

# An Image-based Framework for Intellectual Decision Making in Medical Research

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**Abstract.** Medical decisional problems strongly rely on the analysis and interpretation of diagnostic images acquired by different investigation techniques. Information extraction and correlation from these are usually demanding tasks that can burden the routine work of clinicians. In this paper, an image-based framework is presented which is devoted to the automated extraction of knowledge and data from biomedical images used for intellectual decision making in clinical and medical research. An ontological approach is adopted for encoding different kind of knowledge required in problem solving. Modelling the pre-clinical stage of Parkinson's disease is presented as eligible case study.

## 1 Introduction

An important step in medical diagnostics is analysis of data extracted from digital images, i.e. microscopic, X-ray, MRT and ultrasonic images. For efficient assistance of problem solving is necessary to automate of biomedical image analysis and recognition and to develop suitable information technologies and tools.

During the last years, authors' research has been focused on the investigation and development of image mining methods for solving and improving knowledge in medical tasks, in particular, for disease modelling and diagnostics. In previous works [1, 2], a novel knowledge-based approach to medical image mining was described. The approach is based on an ontological representation of the knowledge needed for problem solving. Ontologies are intended to describe the following kinds of knowledge: 1) knowledge on image processing, analysis, storage, and retrieval; 2) knowledge on processing and analysis of experimental data; 3) knowledge on prediction of diseases progression and treatment results. The proposed approach was developed as a general framework for medical image mining, and then specialized for automation of cytological image analysis used for early diagnostics of oncological blood diseases. In particular, the cell image analysis ontology was developed.

The general value of the framework is allowing the development of a new application devoted to investigate the definition of a preclinical model of Parkinson's disease (PD). More precisely, thanks to the flexible design of the framework, the new

application is just requiring the definition of a specific part of the ontological knowledge base, i.e., the domain ontology and base of rules related to the PD medical domain.

In the next sections, the framework is briefly described, while attention is focused on the specific problem domain and the dedicated knowledge base that is being developed.

## 2 Related Works

Ontologies as an effective way for knowledge representation became very popular last years. Different works related to usage of ontologies for solving image-based tasks have been reported. For example, in [3] an approach for solving the symbol grounding problem involved in semantic image interpretation is presented. The method is based on using the image processing ontology to reduce the gap between the image processing level and the visual level. Authors note that the proposed ontology is not complete and should be considered as a basis for further extension. In [4] a platform dedicated to the knowledge extraction and management for image processing applications is proposed. It includes a system that automatically generates image processing applications on the basis of goal formulations given by a user who is inexperienced in image processing domain. The user defines the goal of processing in terms of his/her application domain and then the system translates this information into image processing terms taken from the image processing ontology. The result of this translation is an image processing request which is sent to the planning system to generate the program that responds to this request.

The main contribution of our work is the development of a sufficiently detailed and well-structured ontology which will cover all important aspects of image processing, analysis and understanding (main categories of concepts, their properties and relations). The proposed ontology can be used as a base for the construction of specialized knowledge bases for supporting image analysis and, then, image mining.

## 3 The Knowledge-based Framework for Image Mining

One of actual and difficult problems in modern medicine is the development of systems for early diseases diagnostics at the preclinical stage. The automation of diagnostics requires automated extraction of information from medical images used for diseases modelling, treatment and intellectual decision making.

To this end, a knowledge-based framework was conceived for aiding the investigation and diagnostic processes of medical problems, by providing local and remote access to known cases, the facility for retrieving images by content, and the possibility of mining image patterns relevant to medical decision making, e.g., diagnoses and prognoses [1].

Several practical scenarios can be supplied for highlighting the usefulness of framework functionalities:

- *Clinicians' case-based reasoning*, i.e., medical decisions made by retrieving similar cases according to patient's information on imagery data;
- *Investigation of concurrent situations*, i.e., exploration of other research results of the same kind but in other fields;
- *Extraction of new knowledge*, i.e., investigation on the relevance of new information (new data, new parameters, new relations among them) for some still debated problems.

A brief overview of the framework can be useful to recall and present its main functionalities and component.

The framework was functionally designed as shown in Fig. 1. More precisely, along with the data storage and management inside a repository, three main and inter-related end-users functionalities were identified:

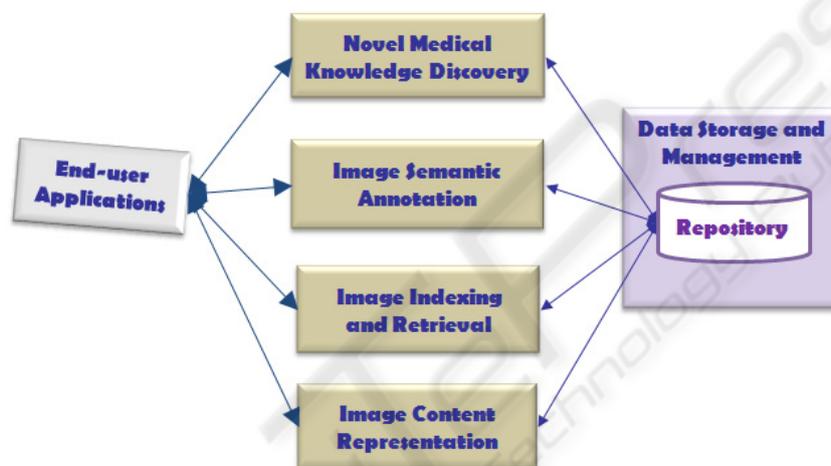


Fig. 1. The knowledge-based framework functionalities.

- Image indexing and retrieval, for finding images ranked in accordance to some requirements on their content. Retrieval can be performed by means of explicit text query or by supplying a reference image. Similarity measures are applied to appropriate image representations for identifying the relevant images to be retrieved;
- Image semantic annotation, for defining a list of semantic keywords to be associated to image and used for their retrieval, in particular for text queries. A structured terminology is supplied for aiding annotations by the users;
- Novel medical knowledge discovery, for extracting valid, novel and understandable knowledge about the diagnostic, prognostic and monitoring processes. Advanced data mining methods can be applied to patterns built by correlating features extracted from images and domain concepts.

The main components of framework, as sketched in Figure 2, are:

- a repository for storing, accessing and retrieving images and information extracted at different levels from them;

- a suite of ontologies including specific domain ontologies related to medical problems and a general image analysis ontology [7];
- a collection of algorithms for image processing, analysis, recognition and mining;
- a user interface for accessing, uploading, browsing and annotating images.

For a detailed description of these components see [1], here we concentrate on the specific application and, hence, on the definition of the *Brain Image Mining Ontology*.

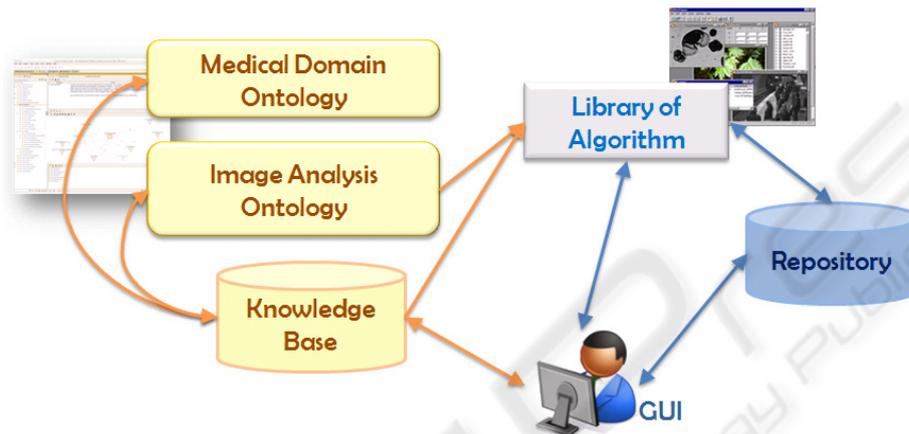


Fig. 2. The main framework components.

## 4 Diagnosis of Neurodegenerative Diseases

The framework is being extended in order to suitably define a model of preclinical stage of Parkinson's disease. In the next subsections, the problem domain and the framework extensions are presented.

### 4.1 Automated Early Diagnostics of Neurodegenerative Diseases

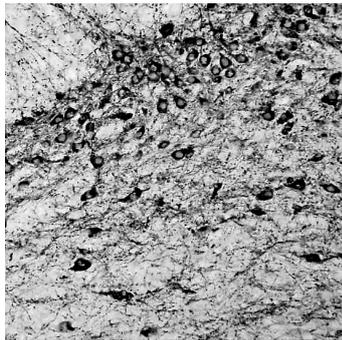
The investigation and modelling of PD is a crucial and always present problem in medicine. Construction of experimental models is crucial for the research of neurodegenerative disease pathogenesis. PD is characterized by a progressive degeneration of dopaminergic (DA-ergic) neurons in the substantia nigra pars compacta (SN) leading to a dopamine (DA) depletion in the striatum. As a result, parkinsonian patients lose the ability to control their movements.

Current tasks are automation of experimental data extraction for filling a model of PD preclinical stage and automation of model investigation by means of computer experiments.

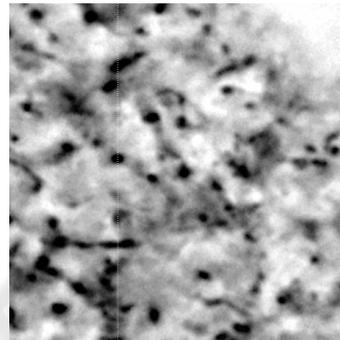
The development of PD preclinical stage model is a complex screening analysis which is being done cooperatively by physiologists, biochemists and morphologists.

The morphological research requires processing and analysis of astronomical quantity of experimental animal serial brain section images. And studying of each of the sections requires quantitative and qualitative feature measurements and analysis of several thousand neurons and axons [5].

The initial data is digital images of the immunostained sections of various brain areas. DA-ergic neurons were labelled on serial sections (a thickness of 20 microns) of the substantia nigra (Fig. 3) and their fibres (axons) on sections of the striatum (a thickness of 12 microns) (Fig. 4) by immunohistochemistry for tyrosinehydroxylase (TH) (TH is the specific enzyme of DA synthesis). Experimental data has been received from digital images of distal parts of axons (terminals).



**Fig. 3.** Neurons.



**Fig. 4.** Terminals.

Terminals are small rounded objects with an area varying from 0.6 to 3 m<sup>2</sup>. Terminals can have an oval, round, prolate or irregular shape. In the presented gray-scale images, the brightness of terminals is lower than the background brightness.

The PD model represents the differences between experimental and control groups. The former is a group of animals injected with a toxin, while the latter is a group of animals not affected by the toxin. A major characteristic of the PD model is the number of DA-ergic axons innervating the striatum in the case of using various schemes for neurotoxin administration (dose, the number of injections, intervals between injections). The extent of degeneration is defined as the difference between the number of terminals of DA-ergic axons in the control and experimental groups. DA-ergic neurons and axons remaining after neurotoxin administration are supposed to demonstrate increased functional activity in order to compensate for the DA deficiency. An indicator of the increased functional activity of neurons and their fibres can be an increase in their sizes. An increase in the concentration of tyrosinehydroxylase (key enzyme in DA synthesis) is supposed to be another specific indicator of the functional activity of DA-ergic axons and neurons.

#### **4.2 The Brain Image Knowledge Base**

The study of PD can be performed from different perspectives which should be considered as a whole to have a better view of the disease. More specifically, in the system physiology perspective, the structures (anatomy) involved in the disease are con-

sidered, while the ways those structures interact or fail to interact belong to the physiology perspective of the disease state. Moreover, the genetic perspective is becoming more and more practised, since the discovery that genetic mutations in the alpha synuclein gene could cause PD has opened new avenues of research in PD.

The idea behind the extension of the knowledge based framework is to provide PD researchers with an instrument that allows them to search for particular patients and particular images of disease states, correlate different kinds of data (of the same patient or of different patients), to formulate hypotheses on the disease state, causes and evolution. Examples of usage of the framework can be the following [6]:

- A researcher interested in anatomy/physiology trying to hypothesize the location of various components on a neuron might want to look at scientific data or hypotheses on the presence of certain type of receptors on a neuron.
- A researcher interested in molecular biology trying to hypothesize the location of some receptors on neurons might want to look at scientific data or hypotheses on the anatomical structure of a neuron or the pharmacology of chemical compounds that bind to a receptor.
- A clinical researcher interested in developing new therapies for a disease may be interested in understanding the mechanisms of how chemical compounds or ligands bind to receptors.

For providing these functionalities, a knowledge base able to integrate different kind of data is being developed. In particular, the information required belongs to:

- anatomy and physiology;
- molecular biology;
- clinical information.

Such knowledge base is being developed by integrating a dedicated ontology and a base of rules which use the ontology concepts and are aimed at encoding more complex relations among them. Several existing biomedical ontologies have been taken into account for reuse [6]. Since the idea is to consider all the information belonging to the three domains listed above, sub-hierarchies related to neuron anatomy (class *Neuron*, and all the meronymy classes), proteins, genes and general information related to disease are being considered. Figure 5 shows a fragment of the ontology. The rules are being encoded by discussing with domain experts who suggest some relevant relations among domain relevant data and information.

Moreover, the library of algorithms is being extended by inserting methods for image segmentation and classification, according to the specifications contained in [5].

In this way, the ontology can be used to annotate images and retrieve particular cases according to users' requirements. On the other hand, the most complex information contained in the knowledge base will allow users to formulate hypotheses and check their validity but retrieving images and data that support and evidence its validity.

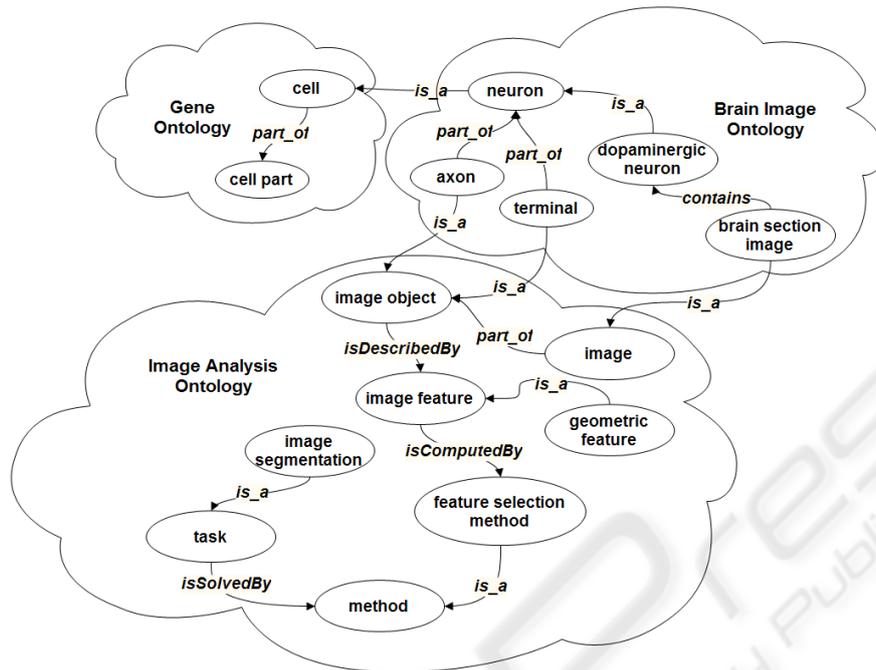


Fig. 5. The fragment of the Brain Image Mining Ontology.

## 5 Conclusions

An image-based framework to biomedical image mining has been presented. The main attention was focused on the problem of extraction and representation of knowledge and data required for intellectual decision making in clinical and medical research. Modelling the preclinical stage of Parkinson's disease has been considered as eligible case study. The requirements to the information represented in the knowledge base on brain image mining have been described.

Future work will be devoted to further development of the knowledge base described, in particular, to the completion of the ontology suite aimed at encoding the knowledge on storing, retrieving, processing, analysis and mining of biomedical images.

## Acknowledgements

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