

NUCLEAR MEDICINE IMAGE MANAGEMENT SYSTEM FOR STORAGE AND SHARING BY USING GRID SERVICES AND SEMANTIC WEB

Daniela Giordano, Carmelo Pino, Concetto Spampinato

Department of Informatics and Telecommunication Engineering, University of Catania, 95125 Catania, Italy

Marco Fargetta

Italian Institute for Nuclear Physics (INFN), Catania, Italy

Angela Di Stefano

Institute of Neurological Science, Italian Research Council (CNR), Catania, Italy

SCITEPRESS

Keywords: Medical data sharing, Logical file catalogue (LFC), Metadata service (AMGA), SPECT, PET, Web 2.0.

Abstract: Large amounts of images (SPECT, PET, scintigraphy) in the nuclear medicine field have been routinely produced in the last decades. In this paper we propose an image management system that allows nuclear medicine physicians to share the acquired images and the associated metadata both locally (i.e. within the same medical institute) and globally with other nuclear medicine physicians located anywhere in the world by using GRID services for data (LFC) and metadata (AMGA) storage. The proposed system guarantees medical data protection by anonymization that removes most sensitive data for unauthorized users, and encryption, that guarantees data protection when it is stored at remote sites. Another important issue is that often nuclear medicine data is associated with other medical data (e.g. neurological data) for diagnosis and therapy follow-up. In order to correlate images with other clinical information, the common metadata are enriched by developing a controlled vocabulary, which integrates known standards such as FOAF, CCR and GeneOntology. All the metadata are stored in an RDF (Resource Description Framework) repository in order to make the system fully compatible with existing metadata storage systems following the semantic web's philosophy.

1 INTRODUCTION

Usually, nuclear medicine physicians carry out examinations, e.g. Single Photon Emission Computed Tomography (SPECT), Positron Emission Tomography (PET) or scintigraphy, without a complete knowledge of the clinical history of the patients. In fact, most of the raw data collected in experiments or clinical trials is usually stored either in different places or in separate paper forms. Therefore there is a real need of integrating heterogeneous data sources in order to have a holistic vision of the patient's health status. Currently, existing methods, e.g. (Gabber et al., 2003), (Xu and Jiang, 2009) barely address medical data management needs beyond the specific depart-

ment's boundaries, while it is known that the patient medical folders are wide spread over many medical sites involved in the patient's healthcare. For this reason, many efforts have been done in last years to develop an interoperable infrastructure for digital health ((Cheung et al., 2009), (Freund, 2006)) by using semantic web concepts, even if the attention is mainly oriented to metadata sharing. A relevant project in this direction is the e-Child project (Freund, 2006) that develops a healthcare platform for pediatrics and aims at integrating ontologies to homogenize biomedical (from genomic, through cellular, disease, patient and population-related) data. The system proposed in this paper follows, and for some aspects, overcomes the above medical data management's approach be-

cause it permits to store images associating them with not directly related metadata (i.e. not included in the DICOM file) by developing an RDF controlled vocabulary, which links the most common ontologies used in the medical field with personal health records on the Internet (e.g. Google Health). This makes our system interoperable with methods that share data following semantic web (e.g. in (Holford et al., 2009)) enriches the common information with personal data, i.e. not present in medical facilities.

Given the huge amount of medical data involved in this process and the need of collaboration among researchers our system is provided with a GRID layer. Many GRID systems have been proposed for handling medical data geographically distributed on various medical centers, such as HOPE (Diarena et al., 2008), MediGrid (Montagnat et al., 2005) or MAMMOGRID (Amendolia, 2004), but most of them deal only with data files and do not provide higher level services for manipulating medical data or for handling the associated metadata, and also lack in the integration with other systems.

Therefore, in this paper we propose an approach for sharing and analyzing nuclear medicine images where 1) metadata are stored in RDF, thus giving the possibility to easily grab information coming from locations who share data according to the web semantic approach and 2) image sharing among different medical institutes is handled by GRID services in a transparent way for the users. The key features of the proposed system are:

- Full functionality in cases of network malfunctioning is ensured since data are stored both locally (in the user's computer and in the medical institute server) and globally on GRID;
- It is interoperable with existing sharing methods that use semantic web;
- It is flexible since researchers can share content choosing when and what to share;
- The metadata, usually contained in the DICOM files, are enriched in order to take into account the whole clinical history of patients. Users may decide to export all the information (data and metadata), only data, only metadata or only part of it;
- Data encryption is implemented using SSH, which ensures data integrity over Internet transfers;
- Data privacy is guaranteed by removing sensitive information before any transmission.

The paper is so organized: in section 2 the storage and sharing system is described. In detail, we present how the storage and the sharing work inside a medical institute using a client-server architecture and how

it works among different institutes using GRID services. Sect. 3 and Sect. 4 describes, respectively, the high level features for managing the interoperability with other systems, for image analysis and for system querying and the user interface. Finally, concluding remarks are given.

2 STORAGE AND SHARING SYSTEM OVERVIEW

In order to develop a distributed environment for image and information sharing to support the diagnosis, the treatment of patients and for statistical evaluation, the system is provided with two levels of storage and sharing: the first is locally managed by a client-server architecture, deployed in the medical institute where the nuclear medicine physicians belong to, whereas the second one is on GRID and allows global data sharing, i.e. data may be shared among different institutes using the services offered by the GRID computing. Fig. 1 shows the architecture of the proposed system for the local and global data sharing.

The typical use case is the following: a nuclear medicine physician stores the images and the metadata of a performed examination in its own local database (located in his/her computer). Afterwards, the client creates an anonymous version of the data removing all the confidential information so they can be sent to the main server in the respect of privacy issues. Additionally, the client allows users to define the set of metadata he/she wants to share both in GRID and in his/her medical institute.

The data transmission between client and server runs asynchronously in order 1) to make the system robust, in fact in case that, Internet connection is unavailable, data are locally stored and subsequently sent to the main server and to GRID when the connection will be available again and 2) to keep users unaffected by the actual time needed for the data transfer.

The server is provided with repositories where all the metadata produced within the same institute are stored. The communication with GRID is delegated to the server, thus optimizing the bandwidth's use.

2.1 Local Data Storage and Sharing

Inside a medical institute, data are stored and shared using a standard client-server architecture, as shown in fig. 2. The client and the server are connected by a local network or a VPN (Virtual Private Network).

The client contains the user interfaces and implements the logical communication with the GRID infrastructure. It also contains a file repository (for image stor-

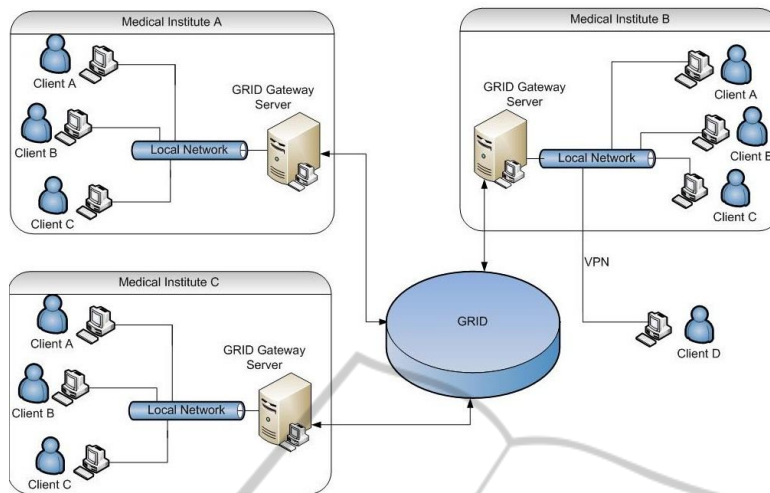


Figure 1: Architecture for Local and Global Data Storage and Sharing.

age) and a SESAME server (Broekstra et al., 2001) (for RDF metadata storage), in order to save patient's data locally.

The server also includes a file repository and a SESAME metadata repository for the data produced by all the nuclear medicine physicians within the same institute. Data are sent from the client to the server using File Transfer Protocol (FTP), whereas metadata is transmitted to SESAME server using Simple Object Access Protocol (SOAP) requests by means of a webservice, as shown in fig. 2.

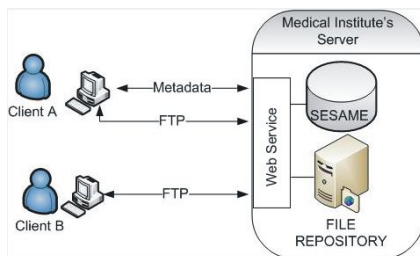


Figure 2: Local Data Storage and Sharing.

By using the client interface, a nuclear medicine physician can record and manage patients, add information to patient's clinical history (according to the schema shown in the next section), include any relevant documents (textual reports, generic images, DICOM images, etc.), run queries locally or on GRID data, associate the metadata deriving from the queries to the data locally stored and perform statistical analysis on sets of data and virtual data (i.e. coming from the main institute center or from GRID).

2.2 Global Data Sharing on GRID using LFC and AMGA

Data sharing among different institutes, geographically distributed, is implemented on GRID and it is based on the paradigm to create virtual environments where large amount of data and complex computations can be performed by different communities grouped in *Virtual Organisation (VO)*. Usually, each VO offers several services for GRID participants in order to simplify the data management providing them basic functionality to store and retrieve files.

The access to the GRID is hidden by a middleware. In this work we used the EGEE Grid and the G-Lite middleware¹. The two main services of the G-lite middleware for data and metadata storage, used in our system, are: the *Logical File Catalogue (LFC)* (Venugopal et al., 2004) and the *AMGA Metadata Service* (Nuno, 2006). The LFC allows users to associate a logical name to a file in a hierarchy format like a local file system, hiding the real location of the storage. Moreover, a logical name may refer many replicas, so a user can retrieve the file through its logical name from the nearest location in a transparent way. The AMGA Metadata Service is a special database designed to store metadata associated with files. Therefore, its internal structure reproduces a file system hierarchy where the directories are collections of metadata defined in a custom schema and each file in the directory is an entry containing the values for the metadata. This approach allows to easily map a file name with a set of metadata inside AMGA.

¹EGEE website - <http://www.eu-egee.org/> and gLite Grid middleware website - <http://www.glite.org/>

The communication with GRID infrastructure is managed by the medical institute server, which also aims at maintaining aligned the information stored in nuclear medicine physicians' computers and the one stored in GRID. In each institute the main server represents the bridge between clients and GRID, as shown in fig. 3.

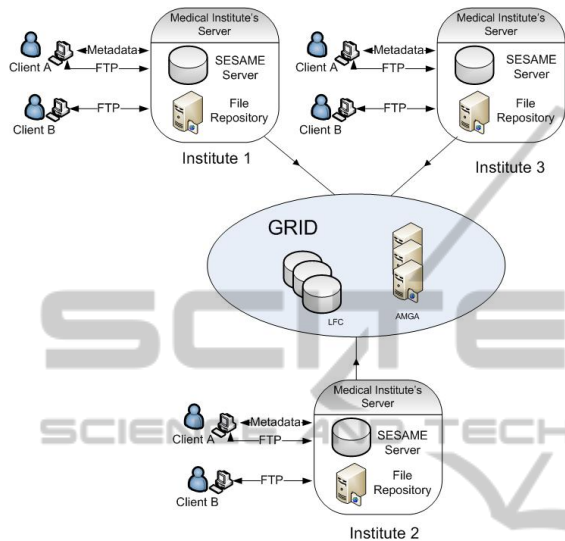


Figure 3: Global Data Storage and Sharing.

The flow diagram of the interaction between a generic medical institute and the GRID infrastructure is shown in fig. 4, when a nuclear medicine physician requires the storage of a specific image on GRID by using a proper GUI (Graphical User Interface). The steps performed by our system are: 1) the client sends to the server the proxy certificate (needed for the access to GRID) previously created and the identifier of the image to be stored on GRID, 2) the server of the medical institute (for simplicity called proxy server) queries the file repository to check if the file is present or not, if not it asks the client to send the image, then it sets the necessary permissions to read, write for the GRID, 3) the server queries the GRID database using the medical institute Grid ID given by the client in order to see if the image was already stored, if not 4) the server removes sensitive data, sends it to a Grid Storage Element, writes in the LFC catalog an appropriate logical name and writes (uploads if the image was previously stored) the metadata in the AMGA server.

3 HIGH LEVEL FEATURES

In order to provide useful information about the stored images and to make them available with the related

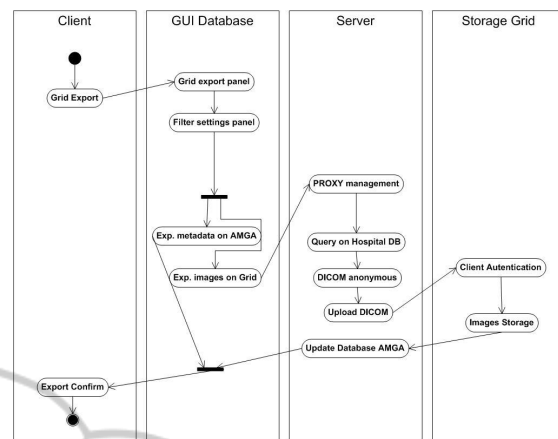


Figure 4: Interaction with Medical Institute - GRID Infrastructure.

metadata to the nuclear medicine community, the system is provided with high level features. More in detail, the system contains three processing levels:

- A semantic layer that enriches patient metadata by a controlled vocabulary using RDF/XML standard. This level guarantees the interoperability with existing frameworks;
- An image processing layer that analyzes the stored images. This is an important layer, since sometimes is desirable to share only the processing results and not the entire image. This level performs the image analysis and interacts with the semantic layer for the storage of the processing results in RDF/XML;
- Query Composition for performing complex queries both locally and on GRID. This module allows users to search useful information by processing only the metadata available locally or in GRID.

The interaction between the three levels and the system's architecture is shown in fig. 5.

3.1 Semantic Layer

Usually nuclear medicine images (e.g. in PET, SPECT) are stored in DICOM format, containing the metadata provided with the standard. These metadata are not sufficient for describing the clinical history of patients. For this reason we enrich the information available in order to give the nuclear medicine physicians the possibility to better figure out a specific disease by developing a model that represents the medical data so that it can be analyzed by semantic tools. In detail, the system stores concepts, specifies typed relationships between these concepts using

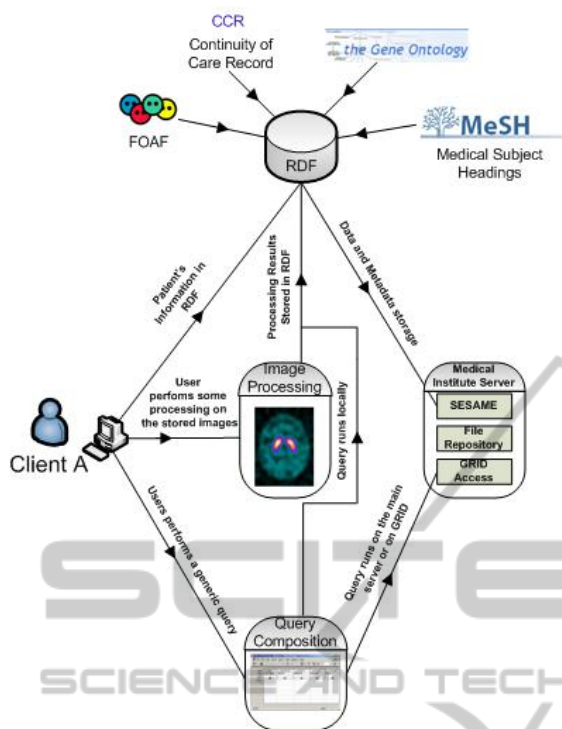


Figure 5: Interaction between the high level features' systems.

RDF² (Resource Description Framework) with XML syntax format. More in detail, we enrich the DICOM metadata by developing a controlled vocabulary that includes:

- Personal data by using FOAF ontology³;
- Generic Health Information according to the Continuity of Care Record (CCR) standard (Detmer et al., 2008) such as: diagnoses, allergies, medication list, immunizations, family history, social history, vital signs, procedures, symptoms, plan of care, functional status, biosignals (EEG, ECG, etc...);
- Genetic information using GeneOntology (Ashburner, 2000);
- Neurological detailed information by using Mesh (Soualmia et al., 2004);
- Image processing information that represents the output of the implemented image processing algorithms and which introduces a new semantic level to the stored metadata.

An example of RDF file for a patient is shown in fig. 6. It is notable that this information is inserted by the users, but it can be easily obtained by

²<http://www.w3.org/RDF/>

³<http://www.foaf-project.org/>

```
<rdf:RDF
xmlns:rdf="http://www.w3.org/1999/02/22-rdf-syntax-ns#"
xmlns:rdfs="http://www.w3.org/2000/01/rdf-schema#"
xmlns:foaf="http://xmlns.com/foaf/0.1/"
xmlns:ccr="http://www.ccr.com"
xmlns:go="http://www.geneontology.org/dtd/go.dtd#"
xmlns:mesh="http://org.ann.bike/MeSH#"
<foaf:Person>
<foaf:name>John Smith</foaf:name>
<foaf:firstName>John</foaf:firstName>
<foaf:surname>Smith</foaf:surname>
<foaf:mbox_sha1sum>71b88e951cb5f07518d69e5bb49a5100fbc3ea5</foaf:mbox_sha1sum>
<foaf:knows_rdf:resource="#Pippo">
</foaf:Person>
<foaf:Person_rdf:ID="#Pippo">
<ccr:Medication>
<ccr:Medication>
<ccr:Status><Text>Active</ccr:Text></ccr:Status>
<ccr:DateTime>
<ccr:Type><Text>Prescription Date</ccr:Text></ccr:Type>
<ccr:ExactDateTime>2007-04-01T05:00:00Z</ccr:ExactDateTime>
</ccr:DateTime>
<ccr:Source>
<ccr:Actor>
<ccr:ActorID>John Smith</ccr:ActorID>
</ccr:Actor>
<ccr:ActorRole>Prescribing clinician</ccr:ActorRole>
</ccr:Source>
<ccr:Product>
<ccr:ProductName>
<ccr:Text>Ibuprofen</ccr:Text>
<ccr:Code>
<ccr:Value>198405</ccr:Value>
<ccr:CodingSystem>RxNorm</ccr:CodingSystem>
</ccr:Code>
</ccr:ProductName>
</ccr:Product>
</ccr:Medication>
</ccr:Medications>
<go:dbxref>
<go:database_symbol>interpro</go:database_symbol>
<go:reference>IIR003999 Staphylococcal toxic shock syndrome toxin
</go:reference>
</go:dbxref>
</foaf:Person>
</rdf:RDF>
```

Figure 6: Example of produced RDF File.

querying systems that share data using RDF. For instance, personal data in FOAF can be derived from a generic social network or by using a v-card, whereas generic health information can be obtained by the user's Google Health Account⁴ or other systems that aim at storing online health care data such as the one proposed in (Bieliková and Moravčík, 2008). Metadata storage has been carried out by using SESAME server (Broekstra et al., 2001) so that these information may be available also for other purposes. The sensitive data, such as Name, Surname, SSN must be available only for the nuclear medicine physician who carries out the examination, and are not exported in RDF in order to ensure data privacy.

3.2 Image Processing Layer

This level is provided with a set of processing methods for medical image analysis. The output of this processing is stored according to the semantic layer and is related to the specific processed image. This allows users to also share the results of the processing avoiding to send the original images when it is not required, resulting in less bandwidth usage. The implemented methods for nuclear image analysis, also available for MRI, X-Rays images, are:

- Measurement of distances, angles and some parameters within the images;
- The contrast absorption curve over time;

⁴<https://www.google.com/health/>

- Image Texture and Image Contour Analysis for specific organs;
- Pattern recognition for identifying brain structures.

Example of such algorithms have been proposed by the authors in (Faro et al., 2010a), (Faro et al., 2010b) and (Giordano et al., 2009). Therefore, when a user performs one of the above methods, the output is treated as metadata and stored in the SESAME server.

3.3 Query Composition

The query composition level aims at building complex queries both locally and on GRID. The queries are performed only on the metadata (stored in the SESAME server and in the AMGA server) since a content based image retrieval module is not present. This level receives the user's queries (by using a controlled GUI) and interacts both with the local storage, performing SPARQL queries on the SESAME server, and with the GRID, where queries are performed following the approach proposed in (Montagnat et al., 2008).

4 CLIENT INTERFACE

The client interface consists of a multiform Java application, allowing users to manage any type of information concerning the patients. The functionalities available for nuclear medicine physicians are:

- view information about their patients ;
- search a patient by typing the name;
- insert, modify or delete a patient;
- print a patient's report;
- show the patient exams;
- insert, modify or delete an exam;
- management of medical history;
- view and process DICOM files;
- export the patient's data on GRID;
- query the system both locally and globally on GRID.

For each functionality a specific GUI has been designed. For example, fig. 7 shows the main interface where a list of patients and a tool to manage patients' data is available. In the bottom area, there is also a section to handle the exams for each patient. Fig. 8, instead, shows the interface for DICOM files and image processing.

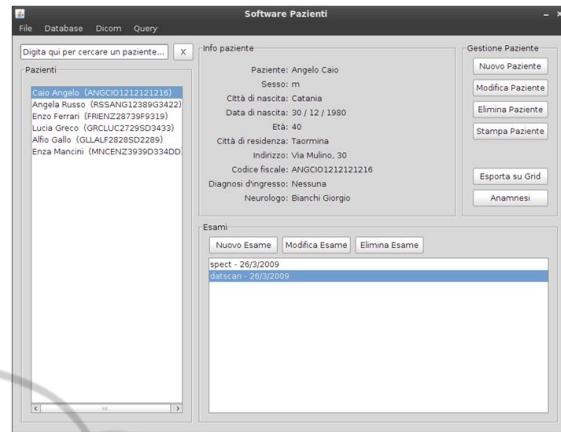


Figure 7: Main GUI.

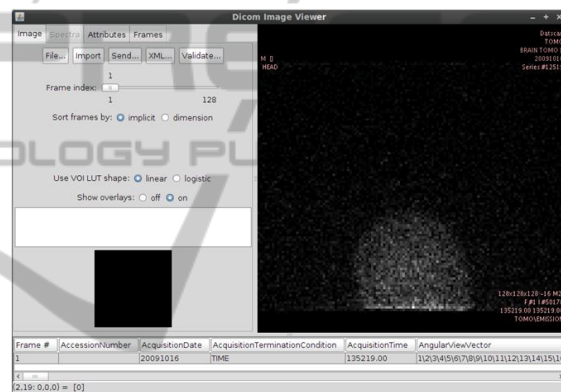


Figure 8: GUI for DICOM File Processing.

Moreover, for expert users the client application contains a form (fig. 9) to set parameters for the FTP, GRID and AMGA.

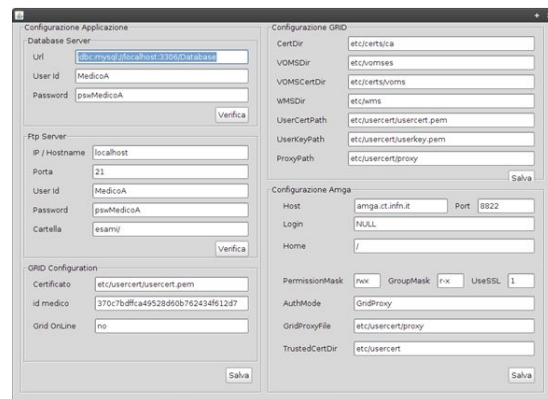


Figure 9: GUI for parameters' setting.

5 CONCLUDING REMARKS

In this paper we presented a distributed system of data and metadata storage, sharing and processing. Web semantic data modeling using RDF permits that all the information can be stored and reused. The RDF modeling allows us to efficiently integrate heterogeneous data into a coherent whole and to provide description of data elements. The area of application, is still confined to the specific sector of nuclear medicine but the flexibility of the architecture permits the application to be used on a greater number of medical fields by a simple re-modeling. The platform will be integrated with an image retrieval system in order to assist the physicians in the retrieval of images by content. A future development is to provide the system with an ontology layer that will map high level requests to low level queries.

REFERENCES

- Amendolia, S. R. (2004). Mammogrid: A service oriented architecture based medical grid application. In *GCC*, pages 939–942.
- Ashburner, M. (2000). Gene ontology: Tool for the unification of biology. *Nature Genetics*, 25:25–29.
- Bieliková, M. and Moravčík, M. (2008). Modeling the reusable content of adaptive web-based applications using an ontology. In *Advances in Semantic Media Adaptation and Personalization*, pages 307–327.
- Broekstra, J., Kampman, A., and Harmelen, F. V. (2001). Sesame: An architecture for storing and querying rdf data and schema information. In *Semantics for the WWW*. MIT Press.
- Cheung, K. H., Prud'hommeaux, E., Wang, Y., and Stephens, S. (2009). Semantic Web for Health Care and Life Sciences: a review of the state of the art. *Brief. Bioinformatics*, 10:111–113.
- Detmer, D., Bloomrosen, M., Raymond, B., and Tang, P. (2008). Integrated personal health records: Transformative tools for consumer-centric care. *BMC Medical Informatics and Decision Making*, 8(1):45.
- Diarena, M., Nowak, S., Boire, J. Y., Bloch, V., Donnarieix, D., Fessy, A., Grenier, B., Irrthum, B., Legre, Y., Maigne, L., Salzemann, J., Thiam, C., Spalinger, N., Verhaeghe, N., de Vlieger, P., and Breton, V. (2008). HOPE, an open platform for medical data management on the grid. *Stud Health Technol Inform*, 138:34–48.
- Faro, A., Giordano, D., Pino, C., Spampinato, C., and Di Stefano, A. (2010a). 3D striatum reconstruction of 123ioflupane MRI images for quantitative assessments on the dopaminergic neurotransmission system. In *IEEE International Workshop on Medical Measurements and Applications (MeMeA 2010)*, Ottawa, Canada.
- Faro, A., Giordano, D., Spampinato, C., and Pennisi, M. (2010b). Statistical texture analysis of MRI images to classify patients affected by multiple sclerosis. In *12th Mediterranean Conference on Medical and Biological Engineering and Computing MEDICON 2010*, pages 236–239, Porto Carras, Chalkidiki, Greece. International Proceedings of the IFBME, Springer.
- Freund, J. (2006). Health-e-child: an integrated biomedical platform for grid-based paediatric applications. *Stud Health Technol Inform*, 120:259–270.
- Gabber, E., Fellin, J., Flaster, M., Gu, F., Hillyer, B., Ng, W. T., Özden, B., and Shriver, E. A. M. (2003). Starfish: highly-available block storage. In *USENIX Annual Technical Conference, FREENIX Track*, pages 151–163.
- Giordano, D., Scarciofalo, G., Spampinato, C., and Leonardi, R. (2009). Automatic skeletal bone age assessment by integrating EMROI and CROI processing. In *IEEE International Workshop on Medical Measurements and Applications (MeMeA 2009)*, Cetraro, Italy.
- Holford, M. E., Rajeevan, H., Zhao, H., Kidd, K. K., and Cheung, K. H. (2009). Semantic Web-based integration of cancer pathways and allele frequency data. *Cancer Inform*, 8:19–30.
- Montagnat, J., Breton, V., and Magnin, I. (2005). Partitioning medical image databases for content-based queries on a Grid. *Methods Inf Med*, 44:154–160.
- Montagnat, J., Frohner, Á., Jouvenot, D., Pera, C., Kunszt, P., Koblitz, B., Santos, N., Loomis, C., Texier, R., Lingrand, D., Guio, P., Rocha, R. B. D., de Almeida, A. S., and Farkas, Z. (2008). A secure grid medical data manager interfaced to the glide middleware. *J. Grid Comput.*, 6(1):45–59.
- Nuno, N. S. (2006). Distributed metadata with the amga metadata catalog. In *Workshop on NextGeneration Distributed Data Management - HPDC-15*.
- Soualmia, L., Golbreich, C., and Darmoni, S. (2004). Representing the mesh in owl: Towards a semi-automatic migration. In *Proceedings of the KR 2004 Workshop on Formal Biomedical Knowledge Representation*, pages 81–87.
- Venugopal, S., Buyya, R., and Winton, L. (2004). A grid service broker for scheduling distributed data-oriented applications on global grids. In *MGC '04: Proceedings of the 2nd workshop on Middleware for grid computing*, pages 75–80, New York, NY, USA. ACM.
- Xu, Z. and Jiang, H. (2009). Hass: Highly available, scalable and secure distributed data storage systems. *Computational Science and Engineering, IEEE International Conference on*, 2:772–780.