

# João D. Ferreira

COMPUTER SCIENTIST · BIOINFORMATICIAN

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## Summary

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### Professional Experience

- 2015–Present** Professor Auxiliar Convitado (50%)  
*at Faculdade de Ciências da Universidade de Lisboa*
- 2016–Present** Collaborator  
*at Laboratório de Sistemas de Informação de Grande Escala*

### Scientific Performance

- Education** Ph.D., Computer Science, Universidade de Lisboa
- Publications** 6 peer-reviewed journal publications;  
6 conference and workshop publications;  
total citations: 92  
*h*-index: 6
- Software** 3 open-source software programmes;  
1 online web application
- Knowledge resources** 2 knowledge-representation artefacts
- Projects** participation in 2 national projects and 1 European project
- Academic Supervision** 1 M.Sc. in Informatics Engineering
- Conference organization** ICBO 2015 Workshop & tutorial Chain  
ICBO 2015 Session chair
- Reviewer** 3 international journals;  
2 international conferences;  
2 national conferences
- Honors and Awards** 2 scholarships (PhD and MSc)

### Pedagogical Performance

- General topics** Bioinformatics;  
Web applications;  
Human-Computer Interaction;  
Databases;  
General programming.
- Courses** 4 first-cycle
- Pedagogical Surveys** Average of 3.78 out of 4

### University Mission

- Relevant projects** FCUL Open Day;  
Semana da Ciência em Movimento;  
2<sup>as</sup> Jornadas GIMGAS

## A Scientific performance

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### A.1 Scientific production

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#### Selected list of publications

[1] Andre Lamurias, **João D. Ferreira**, and Francisco M. Couto. “Improving chemical entity recognition through h-index based semantic similarity.” In: *Journal of Cheminformatics* 7.Suppl 1 Text mining for chemistry and the CHEMDNER track (Jan. 2015), S13. ISSN: 1758-2946. DOI: [10.1186/1758-2946-7-S1-S13](https://doi.org/10.1186/1758-2946-7-S1-S13)

- **Impact Factor:** 4.550
- **Non-self citations:** 4

[3] Catia Pesquita, **João D. Ferreira**, Francisco M. Couto, and Mário J. Silva. “The epidemiology ontology: an ontology for the semantic annotation of epidemiological resources.” In: *Journal of Biomedical Semantics* 5.1 (2014), p. 4. ISSN: 2041-1480. DOI: [10.1186/2041-1480-5-4](https://doi.org/10.1186/2041-1480-5-4)

- **Impact Factor:** 2.262
- **Non-self citations:** 2

[4] **João D. Ferreira**, Janna Hastings, and Francisco M. Couto. “Exploiting disjointness axioms to improve semantic similarity measures.” In: *Bioinformatics* 29.21 (2013), pp. 2781–2787. ISSN: 1367-4811. DOI: [10.1093/bioinformatics/btt491](https://doi.org/10.1093/bioinformatics/btt491)

- **Impact Factor:** 4.981
- **Non-self citations:** 9

[5] **João D. Ferreira**, Daniela Paolotti, Francisco M. Couto, and Mário J. Silva. “On the usefulness of ontologies in epidemiology research and practice.” In: *Journal of epidemiology and community health* (Nov. 2012), pp. –3. ISSN: 1470-2738. DOI: [10.1136/jech-2012-201142](https://doi.org/10.1136/jech-2012-201142)

- **Impact Factor:** 3.501
- **Non-self citations:** 5

[6] **João D. Ferreira** and Francisco M. Couto. “Semantic Similarity for Automatic Classification of Chemical Compounds”. In: *PLoS Computational Biology* 6.9 (Sept. 2010). Ed. by John B. O. Mitchell, e1000937. ISSN: 1553-7358. DOI: [10.1371/journal.pcbi.1000937](https://doi.org/10.1371/journal.pcbi.1000937)

- **Impact Factor:** 4.620
- **Non-self citations:** 33

#### Selected list of citations

Janna Hastings et al. “The ChEBI reference database and ontology for biologically relevant chemistry: enhancements for 2013”. In: *Nucleic acids research* 41.D1 (2013), pp. D456–D463

- This paper is a major milestone in Cheminformatics, describing ChEBI (an ontology of chemical compounds). It has itself 163 citations. Cites my paper [6] as an example of an application that makes use of the semantic information encoded in ChEBI for automatic classification.

Janna Hastings et al. “Structure-based classification and ontology in chemistry.” In: *J. Cheminformatics* 4 (2012), p. 8

- This paper expresses the need and usefulness of systems that exploit the machine-readable information provided by reference ontologies, and cites my paper [6] as an example.

Joerg Kurt Wegner et al. “Cheminformatics”. In: *Communications of the ACM* 55.11 (2012), pp. 65–75

- This paper advocates the use of open-source solutions to deal with the vast amount of existing chemical information, mentioning semantic similarity in ChEBI and citing my paper [6].

Robert Hoehndorf et al. “Thematic series on biomedical ontologies in JBMS: challenges and new directions.” In: *J. Biomedical Semantics* 5 (2014), p. 15

- This paper aims “to disseminate the latest developments in research on biomedical ontologies and provide a venue for publishing newly developed ontologies, updates to existing ontologies as well as methodological advances, and selected contributions from conferences and workshops”. It cites my paper [6] as an example of a semantic similarity application and my paper [4] as the start of using more than taxonomies in semantic similarity in the biomedical domain.

Stefan Schulz et al. “From concept representations to ontologies: a paradigm shift in health informatics?” In: *Healthcare informatics research* 19.4 (2013), pp. 235–242

- Lists my paper [5] as one of the twenty most influential papers in the areas of knowledge-representation, biomedical ontologies and electronic health issues, as listed by the members of the LinkedIn Working Group on Medical Concept Representation.

## Histograms of publications

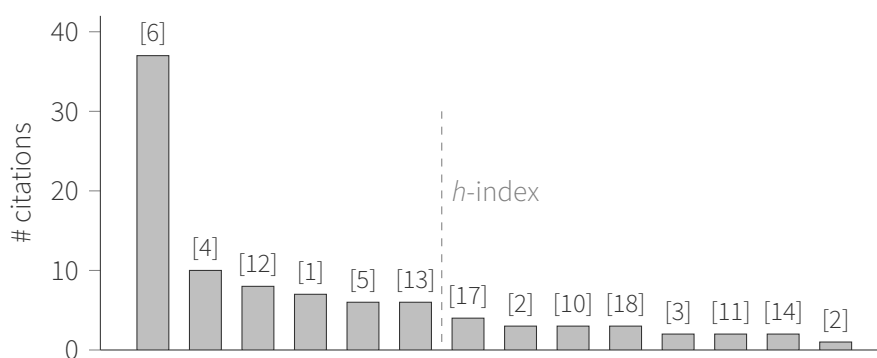


Figure 1: Histogram of citations for each of my cited papers, with papers sorted from most cited to least cited. Top labels specify the publication according to Section D. The vertical dashed line shows the associated  $h$ -index.

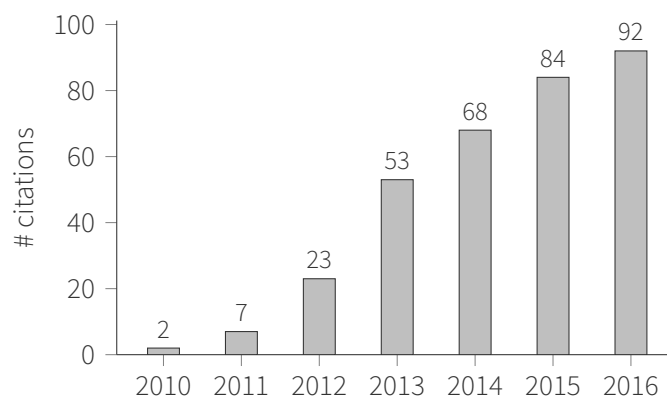


Figure 2: Histogram showing the number of citations per year.

## Citation metrics

Table 1: Citation metrics computed using Google Scholar and DBLP data.

	Metric	Value
Google Scholar	<i>h</i> -index	6
	i10-index	2
	citations	92
	cited papers	14
DBLP	co-authors	15
	#journal articles	5
	# conference papers	6

## Open Software

### OWLtoSQL

2015

URL: <http://github.com/jdferreira/owltoSQL>

- This software is one of the results of my PhD work. It is used to convert an ontology written in OWL (Web Ontology Language) into a relational database, which offers several advantages, the most prominent of which is that it enables *random* access to the ontology constituents (classes, properties, axioms, etc.). It has been used by Bruno Inácio in his M.Sc work (see Section A.3).

### Multi-domain Ontology-based Semantic Similarity (MOSSy)

2015

URL: <http://github.com/jdferreira/mossy>

- This software also resulted from my PhD work. It is responsible for calculating semantic similarity between resources annotated with ontology concepts. The program is configurable to use any OWL ontology and can handle any type of annotated resource, both in single- and multi-domain contexts. It is extensible, since it allows the implementation of semantic similarity measures as Python classes.

### Annota

2010

URL: <https://code.google.com/archive/p/annota/>

- This software was used internally in the GREASE-II project in order to manually disambiguate ambiguous automatic tags produced from free-text news reports.

## Web applications

### MOSSy

2015

URL: <http://lasige.di.fc.ul.pt/webtools/mossy>

- This web application is a gateway to apply MOSSy (the program, see above) to a collection of annotated resources. The application first requests from the user a set of models which can be given in free text or already annotated resources. If free-text models are provided, a text-mining service is used to extract annotations. Given the set of models, the application compares them according and outputs a cluster of the resources as well as the similarity matrix between them.

The software was used by a research team in University College London to showcase similarity between physiological models of tissue, but is available for use by the whole scientific community.

## Knowledge representation artefacts

### Network of Epidemiology-Related Ontologies

2013

SEE MY PAPER [5]

- This work is a collection of ontologies related to the epidemiology domain. It contains 13 ontologies, from domains such as biochemistry, diseases, environment, transmission, vaccines, and geography,

which provide the necessary concepts to annotate a digital resource of the epidemiological domain. The network was later included in the Epimarketplace, a repository of epidemic resources, in order to facilitate the annotation process.

- **Developed:** during my work in the EPIWORK project.

### **Epidemiology Ontology**

2014

URL: <https://code.google.com/p/epidemiology-ontology/>

- This ontology represents both disease transmission methods and epidemiology parameters, containing over 300 concepts. This ontology fills the previously existing gap in these areas, which were under-represented in biomedical ontologies. It supports the semantic annotation of resources.
- **Developed:** during my work in the EPIWORK project.

## **A.2 Research projects**

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### **Past projects**

#### **Semantic Ontology Matching using External Resources (SOMER)**

2012-2014

URL: <http://somer.di.fc.ul.pt/>

- **Position:** Member of research team
- **Funding:** FCT (84 000€)
- **Grant:** PTDC/EIA-EIA/119119/2010
- **Description of project:** The SOMER project aimed to develop ontology matching methods that exploit external knowledge resources through evidence and information content provided by unstructured text and annotation corpora. The results were applied to real world applications based on existing ontologies in the biomedical and geospatial areas.
- **My contributions:** I was involved in writing the proposal for this project. I was also responsible for exploring ways that allowed the use of semantic similarity measures in the process of ontology matching. I contributed to 1 peer-reviewed publication.

#### **Developing the Framework for an Epidemic Forecast Infrastructure (EPIWORK)**

2012-2014

- **Position:** Hired researcher
- **Funding:** FP7 (5 000 000€)
- **Grant:** 231807
- **Description of project:** The EPIWORK project proposed a framework of tools and knowledge for the design of epidemic forecast infrastructures, including: development of the mathematical and computational methods needed to achieve prediction of disease spreading in complex social systems; development of large scale, data driven computational models aimed at epidemic scenario forecast; design and implementation of data-collection mechanisms, such as the collection of real-time disease incidence, through innovative web and ICT applications; and the implementation of a computational platform for epidemic research and data sharing.
- **My contributions:** The LASIGE team was responsible for the creation of the Epidemic Marketplace, an online platform for epidemic research and data sharing was the data-hub for the multiple research communities and countries involved in the project. I was hired by the LASIGE team as a consultant in semantic web, specifically to design the semantic metadata model that was used to annotate the resources of the marketplace, as well as to design the Network of Epidemiology-Related Ontologies, which contributed to the repository as a collection of concepts used to annotate the resources. I contributed to 5 peer-reviewed publications and 1 open-source ontology.

## Geographic Reasoning for Search Engines II (GREASE-II)

2010

- **Position:** M.Sc fellowship
- **Funding:** FCT (117 300 €)
- **Grant:** PTDC/EIA/73614/2006
- **Description of project:** This project aimed at researching information access methods to large collections of documents and objects having geographically rich text and meta-data. One of the main ideas of the project was that geospatial information can be mapped into ontology concepts.
- **My contributions:** One of the tasks of this project was to perform text-mining directly onto news reports in order to find geographical references in text; to assist in this task, a disambiguation module had to be created. I was assigned the creation of software to help disambiguate annotations manually, which would then later be used in a machine-learning step. I was also tasked with the alignment of Geo-Net-PT (a geographical ontology of the Portuguese territory) to Yahoo! GeoPlanet™ (an infrastructure for geo-referencing data on the Internet). I contributed to 1 technical report and to 1 open-source software

## International Collaborations

### European Bioinformatics Institute (EBI)

2013-2016

- I collaborated with J. Hastings, former group coordinator of the Cheminformatics and Metabolism Team at EBI, in devising a semantic similarity measure that can effectively capture the information provided by the disjointness axioms of an ontology. This collaboration resulted in 1 peer-reviewed publication.
- I am also currently collaborating with the team behind Metabolights, a repository used to store details of experiments related to research in metabolism and information derived from the research. My role in this collaboration is to help devise ways to measure annotation quality in order find resources under-annotated and to help authors submit properly annotated data. This collaboration is still ongoing and is expected to result in at least 1 peer-reviewed publication.

## A.3 Academic supervision

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### Current supervision

#### M.Sc in Informatics Engineering

2015-2016

UNIVERSIDADE DE LISBOA

- **Student:** Bruno Inácio
- **Position:** Co-supervision with Francisco M. Couto, Universidade de Lisboa
- **Thesis:** “Quanto valem os metadados?”
- **Topic:** The work focussed (i) on the study of the quality of metadata as a way to sustain semantic integration and to facilitate data-sharing, based on the specificity of the ontology concepts used to annotate the resources and on the thoroughness of these annotations; and (ii) on the development of a platform that allows the assessment of metadata quality of the resources in a scientific data repository.

#### Internship of a B.Sc in Biochemistry

2011-2012

UNIVERSIDADE DE LISBOA

- **Student:** Hugo Ferreira
- **Position:** Co-supervision with Francisco M. Couto, Universidade de Lisboa
- **Topic:** Creation of an ontology of epidemiology.
- **Topic:** The work consisted in using the textual descriptions in a Dictionary of Epidemiology to find relationships between the concepts and thus create an ontology of epidemiology.

## A.4 Knowledge transfer

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None

## A.5 Grants, Honors and Awards

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### Grants

**2010–2015** PhD Grant by Fundação para a Ciência e Tecnologia  
**2009–2010** Pre-PhD grant by LASIGE

### Awards

**2014** BPH Travel Award by the VPH Institute  
**2006** Best Student in 1<sup>st</sup> year by Universidade de Lisboa  
**2006** Merit Scholarship by Universidade de Lisboa

### Conference organisation

#### 6<sup>th</sup> International Conference on Biomedical Ontology (ICBO)

2015

- **Description:** The sixth International Conference on Biomedical Ontology (ICBO) was held in Lisbon in 2015 (<http://icbo2015.fc.ul.pt>). This prestigious and well-attended conference gathered multi-disciplinary researchers at the Faculdade de Ciências da Universidade de Lisboa to present and discuss the latest research breakthroughs in exploring ontologies in a biomedical and clinical context.
- **Position:** Part of the organization: I was Workshop & tutorial chair and Session chair. I was also part of the Proceedings team, having compiled the Proceedings, and organized the submission to CEUR-WS.

### Reviewer

**2016** Journal of Biomedical Semantics  
**2015** Journal of Biomedical Semantics  
Journal of Epidemiology & Community Health  
**2013** Oxford Bioinformatics  
**2012** Oxford Bioinformatics

### Program Committee

**2015** Bioinformatics Open Days  
Conference on Practical Applications of Computational Biology & Bioinformatics  
**2014** Conference on Practical Applications of Computational Biology & Bioinformatics

### Other reviewer activities

**2015** Portuguese Conference on Artificial Intelligence  
**2014** International Conference on Data Integration & Life Sciences

## B Pedagogical performance

### B.1 Teaching

#### Courses

Table 2: This table shows my contribution to courses from the Departamento de Informática da Faculdade de Ciências da Universidade de Lisboa.

**Legend:** *PL*: Prática de Laboratório (Practice lesson); *TP*: Teórico-Prática (Theoretical & Practical lesson); *PAC*: Professor Auxiliar convidado (Invited Auxiliary Teacher)

Course	Position	2015–2016
ASW	PAC	<b>PL</b>
IC	PAC	<b>TP &amp; PL</b>
ITW	PAC	<b>TP &amp; PL</b>
PD	PAC	<b>TP</b>
Prog. I	PAC	<b>TP</b>

#### ASW (Aplicações e Serviços na Web)

2015–2016

- **Degree:** Licenciatura em Tecnologias de Informação (2<sup>nd</sup> year)
- **My lectures:** PL
- **Topics:**
  - Web application characteristics and features;
  - The development process of web applications;
  - Introduction to the main server-side web technologies: resource addressing, protocols and general architecture;
  - The various data transfer formats (XML, JSON, etc.) and related technologies;
  - Introduction to Web Services and Semantic Web.
- **My contributions:** I produced a guide to explain HTML forms, along with JavaScript access to the content of the forms and the content of the GET arguments passed with the URL. The goal was to teach students how to deal with user-input, and how it travels through the network between web pages.

#### IC (Interacção com Computadores)

2015–2016

- **Degree:** Licenciatura em Tecnologias de Informação (2<sup>nd</sup> year)
- **My lectures:** TP & PL
- **Topics:**
  - Introduction to Human-Computer Interaction (HCI);
  - The foundations of HCI: Human and technological aspects;
  - The design process: user centred design, interaction design basics, guidelines for interaction design, and evaluation techniques;
  - Models and theories: cognitive models, task analysis, dialogue notations
- **My contributions:** I contributed to the class materials produced, including one of the project descriptions, which consisted in the development of an application based on web technologies that assisted people managing their kitchen (kitchen utensils, silverware, food etc.). The goal was to teach students how to implement an application following the principles of human-computer interaction.

#### ITW (Introdução às Tecnologias da Web)

2015–2016

- **Degree:** Licenciatura em Tecnologias de Informação (1<sup>st</sup> year)
- **My lectures:** TP & PL
- **Topics:**
  - Internet and web history;
  - Basic web concepts: the architecture, models, protocols, user agents, and transactions;
  - Text and Hypertext Mark-up;
  - Image Mark-up;



- Introduction to HTML;
- Introduction to Cascading Style Sheets (CSS);
- JavaScript concepts: flow control, data structure and objects;
- Processing user input in HTML Forms;
- Introduction to HTML5.
- **My contributions:** I contributed with several HTML and CSS exercises for the students to complete in their TP lectures. The goal was to introduce the students to the development of web pages with HTML and CSS.

## PD (Processamento de Dados)

2015–2016

- **Degree:** Licenciatura em Biologia (2<sup>nd</sup> year)
- **My lectures:** TP – invited teacher
- **Topics:**
  - Introduction to data processing;
  - Introduction to Python (data types and data structures);
  - Introduction to Regular Expressions;
  - Introduction to Biomedical Web Services;
  - Database management systems.
- **My contributions:** I created all the practice class materials – a tutorial to guide students in their project, which was to process a large collection of protein and metabolic information extracted from widely-known biomedical web services, including protein sequences and metabolic pathway data. The goal was to teach students how to process biology-related data with a programming language and an underlying database.

## Prog I (Programação I a outras licenciaturas)

2015–2016

- **Degree:** Several B.Sc and M.Sc offered by Faculdade de Ciências da Universidade de Lisboa
- **My lectures:** TP
- **Topics:**
  - Computation: computability and Turing machines;
  - Algorithms: exhaustive search, approximation search and bisection search;
  - Programming methods: attribution and verification, decision, iteration and recursion, abstraction and specification, cloning;
  - Programming languages: expressions and types, precedence and associativity, functions, scope, libraries and modules;
  - Data structures: sequences, tuples, lists and dictionaries;
  - Files;
  - Software development: reading and writing, documentation, assertions and exceptions, test and debugging.
- **My contributions:** I contributed to the practice class materials by suggesting new exercises and modifications to existing ones.

## Pedagogical surveys

Table 3: Pedagogical survey results. Students could either not answer each question or answer from 1 (strong disagreement) to 4 (strong agreement). Results for each question show the average for the students that answered the question.

**Legend:** Q1: Did the professor lecture with clarity? Q2: Did the professor answer questions with clarity? Q3: Was the professor available for outside-of-class contact & support? Q4: Was there a good pedagogical relation between professor and students? Q5: What is your global appreciation of the professor? PL: Prática de Laboratório (Practice lesson); TP: Teórico-Prática (Theoretical & Practical lesson); \*: Course still ongoing, results unavailable.

Question	2015–2016					
	1 <sup>st</sup> semester			2 <sup>nd</sup> semester		
	IC		Prog I	ASW		ITW
	TP	PL		PL	TP	PL
Q1	3.86	3.86	3.69	*	*	*
Q2	3.86	3.80	3.72	*	*	*
Q3	3.70	3.71	3.62	*	*	*
Q4	3.93	3.86	3.80	*	*	*
Q5	3.87	3.88	3.79	*	*	*

## Teaching materials

### ASW (Aplicações e Serviços na Web)

- I produced a guide to explain HTML forms, along with JavaScript access to the content of the forms and the content of the GET arguments passed with the URL. The goal was to teach students how to deal with user-input, as well as to introduce them to the idea of information flowing with the HTTP request from a form to the next web page. This material was made available to the students through moodle.

### ITW (Introdução às Tecnologias da Web)

- I contributed with several HTML and CSS exercises for the students to complete in their TP lectures. The goal was to introduce the students to the development of web pages with HTML and CSS. This material was made available to the students through moodle.

### PD (Processamento de Dados)

- I created all the practice class materials – a tutorial to guide students in their project, which was to process a large collection of protein and metabolic information extracted from widely-known biomedical web services, including protein sequences and metabolic pathway data. The goal was to teach students how to process biology-related data with a programming language and an underlying database. This material was made using google-docs and published as an HTML page, which was made available to the students through moodle.

## B.2 Jury and Examinations

### Invitation to M.Sc in Bioinformatics and Computational Biology

2016

- **Student:** Samuel Viana
- **Supervisors:** Daniel Faria (Instituto Gulbenkian de Ciência) and Catia Pesquita (Universidade de Lisboa)
- I was formally invited by the supervisors; the examination is expected to happen in July.

## B.3 Teaching-related activities

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### Invited participation in courses

#### AW (Aplicações na Web)

2014–2015

- **Degree:** Mestrado em Engenharia Informática
- I presented a lecture on Semantic Web (SW) with the following topics:
  - The problem of ambiguity that SW tries to solve;
  - Rule-based inference;
  - RDF statements;
  - Several of the SW languages (RDF, OWL, SPARQL);
  - Objects vs. Classes vs. Instances
  - An introduction to several of the layers of the SW, including URIs, XML, RDF, Ontologies and Rules;
  - Real-world examples of SW in action: Semantic wikis, FOAF project, RDFa, hCalendar, Linked Data Project.

#### Bioinformatics & Computational Modelling

2013–2014

- **Degree:** PhD Program in Biological Systems – Functional & Integrative Genomics
- I presented a practical lecture on Bioinformatics, specifically on the use of Python in biomedical data processing. This included:
  - Basic python datatypes and functions;
  - Introduction to BioPython, a package with access to several functions dedicated (i) to biological data processing and (ii) to widely-known biomedical web services;
  - Exercises directed at learning the inners of BioPython, specifically to process protein sequences (using web services to access the SwissProt database and the BLAST algorithm)
  - Introduction to the Gene Ontology and Semantic Similarity

#### Ontologias Aplicadas às Ciências

2013–2014

- **Degree:** M.Sc. class available to several M.Sc. students at FCUL
- I collaborated on the practical classes by designing a project and supervising a group of students in implementing the project's specifications. The projects were designed so that students would develop an intuition about ontology development, ontology matching, and semantic similarity. Two of these projects resulted in publications at national and international level.

#### Bioinformática

2011–2012

- **Degree:** B.Sc. class available to several B.Sc. students at FCUL
- I was a teaching assistant on the practical classes by invitation of the course's head professor. I created a project specification based on ontology development and text-mining, and supervised the students in their implementations.

## C University Mission

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### C.1 Participation in socially relevant projects

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#### FCUL Open Day

2015

- I helped devise a game-based activity that used two Bioinformatics tools developed at LASIGE to show students the current challenges in exploring textual data in the biomedical domain and how semantic-based tools can assist the research process.

#### Semana da Ciência em Movimento

2014

- I helped create and direct the BioIn4matix Run, a peddypaper activity that took 50 secondary school students throughout a physical journey around the FCUL Campus as well as a conceptual journey several around Bioinformatics research steps.

### C.2 Scientifically relevant projects and organizations of scientific, professional or cultural interest

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#### 2<sup>as</sup> Jornadas GIMGAS (Grupo de Internos de Medicina Geral e Familiar de Almada-Seixal)

2015

- This is a conference organized by internal General Practitioner residents from Almada and Seixal, open to interns and recently specialized doctors. I was responsible for creating a registration website for the conference, including managing the registration for the several workshops presented at the conference. The conference was attended by more than 100 doctors.

## D Full publication list

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### D.1 Journal papers

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- [1] Andre Lamurias, **João D. Ferreira**, and Francisco M. Couto. "Improving chemical entity recognition through h-index based semantic similarity." In: *Journal of Cheminformatics* 7.Suppl 1 Text mining for chemistry and the CHEMDNER track (Jan. 2015), S13. ISSN: 1758-2946. DOI: [10.1186/1758-2946-7-S1-S13](https://doi.org/10.1186/1758-2946-7-S1-S13).
- [2] Andre Lamurias, **João D. Ferreira**, and Francisco M. Couto. "Identifying interactions between chemical entities in biomedical text". In: *Journal of Interactive Bioinformatics (JIB)* 11.3 (2014), pp. 1–18. ISSN: 1613-4516. DOI: [10.2390/biecoll-jib-2014-247](https://doi.org/10.2390/biecoll-jib-2014-247).
- [3] Catia Pesquita, **João D. Ferreira**, Francisco M. Couto, and Mário J. Silva. "The epidemiology ontology: an ontology for the semantic annotation of epidemiological resources." In: *Journal of Biomedical Semantics* 5.1 (2014), p. 4. ISSN: 2041-1480. DOI: [10.1186/2041-1480-5-4](https://doi.org/10.1186/2041-1480-5-4).
- [4] **João D. Ferreira**, Janna Hastings, and Francisco M. Couto. "Exploiting disjointness axioms to improve semantic similarity measures." In: *Bioinformatics* 29.21 (2013), pp. 2781–2787. ISSN: 1367-4811. DOI: [10.1093/bioinformatics/btt491](https://doi.org/10.1093/bioinformatics/btt491).
- [5] **João D. Ferreira**, Daniela Paolotti, Francisco M. Couto, and Mário J. Silva. "On the usefulness of ontologies in epidemiology research and practice." In: *Journal of epidemiology and community health* (Nov. 2012), pp. –3. ISSN: 1470-2738. DOI: [10.1136/jech-2012-201142](https://doi.org/10.1136/jech-2012-201142).
- [6] **João D. Ferreira** and Francisco M. Couto. "Semantic Similarity for Automatic Classification of Chemical Compounds". In: *PLoS Computational Biology* 6.9 (Sept. 2010). Ed. by John B. O. Mitchell, e1000937. ISSN: 1553-7358. DOI: [10.1371/journal.pcbi.1000937](https://doi.org/10.1371/journal.pcbi.1000937).

### D.2 Conferences

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- [7] Andre Lamurias, **João D. Ferreira**, and Francisco M. Couto. "Chemical named entity recognition: Improving recall using a comprehensive list of lexical features". In: *8th International Conference on Practical Applications of Computational Biology & Bioinformatics*. Vol. 294. AISC. 2014, pp. 253–260. DOI: [10.1007/978-3-319-07581-5\\_30](https://doi.org/10.1007/978-3-319-07581-5_30).
- [8] **João D. Ferreira**, Catia Pesquita, Mário J. Silva, and Francisco M. Couto. "Digital preservation of epidemic resources: coupling metadata and ontologies". In: *International Conference on Preservation of Digital Objects*. Ed. by José Borbinha, Michael Nelson, and Steve Knight. 2013. URL: [http://xldb.di.fc.ul.pt/xldb/publications/Ferreira.etal:DigitalPreservationOf:2013\\_document.pdf](http://xldb.di.fc.ul.pt/xldb/publications/Ferreira.etal:DigitalPreservationOf:2013_document.pdf).
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### D.3 Thesis

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- [15] **João D. Ferreira**. “Semantic Similarity Across Biomedical Ontologies”. PhD Thesis. Universidade de Lisboa, 2016.
- [16] **João D. Ferreira**. “Structural and semantic similarity metrics for chemical compound classification”. Master’s Dissertation. Universidade de Lisboa, 2010.

### D.4 Other

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- [17] **João D. Ferreira**, David S. Batista, Francisco M. Couto, and Mário J. Silva. *The Geo-Net-PT / Yahoo ! GeoPlanet TM concordance*. Tech. rep. October. Departamento de Informática – Faculdade de Ciências - Universidade de Lisboa, 2010. DOI: **DOI : 10455/6677**.
- [18] Tiago Grego, **João D. Ferreira**, Catia Pesquita, Hugo Bastos, Diogo Vila Viçosa, João M. Freire, and Francisco M. Couto. *Chemical and Metabolic Pathway Semantic Similarity*. Tech. rep. Department of Informatics, Faculty of Sciences, University of Lisbon, 2010. DOI: **DOI : 10455/3335**.

## E Education

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### Ph.D. in Computer Science – Bioinformatics

*Dec. 2010 – Jan. 2016*

UNIVERSIDADE DE LISBOA

- **Thesis:** Semantic Similarity Across Biomedical Ontologies
- **Supervisor:** Prof. Dr Francisco M. Couto
- **Topic:** I developed several algorithms to measure similarity between biomedical resources annotated with concepts retrieved from biomedical ontologies (community-accepted standard representations of knowledge), as well as methodologies to validate them (based mainly on machine-learning). Along the research, I implemented several pieces of software related to knowledge-representation, semantic web and semantic similarity.
- **Grade:** Aprovado com Distinção e Louvor

### M.Sc. in Biochemistry

*Sep. 2008 – June 2010*

UNIVERSIDADE DE LISBOA

- **Thesis:** Structural and semantic similarity metrics for chemical compound similarity
- **Supervisor:** Prof. Dr Francisco M. Couto
- **Topic:** I developed a hybrid similarity measure to compare molecules. The measure was used as the basis of a machine-learning algorithm to predict biological properties of chemical compounds.
- **Grade:** 19 out of 20

### B.Sc. in Biochemistry

*Sep. 2005 – June 2008*

UNIVERSIDADE DE LISBOA

- **Discrimination of some grades:**
  - Análise e Tratamento de Dados em Bioquímica [*Data Analysis and Processing in Biochemistry*]: 19
  - Bioquímica Computacional [*Computational Biochemistry*]: 19
  - Simulação Computacional [*Computational Simulation*]: 20
- **Final grade:** 18 out of 20