

1. Record Nr.	UNISA996556970303316
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Titolo	Intonation Between Phrasing and Accent : Spanish and Quechua in Huari
Pubbl/distr/stampa	Berlin/Boston : , : Walter de Gruyter GmbH, , 2023 ©2024
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Edizione	[1st ed.]
Descrizione fisica	1 online resource (636 pages)
Collana	Linguistica Latinoamericana Series ; ; v.7
Disciplina	498.32316
Soggetti	FOREIGN LANGUAGE STUDY / Spanish
Lingua di pubblicazione	Inglese
Formato	Materiale a stampa
Livello bibliografico	Monografia
Nota di contenuto	Frontmatter -- Acknowledgements -- Contents -- List of figures -- List of tables -- List of symbols and abbreviations -- Huari Quechua morphological glosses -- Glosses from other languages cited in this work -- 1 Introduction -- 2 Data & general methods -- 3 Theoretical background and literature review -- 4 Refined research questions -- 5 Huari Spanish -- 6 Huari Quechua -- 7 Synthesis -- Appendix A – Weblinks to the audio files -- Appendix B – Maptask maps -- Appendix C – Praat scripts -- References
Sommario/riassunto	Are our concepts from prosodic typology, like word stress, pitch accent, head-/edge-prominence, really that tightly linked to individual languages? How are meanings often signaled via intonation in European languages, like information structure and sentence type, expressed in communicative acts between speakers who are bilingual in such a European language, Spanish, and one in which many of these meanings are expressed by morphology, Quechua? Based on semi-spontaneous dialogical elicitation data in both Spanish and Quechua gathered via fieldwork in the bilingual community of Huari, Peru, this work provides some challenging answers to these questions. Besides being the first detailed description of the prosody of a Central Quechuan language, it provides an in-depth study of the intonational systems and prosodic structures of the two languages and shows that their variation spaces overlap to a large extent, in turns exhibiting or not exhibiting evidence of word stress, pitch accents, lexical pitch accents in loanwords, and

head- or edge-prominence.

2. Record Nr.	UNINA9910778260903321
Titolo	Pacific Symposium on Biocomputing 2007 [[electronic resource]] : Maui, Hawaii, 3-7 January 2007 // edited by Russ B. Altman ... [et al.]
Pubbl/distr/stampa	Hackensack, N.J., : World Scientific, c2007
ISBN	1-281-91203-4 9786611912031 981-277-243-X
Descrizione fisica	xvi, 507 p. : ill
Altri autori (Persone)	AltmanRuss
Disciplina	570.15
Soggetti	Biology - Computer simulation Biology - Mathematical models Molecular biology - Computer simulation Molecular biology - Mathematical models
Lingua di pubblicazione	Inglese
Formato	Materiale a stampa
Livello bibliografico	Monografia
Note generali	Bibliographic Level Mode of Issuance: Monograph
Nota di bibliografia	Includes bibliographical references.
Nota di contenuto	Preface -- Protein interactions and diseases. Session introduction / Maricel Kann ... [et al.]. Graph kernels for disease outcome prediction from protein-protein interaction networks / Karsten M. Borgwardt ... [et al.]. Chalkboard: ontology-based pathway modeling and qualitative inference of disease mechanisms / Daniel L. Cook, Jesse C. Wiley, and John H. Gennari. Mining gene-disease relationships from biomedical literature weighting protein-protein interactions and connectivity measures / Graciela Gonzalez ... [et al.]. Predicting structure and dynamics of loosely-ordered protein complexes: influenza hemagglutinin fusion peptide / Peter M. Kasson and Vijay S. Pande. Protein interactions and disease phenotypes in the ABC transporter superfamily / Libusha Kelly, Rachel Karchin, and Andrej Sali. LTHREADER: prediction of Ligand-receptor interactions using localized threading / Vinay Pulim, Jadwiga Bienkowska, and Bonnie Berger. Discovery of protein interaction networks shared by diseases / Lee Sam

... [et al.]. An iterative algorithm for metabolic network-based drug target identification / Padmavati Sridhar, Tamer Kahveci, and Sanjay Ranka. Transcriptional interactions during smallpox infection and identification of early infection biomarkers / Willy A. Valdivia-Granda, Maricel G. Kann, and Jose Malaga -- Computational approaches to metabolomics. Session introduction / David S. Wishart and Russell Greiner. Leveraging latent information in NMR spectra for robust predictive models / David Chang, Aalim Weljie, and Jack Newton. Bioinformatics data profiling tools: a prelude to metabolic profiling / Natarajan Ganesan, Bala Kalyanasundaram, and Mahe Velauthapllai. Comparative QSAR analysis of bacterial, fungal, plant and human metabolites / Emre Karakoc, S. Cenk Sahinalp, and Artem Cherkasov. BioSpider: a web server for automating metabolome annotations / Craig Knox ... [et al.]. New bioinformatics resources for metabolomics / John L. Markley ... [et al.]. Setup X - a public study design database for metabolomic projects / Martin Scholz and Oliver Fiehn. Comparative metabolomics of breast cancer / Chen Yang ... [et al.]. Metabolic flux profiling of reaction modules in liver drug transformation / Jeongah Yoon and Kyongbum Lee -- New frontiers in biomedical text mining. Session introduction / Pierre Zweigenbaum ... [et al.]. Extracting semantic predications from medline citations for pharmacogenomics / Caroline B. Ahlers ... [et al.]. Annotating genes using textual patterns / Ali Cakmak and Gultekin Ozsoyoglu. A fault model for ontology mapping, alignment, and linking systems / Helen L. Johnson, K. Bretonnel Cohen, and Lawrence Hunter. Integrating natural language processing with flybase curation / Nikiforos Karamanis Y. ... [et al.]. A stacked graphical model for associating sub-images with sub-captions / Zhenzhen Kou, William W. Cohen, and Robert F. Murphy. GeneRIF quality assurance as summary revision / Zhiyong Lu, K. Bretonnel Cohen, and Lawrence Hunter. Evaluating the automatic mapping of human gene and protein mentions to unique identifiers / Alexander A. Morgan ... [et al.]. Multiple approaches to fine-grained indexing of the biomedical literature / Aurelie Neveol ... [et al.]. Mining patents using molecular similarity search / James Rhodes ... [et al.]. Discovering implicit associations between genes and hereditary diseases / Kazuhiro Seki and Javed Mostafa. A cognitive evaluation of four online search engines for answering definitional questions posed by physicians / Hong Yu and David Kaufman -- Biodiversity informatics: managing knowledge beyond humans and model organisms. Session introduction / Indra Neil Sarkar. Biomediator data integration and inference for functional annotation of anonymous sequences / Eithon Cadag ... [et al.]. Absent sequences: nullomers and primes / Greg Hampikian and Tim Andersen. An anatomical ontology for amphibians / Anne M. Maglia ... [et al.]. Recommending pathway genes using a compendium of clustering solutions / David M. Ng, Marcos H. Woehrmann, and Joshua M. Stuart. Semi-automated XML markup of biosystematic legacy literature with the Goldengate editor / Guido Sautter, Klemens Bohm, and Donat Agosti -- Computational proteomics: high-throughput analysis for systems biology. Session introduction / William Cannon and Bobbie-Jo Webb-Robertson. Advancement in protein inference from shotgun proteomics using peptide detectability / Pedro Alves ... [et al.]. Mining tandem mass spectral data to develop a more accurate mass error model for peptide identification / Yan Fu ... [et al.]. Assessing and combining reliability of protein interaction sources / Sonia Leach ... [et al.]. Probabilistic modeling of systematic errors in two-hybrid experiments / David Sontag, Rohit Singh, and Bonnie Berger. Prospective exploration of biochemical tissue composition via imaging mass spectrometry guided by principal component analysis /

Raf Van de Plas ... [et al.] -- DNA-protein interactions: integrating structure, sequence, and function. Session introduction / Martha L. Bulyk ... [et al.]. Discovering motifs with transcription factor domain knowledge / Henry C. M. Leung, Francis Y. L. Chin, and Bethany M. Y. Chan. Ab initio prediction of transcription factor binding sites / L. Angela Liu and Joel S. Bader. Comparative pathway annotation with protein-DNA interaction and operon information via graph tree decomposition / Jizhen Zhao, Dongsheng Che, and Liming Cai.

3. Record Nr.	UNICAMPANIAVAN0275376
Titolo	Trends in Biomathematics: Chaos and Control in Epidemics, Ecosystems, and Cells : Selected Works from the 20th BIOMAT Consortium Lectures, Rio de Janeiro, Brazil, 2020 / Rubem P. Mondaini editor
Pubbl/distr/stampa	Cham, : Springer, 2021
Descrizione fisica	xi, 395 p. : ill. ; 24 cm
Soggetti	00B25 - Proceedings of conferences of miscellaneous specific interest [MSC 2020] 92-XX - Biology and other natural sciences [MSC 2020]
Lingua di pubblicazione	Inglese
Formato	Materiale a stampa
Livello bibliografico	Monografia