GPCALMA: a Grid Approach to Mammographic Screening

S. Bagnasco\textsuperscript{a}\footnote{INFN\textsuperscript{I}.N.F.N., Sezione di Torino, Italy.,}, U. Bottigli\textsuperscript{b}\footnote{SAS\textsuperscript{I}}\textsuperscript{I}.N.F.N. di Cagliari, Italy, P. Cerello\textsuperscript{c}\footnote{INFN\textsuperscript{I}.N.F.N. di Pisa, Italy, M. E. Fantacci\textsuperscript{d}\footnote{PISA\textsuperscript{I}}, E. Lopez Torres\textsuperscript{e}\footnote{CUBA\textsuperscript{I}}, Habana, Cuba, G. L. Masala\textsuperscript{g}\footnote{SAS}, P. Oliva\textsuperscript{h}\footnote{SAS}, A. Retico\textsuperscript{i}\footnote{PISA\textsuperscript{I}}, S. Stumbo\textsuperscript{j}\footnote{SAS}}\textsuperscript{d}\footnote{PISA\textsuperscript{I},}, P. Delogu\textsuperscript{d}\footnote{PISA\textsuperscript{I}}, M. E. Fantacci\textsuperscript{d}\footnote{PISA\textsuperscript{I}}, E. Lopez Torres\textsuperscript{e}\footnote{CUBA\textsuperscript{I}}, CEADEN, Habana, Cuba,

The next generation of High Energy Physics experiments requires a GRID approach to a distributed computing system and the associated data management: the key concept is the Virtual Organisation (VO), a group of geographically distributed users with a common goal and the will to share their resources. A similar approach is being applied to a group of Hospitals which joined the GPCALMA project (Grid Platform for Computer Assisted Library for MAmmography), which will allow common screening programs for early diagnosis of breast and, in the future, lung cancer. HEP techniques come into play in writing the application code, which makes use of neural networks for the image analysis and shows performances similar to radiologists in the diagnosis. GRID technologies will allow remote image analysis and interactive online diagnosis, with a relevant reduction of the delays presently associated to screening programs.

1. Introduction

A reduction of breast cancer mortality in asymptomatic women is possible in case of early diagnosis \cite{1}, which is available thanks to screening programs, a periodical mammographic examination performed for 49-69 years old women. The GPCALMA Collaboration aims at the development of tools that would help in the early diagnosis of breast cancer: Computer Assisted Detection (CAD) would significantly improve the prospects for mammographic screening, by quickly providing reliable information to the radiologists.

A dedicated software to search for massive lesions and microcalcification clusters was developed recently (1998-2001): its best results in the search for massive lesions (microcalcification clusters) are 94% (92%) for sensitivity and 95% (92%) for specificity. Meanwhile, in view of the huge distributed computing effort required by the CERN/LHC collaborations, several GRID projects were started. It was soon understood that the application of GRID technologies to a database of mammographic images would facilitate a large-scale screening program, providing transparent and real time access to the full data set.

The data collection in a mammographic screening program will intrinsically create a distributed database, involving several sites with different functionality: data collection sites and diagnostic sites, i.e. access points from where radiologists would be able to query/analyze the whole distributed database. The scale is pretty similar to that of LHC projects: taking Italy as an example, a full mammographic screening program would act on a target sample of about 6.8 million women, thus generating 3.4 millions mammographic exams/year. With an average size of 60 MB/exam, the amount of raw data would be
in the order of 200 $\text{TB}/\text{year}$: a screening program on the European scale would be a data source comparable to one of the LHC experiments.

GPCALMA was proposed in 2001, with the purpose of developing a GRID application; based on technologies similar to those adopted by the CERN/ALICE Collaboration. In each hospital, digital images will be stored in the local database and registered to a common service (Data Catalogue). Data describing the mammograms, also known as metadata, will also be stored in the Data Catalogue and could be used to define an input sample for any kind of epidemiology study. The algorithm for the image analysis will be sent to the remote site where images are stored, rather than moving them to the radiologist’s sites. A preliminary selection of cancer candidates will be quickly performed and only mammograms with cancer probabilities higher than a given threshold would be transferred to the diagnostic sites and interactively analysed by one or more radiologists.

2. The GPCALMA CAD Station

The hardware requirements for the GPCALMA CAD Station are very simple: a PC with SCSI bus connected to a planar scanner and to a high resolution monitor. The station can process mammograms directly acquired by the scanner and/or images from file and allows human and/or automatic analysis of the digital mammogram. The software configuration for the use in local mode requires the installation of ROOT\cite{2} and GPCALMA, which can be downloaded either in the form of source code from the respective CVS servers. The functionality is usually accessed through a Graphic User Interface, or, for developers, the ROOT interactive shell. The Graphic User Interface (fig. 1) allows the acquisition of new data, as well as the analysis of existing ones. Three main menus drive the creation of (access to) datasets at the patient and the image level and the execution of CADe algorithms. The images are displayed according to the standard format required by radiologists: for each image, it is possible to insert or modify diagnosis and annotations, manually select the Regions of Interest (ROI) corresponding to the radiologists geometrical indication. An interactive procedure allows zooming, either continously or on a selected region, windowing, gray levels and contrast selection, image inversion, luminosity tuning. The

![Figure 1. The GPCALMA Graphic User Interface. Three menus, corresponding to the Patient, the Images and the CAD diagnosis levels, drive it. On the left, the CAD results for microcalcifications and masses are shown in red squares and green circles, together with the radiologist’s diagnosis (blue circle). On the right, the image colours are inverted. The widget drives the update of patient and image related metadata. Human analysis produces a diagnosis of the breast lesions in terms of kind, localization on the image, average dimensions and, if present, histological type. The automatic procedure finds the ROI's on the image with a probability of containing an interesting area larger than a pre-selected threshold value.](image)

3. Grid Approach

The amount of data generated by a national or european screening program is so large that they can’t be managed by a single computing centre. In addition, data are generated according to an instrinsically distributed pattern: any hospital participating to the program will collect a small fraction of the data. Still, that amount would
be large enough to saturate the available network connections.

The availability of the whole database to a radiologist, regardless of the data distribution, would provide several advantages:

- the CAD algorithms could be trained on a much larger data sample, with an improvement on their performance, in terms of both sensitivity and specificity.

- the CAD algorithms could be used as real-time selectors of images with high breast cancer probability (see fig. 2): radiologists would be able to prioritise their work, with a remarkable reduction of the delay between the data acquisition and the human diagnosis (it could be reduced to a few days).

- data associated to the images (i.e., metadata) and stored on the distributed system would be available to select the proper input for epidemiology studies or for the training of young radiologists.

These advantages would be granted by a GRID approach: the configuration of a Virtual Organisation, with common services (Data and Metadata Catalogue, Job Scheduler, Information System) and a number of distributed nodes providing computing and storage resources would allow the implementation of the screening, tele-training and epidemiology use cases. However, with respect to the model applied to High Energy Physics, there are some important differences: the network conditions do not allow the transfer of large amounts of data, the local nodes (hospitals) do not agree on the raw data transfer to other nodes as a standard and, most important, some of the use cases require interactivity.

According to these restrictions, our approach to the implementation of the GPCALMA Grid application is based on two different tools: AliEn [3] for the management of common services, PROOF [2] for the interactive analysis of remote data without data transfer.

3.1. Data Management

The GPCALMA data model foresees several Data Collection Centres, where mammograms are collected, locally stored and registered in the Data Catalogue. In order to make them available to a radiologist connecting from a Diagnostic Centre, it is mandatory to use a mechanism that identifies the data corresponding to the exam in a site-independent way: they must be selected by means of a set of requirements on the attached metadata and identified through a Logical Name which must be independent of their physical location. AliEn implements these features in its Data Catalogue Services, run by the Server: data are registered making use of a hierarchical namespace for their Logical Names and the system keeps track of their association to the actual name of the physical files. In addition, it is possible to attach metadata to each level of the hierarchical namespace. The Data Catalogue is browsable from the AliEn command line as well as from the Web portal; the C++ Application Program Interface (API) to ROOT is under development. Metadata
associated to the images can be classified in several categories: patient and exam identification data, results of the CAD algorithm analysis, radiologist’s diagnosis, histological diagnosis, etc. Some of these data will be directly stored in the Data Catalogue, but some of them may be stored in dedicated files and registered: the decision will be made after a discussion with the radiologists.

A dedicated *AliEn* Server for GPCALMA has been configured [4], in collaboration with the *AliEn* development team. Fig. 3 shows a screenshot from the WEB Portal.

![Screenshot from the GPCALMA AliEn WEB Portal](image)

Figure 3. Screenshot from the GPCALMA AliEn WEB Portal. Making use of the left side frame, the site can be navigated. General Information about the AliEn project, the installation and configuration guides, the status of the Virtual Organisation Services can be accessed. On the main frame, the list of the core services is shown, together with their status.

### 3.2. Remote Data Processing

Tele-diagnosis and tele-training require interactivity in order to be fully exploited, while in the case of screening it would be possible - although not optimal - to live without. The PROOF *Parallel ROOT Facility* system [2] allows to run interactive parallel processes on a distributed cluster of computers. A dedicated cluster of several PCs was configured and the remote analysis of a digitised mammogram without data transfer was recently run. As soon as input selection from the *AliEn* Data Catalogue will be possible, more complex use cases will be deployed. The basic idea is that, whenever a list of input Logical Names will be selected, that will be split into a number of sub-lists containing all the files stored in a given site and each sub-list will be sent to the corresponding node, where the mammograms will be analysed.

### 4. Present Status and Plans

The project is developing according to the original schedule. The CAD algorithms were rewritten in C++, making use of ROOT, in order to be PROOF-compliant; moreover, the ROOT functionality allowed a significant improvement of the Graphic User Interface, which, thanks to the possibility to manipulate the image and the associated description data, is now considered fully satisfactory by the radiologists involved in the project. The GPCALMA application code is available via CVS server for download and installation; a script to be used for the node configuration is being developed. The *AliEn* Server, which describes the Virtual Organisation and manages its services, is installed and configured; some *AliEn* Clients are in use, and they will soon be tested with GPCALMA jobs. The remote analysis of mammograms was successfully accomplished making use of PROOF. Presently, all but one the building blocks required to implement the tele-diagnosis and screening use cases were deployed. As soon as the implementation of the data selection from the ROOT shell through the *AliEn* C++ *API* will be available, GPCALMA nodes will be installed in the participating hospitals and connected to the *AliEn* Server, hosted by INFN. Hopefully, that task will be completed by the end of 2004.

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REFERENCES