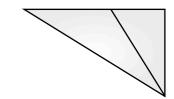


Curriculum Vitae

Florian Leitner, PhD

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Personal Profile and Expertise

- Content analyses based on information retrieval and extraction.
- Applied machine learning for data and text mining.
- Specialization on molecular biology and bioinformatics.
- Distributed web services, computing, and web development expert.
- Excellent communication, planning, and organization skills.

Professional Career Summary

Post-doc, National Cancer Research Centre (CNIO), Madrid, Spain

since Mar. 2013 • Prof. Alfonso Valencia Herrera

- Co-organization and evaluation of international scientific community challenges on information extraction for molecular biology; www.biocreative.org (BioCreative IV: ChemDNER - chemical entities)
- Development of Apache UIMA-based text mining/information extraction pipeline in Java (txtfnnl) with a PostgreSQL/Python support library (libfnl) for Gene Normalization and Transcription Regulation Events

PhD Student, National Cancer Research Centre (CNIO), Madrid, Spain

since Aug. 2006 • Prof. Alfonso Valencia Herrera

- Co-organization and evaluation of three international scientific community challenges on information extraction for molecular biology: BioCreative II and III, and acting main organizer of BioCreative II.5 (www.biocreative.org).
- Design and implementation of the first distributed meta-service for natural language processing in biology, the BioCreative Meta-Server (Python and JavaScript; placed third in the 2009 ICWS Services Cup after submissions from IBM and SAP).

MSc Student, Institute for Molecular Pathology (IMP), Vienna, Austria

Jul. 2003 - Jul. 2006 • Prof. Frank Eisenhaber (now director of the A*STAR Bioinformatics Institute, Singapore)

- Data acquisition and import from biological resources (parser development) and data visualization (on Apache/Struts) for an in-hose bioinformatics framework ("Annotator").
- Data indexing and query interface for the Annotator framework with Lucene.

Developer, CSO Lanifex GmbH., Vienna, Austria

Mar. 2003 - Jun. 2003 • Paul Gillingwater, MBA (now CEO of Anagram Media e.U., U.S.A.)

- Development of a specialized VPN network for the Austrian National Bank (particular contributions: Linux kernel modifications and adapting the Mozilla framework as a GUI for the thin client devices).
- Contributions to the Vienna United Nations HQ intranet framework (MySQL, PHP).

Visiting Student, European Media Labs (EML), Heidelberg, Germany

Jul. 2002 - Feb. 2003 • Rebecca Wade, PhD

- Development of a visualization plugin for annotations of protein structures (Java).
- Installation and configuration of Python libraries for structural biocomputing.

Undergraduate Student, Novartis Research Institute, Vienna, Austria

Jul. 2001 - Jun. 2002 • Markus Jaritz, PhD (now head of the GEN-AU Bioinformatics Division, Austria)

• Development of a Perl-based framework for Bioinformatics (Non-Disclosure Agreement).

Product/Marketing Manager, ProDesign, Natters bei Innsbruck, Austria

Sep. 1997 - Jul. 2000 • Herbert Hofbauer, CEO

- International production (China, Israel, South Korea) and sales management for a Tyrolean paraglider producer.
- Organization of marketing (sports) events and advertisements.
- Contributions to the in-house production/warehouse management tool (VisualBasic).

Executive Manager, Leitner Siebdruck, Völs bei Innsbruck, Austria

Jul. 1996 - Aug. 1997 • Harald Leitner, CEO

- Production and sales management for a Tyrolean printing company.
- Calculations, offers, and bids on tenders for commercial signs, markings, and advertisements.
- Human resource management.

Selected IT Experience

- Over a decade of professional work on POSIX (UNIX, Linux, BSD, Cygwin, Solaris) systems.
- ▶ Performing content analyses with Tika, UIMA, GATE, NLTK, and LingPipe.
- Machine learning with Mahout, SciPy, R, Orange, and several other, specialized libraries.
- ▶ Indexing Big Data with Lucene/Solr, Sphinx, and Xapian.
- Distributed computing experience with Hadoop/HDFS.
- Designing asynchronous messaging solutions with ZeroMQ and RabbitMQ.
- Data management skills with PostgresSQL, CouchDB, Riak, Cassandra, and HBase.
- ▶ Web development with Apache/Struts, Django, Rails, and Node.js.
- ▶ Deep knowledge of Python programming and test-driven RAD.
- Expertise in the design and implementation of efficient Java.
- Excellent C programming, Perl hacking, and Shell scripting skills; budding R mastery.
- Well-grounded programming experience in server- and client-side JavaScript.
- ▶ Applied understanding of (Common) Lisp, Erlang, Ruby, C++, and Scala.

Academic Titles

March 2012 - PhD, Molecular Biology, with distinction Autonomous University of Madrid (UAM), Spain

July 2006 - MSc, Molecular Biology, with distinction Vienna University, Austria

First-author Publications

1. Leitner, F., Chatr-aryamontri, A., Mardis, S. A., Ceol, A., Krallinger, M., Licata, L., Hirschman, L., Cesareni, G. & Valencia, A.

The FEBS Letters/BioCreative II.5 experiment: making biological information accessible *Nature Biotechnology*, 28(9), pp. 897–9, 2010

2. Leitner, F., Krallinger, M., Cesareni, G. & Valencia, A.

The FEBS Letters SDA corpus: a collection of protein interaction articles with high quality annotations for the BioCreative II.5 online challenge and the text mining community FEBS Letters, 584(19), pp. 4129–30, 2010

3. Leitner, F., Mardis, S. A., Krallinger, M., Cesareni, G., Hirschman, L. A. & Valencia, A. An overview of BioCreative II.5

IEEE/ACM Trans Comput Biol Bioinform, 7(3), pp. 385–99, 2010

4. Leitner, F., Krallinger, M., Rodriguez-Penagos, C., Hakenberg, J., Plake, C., Kuo, C.-J., Hsu, C.-N., Tsai, R. T.-H., Hung, H.-C., Lau, W. W., Johnson, C. A., Saetre, R., Yoshida, K., Chen, Y. H., Kim, S., Shin, S.-Y., Zhang, B.-T., Baumgartner, Jr, W. A., Hunter, L., Haddow, B., Matthews, M., Wang, X., Ruch, P., Ehrler, F., Ozgür, A., Erkan, G., Radev, D. R., Krauthammer, M., Luong, T., Hoffmann, R., Sander, C. & Valencia, A.

Introducing meta-services for biomedical information extraction Genome Biology, 9 Suppl 2, S6, 2008

5. Leitner, F. & Valencia, A.

A text-mining perspective on the requirements for electronically annotated abstracts \it FEBS Letters, 582(8), pp. 1178-81, 2008

Other Accomplishments

- 2012 Programa Integral de Lectura Madrid (a 6 months long speed reading course in Spanish) accredited final reading speed: 2000 words/minute
- 2012 Stanford Online Natural Language Processing Course completed with 88.3/100 total score
- 2009 IEEE Services-I World Conference "Services Cup"
 3rd place behind competitors from SAP and IBM

Co-author Publications

- Salgado D., Krallinger M., Depaule M., Drula E., Tendulkar A., Leitner F., Valencia A., Marcelle C. MyMiner: a web application for computer-assisted biocuration and text annotation. Bioinformatics, 2012, [advance access]
- Krallinger, M., Leitner, F., Vazquez, M., Salgado, D., Marcelle, C., Tyers, M., Valencia, A., Chatr-aryamontri, A. How to link ontologies and protein-protein interactions to literature: text-mining approaches and the BioCreative experience Database, 2012, bas017, 2012
- 3. Krallinger, M., Vazquez, M., <u>Leitner</u>, F., Salgado, D., Chatr-Aryamontri, A., Winter, A., Perfetto, L., Briganti, L., Licata, L., Iannuccelli, M., Castagnoli, L., Cesareni, G., Tyers, M., Schneider, G., Rinaldi, F., Leaman, R., Gonzalez, G., Matos, S., Kim, S., Wilbur, W., Rocha, L., Shatkay, H., Tendulkar, A. V., Agarwal, S., Liu, F., Wang, X., Rak, R., Noto, K., Elkan, C., Lu, Z., Dogan, R., Fontaine, J.-F., Andrade-Navarro, M. A. & Valencia, A. The protein-protein interaction tasks of BioCreative III: classification/ranking of articles and linking bio-ontology concepts to full text BMC Bioinformatics, 12(Suppl 8), S3, 2011
- 4. Vazquez, M., Krallinger, M., <u>Leitner</u>, F. & Valencia, A. **Text mining for drugs and chemical compounds: methods, tools and applications** *Molecular Informatics*, 30(6–7), pp. 506-519, 2011
- Krallinger, M., Tendulkar, A., <u>Leitner</u>, F., Chatr-aryamontri, A. & Valencia, A.
 The PPI affix dictionary (PPIAD) and biomethod lexicon: importance of affixes and tags for recognition of entity mentions and experimental protein interactions
 BMC Bioinformatics, 11(Suppl 5), O1, 2010
- Krallinger, M., <u>Leitner</u>, F., Rodriguez-Penagos, C. & Valencia, A.
 Overview of the protein-protein interaction annotation extraction task of BioCreative II Genome Biology, 9(Suppl 2), S4, 2008
- 7. Krallinger, M., Morgan, A., Smith, L., <u>Leitner</u>, F., Tanabe, L., Wilbur, J., Hirschman, L. & Valencia, A. **Evaluation of text-mining systems for biology: overview of the second BioCreative community challenge** *Genome Biology*, 9(Suppl 2), S1, 2008
- 8. Gabdoulline, R.R., Hoffmann, R., <u>Leitner</u>, F. & Wade, R. C. **ProSAT: functional annotation of protein 3D structures** *Bioinformatics*, 19 (13), p. 1723, 2003

Conference Proceedings (see Presentations for First Author Proceedings)

- Krallinger, M., Vazquez, M., <u>Leitner</u>, F. & Valencia A. Results of the BioCreative III (Interaction) Article Classification Task. Proceedings of the BioCreative III Challenge Evaluation Workshop, pp. 17-23, 2010
- Krallinger, M., Vazquez, M., <u>Leitner</u>, F. & Valencia A.
 Results of the BioCreative III Interaction Method Task.
 Proceedings of the BioCreative III Challenge Evaluation Workshop, pp. 9-16, 2010
- Starlinger, J., <u>Leitner</u>, F., Valencia, A. & Leser, U.
 SOA-based integration of text mining services
 2009 World Conference on Services (IEEE SERVICES), pp. 99-106, 2009
- 4. Krallinger, M., <u>Leitner</u>, F. & Valencia, A. **Assessment of the second BioCreative PPI task: automatic extraction of protein-protein interactions** *Proceedings of the second BioCreative Challenge Evaluation Workshop*, pp. 41-54, 2007
- 5. Krallinger, M., Rodriguez-Penagos, C., <u>Leitner</u>, F. & Valencia, A. **Exploiting text mining of biological annotations to extract protein interactions for the Semantic web.** *Seventh Intl. Workshop on Network Tools and Applications in Biology* (NETTAB), pp. 47-53, 2007

Book Chapters

1. Krallinger, R., Leitner, F. & Valencia, A.

Analysis of Biological Processes and Diseases Using Text Mining Approaches

Editor: Matthiesen, R.

Bioinformatics Methods in Clinical Research, Methods in Molecular Biology 593, pp. 341-382, 2010

2. Leitner, F., Hoffmann, R. & Valencia, A.

Biological Knowledge Extraction

Editor: Krawetz, S.

Bioinformatics for Systems Biology, Part V, pp. 413-433, 2009

Presentations

1. Mining cis-Regulatory Transcription Networks from Literature BioLINK @ ISMB/ECCB, 2013, Berlin, Germany (in July)

2. Integration across genomics and medicine

Panel discussion, BioLINK @ ISMB/ECCB, 2011, Vienna, Austria

3. Annotating scientific manuscripts and the role of text-mining

FEBS J Board Meeting, 2010, Cambridge, UK

4. Making biological information accessible

CNIO Progress Reports, June 2010, Madrid, Spain

5. On interoperability in text mining

BioCreative III Workshop, 2010, Bethesda, USA

6. The BioCreative II.5 evaluation results

BioCreative II.5 Workshop, 2009, Madrid, Spain

7. The BCMS in the challenge and as a collaborative platform

BioCreative II.5 Workshop, 2009, Madrid, Spain

8. Comparative community assessments for applied biomedical text mining - BioCreative II challenge and meta-services

ISMB/ECCB, 2009, Stockholm, Sweden

9. The BioCreative II.5 online challenge

BioLINK @ ISMB/ECCB, 2009, Stockholm, Sweden

10. The BioCreative Meta-Server platform

SMBM, 2008, Turku, Finland

11. The interaction pair & interaction method sub-tasks evaluation

BioCreative II Workshop, 2007, Madrid, Spain