

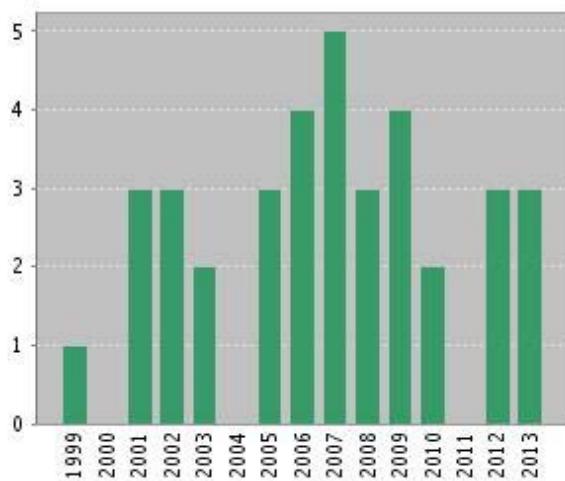
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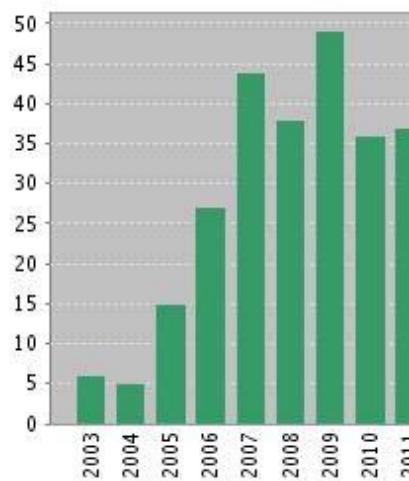
au=zwir i

Timespan=All years. Databases=SCI-EXPANDED, SSCI, A&amp;HCI, CPCI-S, CPCI-SSH.

Published Items in Each Year



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	2010	2011	2012	2013	2014	Total	Average Citations per Year
1.	36	37	43	38	0	338	30.73
1.	<b>Title:</b> Dissecting the PhoP regulatory network of Escherichia coli and Salmonella enterica <b>Author(s):</b> Zwir, I; Shin, D; Kato, A; et al. <b>Source:</b> PROCEEDINGS OF THE NATIONAL ACADEMY OF SCIENCES OF THE UNITED STATES OF AMERICA <b>Volume:</b> 102 <b>Issue:</b> 8 <b>Pages:</b> 2862-2867 <b>DOI:</b> 10.1073/pnas.0408238102 <b>Published:</b> FEB 22 2005	12	11	8	9	0	108
2.	<b>Title:</b> Linguistic modeling by hierarchical systems of linguistic rules <b>Author(s):</b> Cordon, O; Herrera, F; Zwir, I <b>Source:</b> IEEE TRANSACTIONS ON FUZZY SYSTEMS <b>Volume:</b> 10 <b>Issue:</b> 1 <b>Pages:</b> 2-20 <b>DOI:</b> 10.1109/91.983275 <b>Published:</b> FEB 2002	6	7	5	1	0	58
3.	<b>Title:</b> Evolution of a Bacterial Regulon Controlling Virulence and Mg <sup>2+</sup> Homeostasis <b>Author(s):</b> Perez, J. Christian; Shin, Dongwoo; Zwir, Igor; et al. <b>Source:</b> PLOS GENETICS <b>Volume:</b> 5 <b>Issue:</b> 3 <b>Article Number:</b> e1000428 <b>DOI:</b>	8	8	11	5	0	38

		2010	2011	2012	2013	2014	Total	Average Citations per Year
		36	37	43	38	0	338	30.73
10.1371/journal.pgen.1000428 <b>Published:</b> MAR 2009								
4.	<b>Title:</b> Analysis of differentially-regulated genes within a regulatory network by GPS genome navigation <b>Author(s):</b> Zwir, I; Huang, H; Groisman, EA <b>Source:</b> BIOINFORMATICS <b>Volume:</b> 21 <b>Issue:</b> 22 <b>Pages:</b> 4073-4083 <b>DOI:</b> 10.1093/bioinformatics/bti672 <b>Published:</b> NOV 15 2005	2	0	3	2	0	24	2.67
5.	<b>Title:</b> Linguistic modeling with hierarchical systems of weighted linguistic rules <b>Author(s):</b> Alcalá, R; Cano, JR; Cordon, O; et al. <b>Source:</b> INTERNATIONAL JOURNAL OF APPROXIMATE REASONING <b>Volume:</b> 32 <b>Issue:</b> 2-3 <b>Pages:</b> 187-215 <b>Article Number:</b> PII S0888-613X(02)00083-X <b>DOI:</b> 10.1016/S0888-613X(02)00083-X <b>Published:</b> FEB 2003	2	4	1	0	0	20	1.82
6.	<b>Title:</b> A hierarchical knowledge-based environment for linguistic modeling: models and iterative methodology <b>Author(s):</b> Cordon, O; Herrera, F; Zwir, I <b>Source:</b> FUZZY SETS AND SYSTEMS <b>Volume:</b> 138 <b>Issue:</b> 2 <b>Pages:</b> 307-341 <b>Article Number:</b> PII S0165-0114(02)00388-3 <b>DOI:</b> 10.1016/S0165-0114(02)00388-3 <b>Published:</b> SEP 1 2003	2	1	1	0	0	16	1.45
7.	<b>Title:</b> A hybrid promoter analysis methodology for prokaryotic genomes <b>Author(s):</b> Cotik, V; Zaliz, RR; Zwir, I <b>Source:</b> FUZZY SETS AND SYSTEMS <b>Volume:</b> 152 <b>Issue:</b> 1 <b>Pages:</b> 83-102 <b>DOI:</b> 10.1016/j.fss.2004.10.016 <b>Published:</b> MAY 16 2005	0	3	0	0	0	14	1.56
8.	<b>Title:</b> Profile analysis and prediction of tissue-specific CpG island methylation classes <b>Author(s):</b> Previti, Christopher; Harari, Oscar; Zwir, Igor; et al. <b>Source:</b> BMC BIOINFORMATICS <b>Volume:</b> 10 <b>Article Number:</b> 116 <b>DOI:</b> 10.1186/1471-2105-10-116 <b>Published:</b> APR 21 2009	3	0	3	4	0	11	2.20
9.	<b>Title:</b> Automated biological sequence description by genetic multiobjective generalized clustering <b>Author(s):</b> Zwir, I; Zaliz, RR; Ruspini, EH <b>Book Editor(s):</b> Valafar, F <b>Conference:</b> International Conference on Mathematics and Engineering Techniques in Medicine and Biological Sciences <b>Location:</b> LAS VEGAS, NEVADA <b>Date:</b> JUN 25-28, 2001 <b>Sponsor(s):</b> Comp Sci Res, Educ & Appl Press; Int Technol Inst; Korea Informat Proc Soc; World Acad Sci Informat Technol; PACT Corp; Comp Vis Res & Appl Tech; Java High Performance Comp Res Grp; Sundance Digital Signal Proc Inc <b>Source:</b> TECHNIQUES IN BIOINFORMATICS AND MEDICAL INFORMATICS <b>Book Series:</b> ANNALS OF THE NEW YORK ACADEMY OF SCIENCES <b>Volume:</b> 980 <b>Pages:</b> 65-82 <b>Published:</b> 2002	1	0	0	1	0	11	0.92

	2010	2011	2012	2013	2014	Total	Average Citations per Year
	36	37	43	38	0	338	30.73

10. **Title:** A Multiobjective Evolutionary Conceptual Clustering Methodology for Gene Annotation Within Structural Databases: A Case of Study on the Gene Ontology Database  
**Author(s):** Romero-Zaliz, Rocio C.; Rubio-Escudero, Cristina; Cobb, J. Perren; et al.  
**Source:** IEEE TRANSACTIONS ON EVOLUTIONARY COMPUTATION **Volume:** 12 **Issue:** 6 **Pages:** 679-701 **DOI:** 10.1109/TEVC.2008.915995 **Published:** DEC 2008

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