A pageranking based method for identifying characteristic genes of a disease

Hai D.T., Lee W., Thuy H.Q.
SYSlab., College of Technology, Vietnam National University, Hanoi 144, Xuan Thuy, Cau Giay, Ha Noi, Viet Nam; Network Research Lab. (NetLab), 507, A-San Science Building, Korea University, Seoul, South Korea

Abstract: Ranking is an importance task in the research field of complex biological networks, including gene networks [11]. In this article, we propose a method for ranking the genes causing a specific type of disease, special characteristics of samples under a particular condition. Genes with the highest scores may be considered as the key factors leading to the development of disease type in the studied samples. The experience results on the gene expression data of two subtypes of Leukemia cancer disease, i.e., ALB and ALT for B-cells and T-cells are respectively showed.
Author Keywords: Feature set; Gene networks; Rank of genes
Index Keywords: Chlorine compounds; Gene expression; B-cells; Biological networks; Feature set; Gene expression data; Gene networks; International conferences; Key factors; Particular condition; Rank of genes; T-cells; Genes

Year: 2008
Source title: Proceedings of 2008 IEEE International Conference on Networking, Sensing and Control, ICNSC
Art. No.: 4525457
Page : 1496-1499
Cited by: 1
Link: Scorpus Link
Correspondence Address: Hai, D. T.; SYSlab., College of Technology, Vietnam National University, Hanoi 144, Xuan Thuy, Cau Giay, Ha Noi, Viet Nam; email: haidt@vnu.edu.vn
Sponsors: IEEE Systems, Man and Cybernetics Society
Conference name: 2008 IEEE International Conference on Networking, Sensing and Control, ICNSC
Conference date: 6 April 2008 through 8 April 2008
Conference location: Sanya
Conference code: 72895
ISBN: 9.78E+12
DOI: 10.1109/ICNSC.2008.4525457
Language of Original Document: English
Abbreviated Source Title: Proceedings of 2008 IEEE International Conference on Networking, Sensing and Control, ICNSC
Document Type: Conference Paper
Source: Scopus

Authors with affiliations:

- Hai, D.T., SYSlab., College of Technology, Vietnam National University, Hanoi 144, Xuan Thuy, Cau Giay, Ha Noi, Viet Nam
- Lee, W., Network Research Lab. (NetLab), 507, A-San Science Building, Korea University, Seoul, South Korea
- Thuy, H.Q., SYSlab., College of Technology, Vietnam National University, Hanoi 144, Xuan Thuy, Cau Giay, Ha Noi, Viet Nam

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