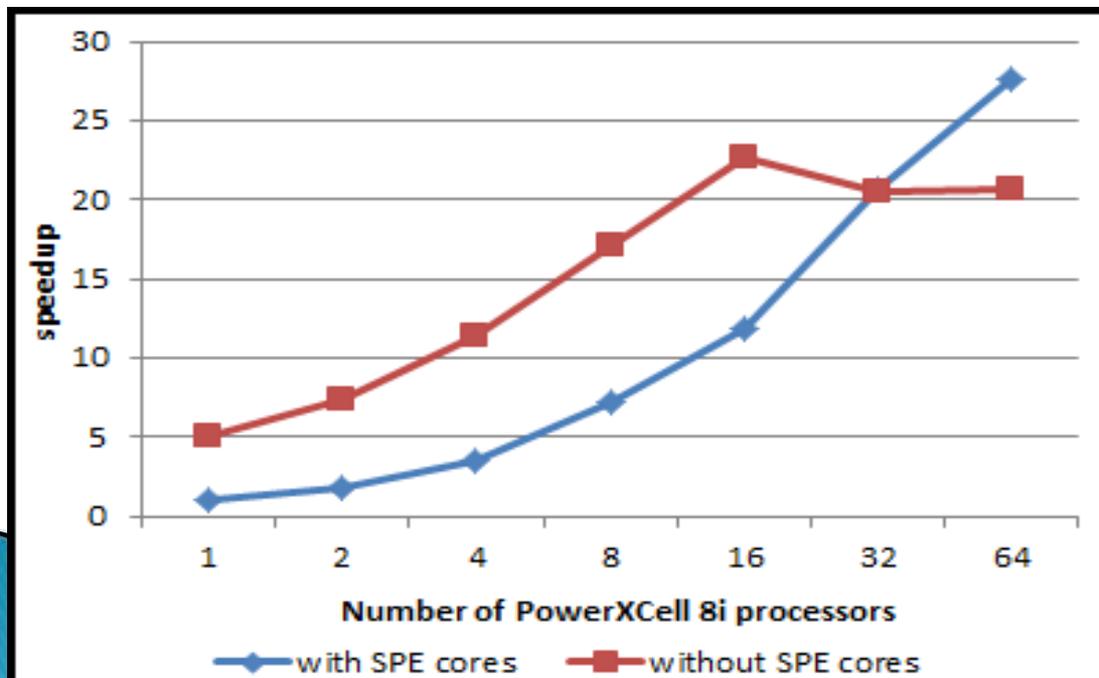


Figure 4. Graphical representation of speeds-up obtained.

7. CONCLUSIONS

- ▶ In this paper, we present the methods used in order to design, program, and analyze a multilevel parallel variant so as to determine a model for a type-3 classifier.
- ▶ A comparison was performed between the sequential variant and parallelized variant was carried out on a single level, as well as for the parallelized variant on two levels.

7. CONCLUSIONS

- ▶ The performances achieved by algorithm execution were analyzed for the USV-RoadRunner supercomputer. For the moment, we obtain a speed-up of 28.
- ▶ Algorithm can be further optimized by using the SIMD facilities held of SPE cores. Using these facilities, the speedup can increase with an order size (x10).



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Thank you!