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Research Areas

Computer Sciences, Engineering and Technology

Courses

Küresel Meseleler (GLB), Undergraduate, 2022 - 2023

Computational Genomics, Undergraduate, 2022 - 2023

Published journal articles indexed by SCI, SSCI, and AHCI

- I. **PriPath: identifying dysregulated pathways from differential gene expression via grouping, scoring, and modeling with an embedded feature selection approach**
Yousef M., ÖZDEMİR F., Jaber A., Allmer J., GÜNGÖR B.
BMC Bioinformatics, vol.24, no.1, 2023 (SCI-Expanded)
- II. **AMP-GSM: Prediction of Antimicrobial Peptides via a Grouping-Scoring-Modeling Approach**
Söylemez Ü. G., Yousef M., GÜNGÖR B.
Applied Sciences (Switzerland), vol.13, no.8, 2023 (SCI-Expanded)
- III. **Ensemble feature selection and classification methods for machine learning-based coronary artery disease diagnosis**
KOLUKISA B., Bakir-Gungor B.
Computer Standards and Interfaces, vol.84, 2023 (SCI-Expanded)
- IV. **miRdisNET: Discovering microRNA biomarkers that are associated with diseases utilizing biological knowledge-based machine learning**
Jabeer A., Temiz M., GÜNGÖR B., Yousef M.
Frontiers in Genetics, vol.13, 2023 (SCI-Expanded)
- V. **Review of feature selection approaches based on grouping of features**
Kuzudisli C., GÜNGÖR B., BULUT N., Qaqish B., Yousef M.
PeerJ, vol.11, 2023 (SCI-Expanded)
- VI. **microBiomeGSM: the identification of taxonomic biomarkers from metagenomic data using grouping, scoring and modeling (G-S-M) approach**
GÜNGÖR B., Temiz M., Jabeer A., Wu D., Yousef M.
Frontiers in Microbiology, vol.14, 2023 (SCI-Expanded)
- VII. **TextNetTopics Pro, a topic model-based text classification for short text by integration of semantic**

and document-topic distribution information

Voskerghian D., GÜNGÖR B., Yousef M.

Frontiers in Genetics, vol.14, 2023 (SCI-Expanded)

- VIII. **GeNetOntology: identifying affected gene ontology terms via grouping, scoring, and modeling of gene expression data utilizing biological knowledge-based machine learning**
Ersoz N. S., GÜNGÖR B., Yousef M.
Frontiers in Genetics, vol.14, 2023 (SCI-Expanded)
- IX. **A toolbox of machine learning software to support microbiome analysis**
Marcos-Zambrano L. J., López-Molina V. M., GÜNGÖR B., Frohme M., Karadzovic-Hadziabdic K., Klammsteiner T., Ibrahim E., Lahti L., Loncar-Turukalo T., Dharmo X., et al.
Frontiers in Microbiology, vol.14, 2023 (SCI-Expanded)
- X. **Invention of 3Mint for feature grouping and scoring in multi-omics**
ÜNLÜ YAZICI M., Marron J., GÜNGÖR B., Zou F., Yousef M.
Frontiers in Genetics, vol.14, 2023 (SCI-Expanded)
- XI. **Topological feature generation for link prediction in biological networks**
Temiz M., GÜNGÖR B., Güner Şahan P., COŞKUN M.
PeerJ, vol.11, 2023 (SCI-Expanded)
- XII. **Clinical and molecular evaluation of MEFV gene variants in the Turkish population: a study by the National Genetics Consortium**
DÜNDAR M., FAHRİOĞLU U., Yıldız S. H., Bakir-Gungor B., TEMEL Ş. G., AKIN H., ARTAN S., Cora T., ŞAHİN F. İ., DURSUN A., et al.
FUNCTIONAL & INTEGRATIVE GENOMICS, vol.22, no.3, pp.291-315, 2022 (SCI-Expanded)
- XIII. **The Determination of Distinctive Single Nucleotide Polymorphism Sets for the Diagnosis of Behcet's Disease**
IŞIK Y. E., GÖRMEZ Y., AYDIN Z., Bakir-Gungor B.
IEEE-ACM TRANSACTIONS ON COMPUTATIONAL BIOLOGY AND BIOINFORMATICS, vol.19, no.3, pp.1909-1918, 2022 (SCI-Expanded)
- XIV. **Prediction of Linear Cationic Antimicrobial Peptides Active against Gram-Negative and Gram-Positive Bacteria Based on Machine Learning Models**
Soylemez U. G., Yousef M., KESMEN Z., Buyukkiraz M. E., Bakir-Gungor B.
APPLIED SCIENCES-BASEL, vol.12, no.7, 2022 (SCI-Expanded)
- XV. **Inflammatory bowel disease biomarkers of human gut microbiota selected via different feature selection methods**
Bakir-Gungor B., Lar H. H., Jabeer A., NALBANTOĞLU Ö. U., Aran O., Yousef M.
PEERJ, vol.10, 2022 (SCI-Expanded)
- XVI. **miRModuleNet: Detecting miRNA-mRNA Regulatory Modules**
Yousef M., GÖY G., Bakir-Gungor B.
FRONTIERS IN GENETICS, vol.13, 2022 (SCI-Expanded)
- XVII. **Developing a label propagation approach for cancer subtype classification problem**
GÜNER P., Bakir-Gungor B., Coskun M.
TURKISH JOURNAL OF BIOLOGY, vol.46, no.2, pp.145-161, 2022 (SCI-Expanded)
- XVIII. **Enlightening the molecular mechanisms of type 2 diabetes with a novel pathway clustering and pathway subnetwork approach**
Bakir-Gungor B., ÜNLÜ YAZICI M., GÖY G., Temiz M.
TURKISH JOURNAL OF BIOLOGY, vol.46, no.4, pp.318-351, 2022 (SCI-Expanded)
- XIX. **Blockchain for genomics and healthcare: a literature review, current status, classification and open issues**
ADANUR DEDETÜRK B., SORAN A., Bakir-Gungor B.
PEERJ, vol.9, 2021 (SCI-Expanded)
- XX. **Discovering Potential Taxonomic Biomarkers of Type 2 Diabetes From Human Gut Microbiota via Different Feature Selection Methods**

- Bakir-Gungor B., Bulut O., Jabeer A., NALBANTOĞLU Ö. U., Yousef M.
FRONTIERS IN MICROBIOLOGY, vol.12, 2021 (SCI-Expanded)
- XXI. **miRcorrNet: machine learning-based integration of miRNA and mRNA expression profiles, combined with feature grouping and ranking**
Yousef M., GÖY G., Mitra R., Eischen C. M., Jabeer A., Bakir-Gungor B.
PEERJ, vol.9, 2021 (SCI-Expanded)
- XXII. **Statistical and Machine Learning Techniques in Human Microbiome Studies: Contemporary Challenges and Solutions**
Moreno-Indias I., Lahti L., Nedyalkova M., Elbere I., Roshchupkin G., Adilovic M., AYDEMİR Ö., Bakir-Gungor B., Santa Pau E. C., D'Elia D., et al.
FRONTIERS IN MICROBIOLOGY, vol.12, 2021 (SCI-Expanded)
- XXIII. **Application of Biological Domain Knowledge Based Feature Selection on Gene Expression Data**
Yousef M., Kumar A., Bakir-Gungor B.
ENTROPY, vol.23, no.1, 2021 (SCI-Expanded)
- XXIV. **Multi fragment melting analysis system (MFMAS) for one-step identification of lactobacilli**
KESMEN Z., Kilic O., GÖRMEZ Y., ÇELİK M., Bakir-Gungor B.
JOURNAL OF MICROBIOLOGICAL METHODS, vol.177, 2020 (SCI-Expanded)
- XXV. **In silico analyses and global transcriptional profiling reveal novel putative targets for Pea3 transcription factor related to its function in neurons**
KANDEMİR B., Dag U., Gungor B., Durasi I. M., Erdogan B., Sahin E., Sezerman U., KURNAZ I.
PLOS ONE, vol.12, no.2, 2017 (SCI-Expanded)
- XXVI. **Active Subnetwork GA: A Two Stage Genetic Algorithm Approach to Active Subnetwork Search**
ÖZİŞİK O., Bakir-Gungor B., DİRİ B., Sezerman O. U.
CURRENT BIOINFORMATICS, vol.12, no.4, pp.320-328, 2017 (SCI-Expanded)
- XXVII. **'Epistatic interactions between autoimmunity and genetic thrombophilia' Reply**
Bakir-Gungor B., REMMERS E. F., MEGURO A., MIZUKI N., KASTNER D. L., Gül A., Sezerman O. U.
EUROPEAN JOURNAL OF HUMAN GENETICS, vol.23, no.10, pp.1279-1280, 2015 (SCI-Expanded)
- XXVIII. **Identification of possible pathogenic pathways in Behcet's disease using genome-wide association study data from two different populations**
Bakir-Gungor B., REMMERS E. F., MEGURO A., MIZUKI N., KASTNER D. L., Gül A., Sezerman O. U.
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- XXIX. **PANOGA: a web server for identification of SNP-targeted pathways from genome-wide association study data**
Bakir-Gungor B., Egemen E., Sezerman O. U.
BIOINFORMATICS, vol.30, no.9, pp.1287-1289, 2014 (SCI-Expanded)
- XXX. **HomSI: a homozygous stretch identifier from next-generation sequencing data**
GÖRMEZ Z., Bakir-Gungor B., SAGIROGLU M. S.
BIOINFORMATICS, vol.30, no.3, pp.445-447, 2014 (SCI-Expanded)
- XXXI. **Identifying SNP targeted pathways in partial epilepsies with genome-wide association study data**
Bakir-Gungor B., BAYKAN B., Iseri S. U., Tuncer F. N., Sezerman O. U.
EPILEPSY RESEARCH, vol.105, pp.92-102, 2013 (SCI-Expanded)
- XXXII. **The Identification of Pathway Markers in Intracranial Aneurysm Using Genome-Wide Association Data from Two Different Populations**
GÜNGÖR B., Sezerman O. U.
PLOS ONE, vol.8, no.3, 2013 (SCI-Expanded)
- XXXIII. **A new methodology to associate snps with human diseases according to their pathway related context**
GÜNGÖR B., Sezerman O. U.
PLOS ONE, vol.6, no.10, 2011 (SCI-Expanded)
- XXXIV. **The Rat Genome Database 2009: Variation, ontologies and pathways**
Dwinell M. R., Worthey E. A., Shimoyama M., GÜNGÖR B., Depons J., Laulederkind S., Lowry T., Nigram R., Petri V.,

Smith J., et al.

Nucleic Acids Research, vol.37, no.SUPPL. 1, 2009 (SCI-Expanded)

Articles Published in Other Journals

- I. **Comparative Performance Analysis of ARIMA, Prophet and Holt-Winters Forecasting Methods on European COVID-19 Data**
ERSÖZ N. Ş., GÜNER P., AKBAŞ A., BAKİR-GUNGOR B.
International Journal of 3D Printing Technologies and Digital Industry, vol.6, no.3, pp.556-565, 2022 (Peer-Reviewed Journal)
- II. **Impact of Image Preprocessing on Skin Cancer Image Classification Using Convolutional Neural Networks**
Güngör B., Adanur Dedetürk B., Taşdemir K.
Kocaeli Journal of Science and Engineering, vol.38, no.2, pp.190-200, 2022 (Peer-Reviewed Journal)
- III. **Evaluation of Sub-Network search programs in epilepsy-related GWAS dataset**
ADANUR DEDETÜRK B., Bakir Gungor B.
PAMUKKALE UNIVERSITY JOURNAL OF ENGINEERING SCIENCES-PAMUKKALE UNIVERSITESI MUHENDISLIK BILIMLERI DERGISI, vol.28, no.2, pp.292-298, 2022 (ESCI)
- IV. **Recursive Cluster Elimination based Rank Function (SVM-RCE-R) implemented in KNIME**
Yousef M., GÜNGÖR B., Jabeer A., GÖY G., Qureshi R., C. Showe L.
F1000Research, vol.9, 2021 (Scopus)
- V. **Correlation of PAPP-A values with maternal characteristics, biochemical and ultrasonographic markers of pregnancy**
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- VI. **Classification of Cardiovascular Artery Diseases Using Artificial Neural Network**
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International Journal of Bioscience, Biochemistry and Bioinformatics, vol.10, no.1, pp.34-41, 2020 (Peer-Reviewed Journal)
- VII. **Coronary Artery Disease Diagnosis Using Optimized Adaptive Ensemble Machine Learning Algorithm**
Kolukisa B., Yavuz L., Soran A., Güngör B., Tuncer D., Önen A., Güngör V. Ç.
International Journal of Bioscience, Biochemistry and Bioinformatics, vol.10, no.1, pp.58-65, 2019 (Peer-Reviewed Journal)
- VIII. **Diagnosis of Coronary Heart Disease via Classification Algorithms and a New Feature Selection Methodology**
KOLUKISA B., Hacilar H., GÖY G., Kuş M., GÜNGÖR B., ARAL A., GÜNGÖR V. Ç.
International Journal of Data Mining Science, vol.1, no.1, pp.8-15, 2019 (Scopus)
- IX. **Weighted Association Rules and Scoring Methodology for Cardiovascular Diseases**
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International Journal of Bioscience, Biochemistry and Bioinformatics, vol.9, no.4, pp.222-230, 2019 (Peer-Reviewed Journal)
- X. **Identification of SNP Targeted Pathways From Genome wide Association Study GWAS Data**
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Nature Protocol Exchange, 2012 (Peer-Reviewed Journal)

Refereed Congress / Symposium Publications in Proceedings

- I. **Effect of Recursive Cluster Elimination with Different Clustering Algorithms Applied to Gene Expression Data**

Kuzudisli C., GÜNGÖR B., Qaqish B. F., Yousef M.

2023 Innovations in Intelligent Systems and Applications Conference, ASYU 2023, Sivas, Turkey, 11 - 13 October 2023

- II. **Population Specific Classification of Colorectal Cancer with Meta-Analysis of Metagenomic Data**
Metagenomik Verilerin Meta-Analiziyle Kolorektal Kanserin Pop lasyonuna zg Siniflandirilmesi
Temiz M., Yousef M., GÜNGÖR B.
2023 Innovations in Intelligent Systems and Applications Conference, ASYU 2023, Sivas, Turkey, 11 - 13 October 2023
- III. **Integrative analyses in omics data: Machine learning perspective**
Integrative Analysen von Omics-Daten: Perspektive des maschinellen Lernens
Yazici M., GÜNGÖR B., Yousef M.
68. Jahrestagung der Deutsche Gesellschaft für Medizinische Informatik, Biometrie und Epidemiologie e.V., GMDS 2023 - 68th Annual Conference of the German Association for Medical Informatics, Biometry and Epidemiology, GMDS 2023, Heilbronn, Germany, 17 - 21 September 2023, vol.19
- IV. **Identifying Taxonomic Biomarkers of Colorectal Cancer in Human Intestinal Microbiota Using Multiple Feature Selection Methods**
Jabeer A., Kocak A., AKKAŞ H., Yeniser F., NALBANTOĞLU Ö. U., Yousef M., GÜNGÖR B.
2022 Innovations in Intelligent Systems and Applications Conference, ASYU 2022, Antalya, Turkey, 7 - 09 September 2022
- V. **The relationship between TSH levels, maternal characteristics and racial group of the aneuploidy screening**
Soylemez U. G., Celebiler H. K., Harkonen J., GÜNGÖR B.
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- VI. **A comparative study on psychiatric disorders: Identification of shared pathways and common agents**
Kuzudisli C., GÜNGÖR B.
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- VII. **Integrating Gene Ontology Based Grouping and Ranking into the Machine Learning Algorithm for Gene Expression Data Analysis**
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- VIII. **SVM-RCE-R-OPT: Optimization of Scoring Function for SVM-RCE-R**
Yousef M., Jabeer A., Bakir-Gungor B.
32nd International Conference on Database and Expert Systems Applications (DEXA), ELECTR NETWORK, 27 - 30 September 2021, vol.1479, pp.215-224
- IX. **Blockchain-based Fog Computing Applications in Healthcare**
Saglikta Blokzincir Tabanlı Sistemlerin Uygulamaları
Adanur B., GÜNGÖR B., SORAN A.
28th Signal Processing and Communications Applications Conference, SIU 2020, Gaziantep, Turkey, 5 - 07 October 2020
- X. **A Top-Down Approach for Finding Affected Pathway Subnetworks in Type 2 Diabetes**
Tip 2 Diyabet'te Etkilenen Yolak Alt Ağlarını Bulmak İçin Yukarıdan Aşağıya İnceleyen Bir Yaklaşım
Ünlü Yazıcı M., Güngör B.
28th Signal Processing and Communications Applications Conference, SIU 2020, Gaziantep, Turkey, 5 - 07 October 2020
- XI. **The identification of affected pathway subnetworks and pathway clusters in colon cancer**
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- XII. **A New Method to Identify Affected Pathway Subnetworks and Clusters in Colon Cancer**

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- XIII. **Kolon kanserinde etkilenen genlerin ve yolakların omik veri entegrasyonu ile in siliko yöntemler kullanılarak belirlenmesi**
ÜNLÜ M., GÜNGÖR B.
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- XIV. **Performance Evaluations of Active Subnetwork Search Methods in Protein-Protein Interaction Networks Protein-Protein Etkileşim Ağlarında Aktif Alt Ağ Arama Yöntemlerinin Performans Değerlendirmeleri**
GÜNER P., GÜNGÖR B.
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- XV. **In-Silico Methods to Identify Common MicroRNAs and Pathways of Neuromuscular Diseases**
Ünlü M., Aksu E., Akkaya Ulum Z. Y., Hayta B., Güngör B.
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- XVI. **In-silico Identification of Papillary Thyroid Carcinoma Molecular Mechanisms**
Ersöz N. Ş., GÜZEL Y., GÜNGÖR B.
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- XVII. **Dört Farklı Kanser Tipine Duyarlı Genlerle İlişkili Fonksiyonel İnteraktomların Karşılaştırmalı Çalışması**
ÜNLÜ M., GÜNGÖR B.
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- XVIII. **Weighted Association Rules and Scoring Methodology for Cardiovascular Diseases**
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- XIX. **Coronary Artery Disease Diagnosis Using Optimized Adaptive Ensemble Machine Learning Algorithm**
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- XX. **Classification of Cardiovascular Artery Diseases based on Deep Neural Network**
Hacılar H., GÖRMEZ Y., GÜNGÖR B., Gezer C., AYDIN Z., GÜNGÖR V. Ç.
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- XXI. **The Identification of Discriminative Single Nucleotide Polymorphisms for the Classification of Behçets Disease**
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- XXII. **The relationship between PAPP-A levels, maternal characteristics, demographics, gestational diabetes and other biochemical markers of the aneuploidy screening**
UZUT Ö. G., KAYMAKÇALAN ÇELEBİLER H., HARKONEN J., GÜNGÖR B.
International Conference on Computer Sciences and Engineering (UBMK), 20 - 23 September 2018
- XXIII. **Paths Toward the Comprehensive Functional Enrichment of Multi-omics Data**
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- XXIV. **Machine Learning Analysis of Inflammatory Bowel Disease-Associated Metagenomics Dataset**
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- XXV. **The Identification of Discriminative Single Nucleotide Polymorphism Sets for the Classification of**

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XXVI. Identify Commonly Affected Pathways in Psychiatric Diseases

Bulut U., Bakir-Gungor B.

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XXVII. The relationship between PAPP-A levels, maternal characteristics, gestational diabetes and other biochemical markers of the aneuploidy screening

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XXVIII. Common miRNA signatures in a group of rare neuromuscular disorders

AKSU E., AKKAYA ULUM Z. Y., PEYNİRCİOĞLU B., DAYANGAÇ ERDEN D., YÜZBAŞIOĞLU A., GÜNGÖR B., TALİM B., HAYTA B.

43rd FEBS 2018 Congress, 7 - 12 July 2018

XXIX. Evaluation of Classification Algorithms, Linear Discriminant Analysis and a New Hybrid Feature Selection Methodology for the Diagnosis of Coronary Artery Disease

KOLUKISA B., Hacilar H., Goy G., Kus M., Bakir-Gungor B., Aral A., GÜNGÖR V. Ç.

2018 IEEE International Conference on Big Data, Big Data 2018, Washington, United States Of America, 10 - 13 December 2018, pp.2232-2238

XXX. Comparison of Disease Specific Sub-Network Identification Programs

Adanur B., GÜNGÖR B.

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XXXI. GWAS Analizi ve Kompleks Hastalıklarda Etkilenen Yolakların Belirlenmesi

GÜNGÖR B.

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XXXII. The Identification of Pathway Markers in Behcet's Disease Using Genomewide Association Data From Two Different Populations.

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