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Education

Doctorate
2000 - 2007 Ohio State University, The Molecular, Cellular And Developmental Biology Graduate Program, United States Of America

Undergraduate
1996 - 2000 İhsan Dogramaci Bilkent University, Faculty Of Science, Department Of Molecular Biology And Genetics, Turkey

Supported Projects

1. YILMAZ A., YENİDÜNYA YALIN G., Özkurt K., Project Supported by Higher Education Institutions, Retina Görüntülerinden ve Konuşma Sesi Analizinden Gestasyonel Diyabet Tahmini için Derin Öğrenme Modellerinin Geliştirilmesi, 2024 - Continues
2. Çakır R., Yılmaz A., Çalık H., Arı Yuka S., Maden H., Emergency R&D Project of Group A, Kolorektal Kanserde Çoklu İlaç Direncine Karşı In Siliko Terapötik Peptit Tasarlanması ve Kitosan Nanopartikülleri ile Etkinliğinin In Vitro İncelenmesi, 2024 - 2026
3. Pişkin M. B., Üstündağ C. B., Çakır Koç R., Yılmaz A., Yeşilkır S., TUBITAK Project, Kompozit Nanofiber Kardiyak Yamaların Adipoz Doku Kaynaklı Mezenkimal Kök Hücrelerde Kardiyojenik Farklılaşma Ve Transkriptomik Değişiklikler Üzerine Etkisinin İncelenmesi, 2022 - 2025
4. YILMAZ A., YÖNET N., TÜRKER M., Project Supported by Higher Education Institutions, Bitki Sekonder Metabolitlerinin Üretilimi İçin Aday Moleküllerin Retrosentez Yaklaşımı ile Ortaya Çıkarılması, 2021 - 2024
5. ÇAKIR R., YILMAZ R., ÇALIK H., ARI YUKA S., YILMAZ A., YÜCEL B., YAMAN S., KÜÇÜKBAŞ M., Project Supported by Higher Education Institutions, Paklitaksel Dirençli Yumurtalık Kanseri Hücre Hattında İlaç Direncini Tersine Çevirebilecek Nitidine-Paklitaksel Yüklü Lipit-Kitosan Hibrit Nanopartiküllerinin Geliştirilmesi, 2021 - 2023
6. YILMAZ A., Project Supported by Higher Education Institutions, İnsan Nöroblastoma Hücrelerinde Metal-Uyarıcılı Oksidatif Stresle İndüklenen Alzheimer Modeli, 2016 - 2017
7. YILMAZ A., BULUT YILGÖR M., ÖZTAŞ E. S., TUBITAK Project, Bazı Sonlu Halkalar Üzerinde Tanımlı Ters Sıralı-Tamlama Kodları Ve DNA\ya Uygulamaları, 2013 - 2015

Published journal articles indexed by SCI, SSCI, and AHCI

1. **The role of ceRNAs in breast cancer microenvironmental regulation and therapeutic implications**
YILMAZ A., Ari Yuka S.
Journal of Molecular Medicine, vol.103, no.1, pp.33-49, 2025 (SCI-Expanded)
2. **Decoding dynamic miRNA:ceRNA interactions unveils therapeutic insights and targets across**

predominant cancer landscapes.

Ari Yuka S., Yilmaz A.

BIODATA MINING, vol.17, no.1, pp.11, 2024 (SCI-Expanded)

3. **Efficient Storage and Analysis of Genomic Data: A k-mer Frequency Mapping and Image Representation Method**

Luleci H. B., Yuka S. A., YILMAZ A.

Interdisciplinary Sciences - Computational Life Sciences, 2024 (SCI-Expanded)

4. **Comparative analysis of retrosynthesis applications for predicting of pathways for plant Secondary metabolite production**

Yonet N., Ari Yuka S., Türker M., Yilmaz A.

Biotechnology and Biotechnological Equipment, vol.38, no.1, 2024 (SCI-Expanded)

5. **Biofabrication of Acer palmatum-Mediated Multifunctional CuO Nanoparticles for Dye Removal, Antibacterial-Antifungal Activity, and Molecular Docking**

Sazak C., ATTAR A., YILMAZ A., ALTIKATOĞLU YAPAÖZ M.

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6. **PGT for structural chromosomal rearrangements in 300 couples reveals specific risk factors but an interchromosomal effect is unlikely**

Ogur C., Kahraman S., Griffin D. K., Cinar Yapan C., Tufekci M. A., Cetinkaya M., TEMEL Ş. G., YILMAZ A.

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7. **Robust and rigorous identification of tissue-specific genes by statistically extending tau score**

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8. **Effect of SARS-CoV-2 infection on host competing endogenous RNA and miRNA network.**

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PeerJ, vol.9, no.e12370, 2021 (SCI-Expanded)

9. **Network based multifactorial modelling of miRNA-target interactions**

Ari Yuka S., Yilmaz A.

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10. **Important biological information uncovered in previously unaligned reads from chromatin immunoprecipitation experiments (ChIP-Seq)**

Ouma W. Z., Mejia-Guerra M. K., YILMAZ A., PAREJA-TOBES P., Li W., Doseff A. I., Grotewold E.

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11. **Molecular basis for the action of a dietary flavonoid revealed by the comprehensive identification of apigenin human targets**

Arango D., Morohashi K., Yilmaz A., Kuramochi K., Parihar A., Brahimaj B., Grotewold E., Doseff A. I.

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12. **Unraveling the KNOTTED1 regulatory network in maize meristems**

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13. **A Genome-Wide Regulatory Framework Identifies Maize Pericarp Color1 Controlled Genes**

Morohashi K., Casas M. I., Falcone Ferreyra L., Mejia-Guerra M. K., Pourcel L., Yilmaz A., Feller A., Carvalho B., Emiliani J., Rodriguez E., et al.

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14. **Thriving under Stress: Selective Translation of HIV-1 Structural Protein mRNA during Vpr-Mediated Impairment of eIF4E Translation Activity**

Sharma A., Yilmaz A., Marsh K., Cochrane A., Boris-Lawrie K.

PLOS PATHOGENS, vol.8, no.3, 2012 (SCI-Expanded)

15. **AGRIS: the Arabidopsis Gene Regulatory Information Server, an update**

Yilmaz A., Mejia-Guerra M. K., Kurz K., Liang X., Welch L., Grotewold E.

NUCLEIC ACIDS RESEARCH, vol.39, 2011 (SCI-Expanded)

16. **Genome sequencing and analysis of the model grass *Brachypodium distachyon***
 Vogel J. P., Garvin D. F., Mockler T. C., Schmutz J., Rokhsar D., Bevan M. W., Barry K., Lucas S., Harmon-Smooth M., Lail K., et al.
 NATURE, vol.463, no.7282, pp.763-768, 2010 (SCI-Expanded)
17. **Components and Mechanisms of Regulation of Gene Expression**
 YILMAZ A., Grotewold E.
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18. **The word landscape of the non-coding segments of the *Arabidopsis thaliana* genome**
 Lichtenberg J., Yilmaz A., Welch J. D., Kurz K., Liang X., Drews F., Ecker K., Lee S. S., Geisler M., Grotewold E., et al.
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19. **GRASSIUS: A Platform for Comparative Regulatory Genomics across the Grasses**
 Yilmaz A., Nishiyama M. Y., Fuentes B. G., Souza G. M., Janies D., Gray J., Grotewold E.
 PLANT PHYSIOLOGY, vol.149, no.1, pp.171-180, 2009 (SCI-Expanded)
20. **NOD/SCID mouse model of canine T-cell lymphoma with humoral hypercalcaemia of malignancy: cytokine gene expression profiling and in vivo bioluminescent imaging**
 Nadella M. V. P., Kisseberth W. C., Nadella K. S., Thudi N. K., Thamm D. H., McNeil E. A., Yilmaz A., Boris-Lawrie K., Rosol T. J.
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21. **RNA helicase A interacts with divergent lymphotropic retroviruses and promotes translation of human T-cell leukemia virus type 1**
 Bolinger C., Yilmaz A., Hartman T. R., Kovacic M. B., Fernandez S., Ye J., Forget M., Green P. L., Boris-Lawrie K.
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22. **Retrovirus translation initiation: Issues and hypotheses derived from study of HIV-1**
 Yilmaz A., BOLINGER C., BORIS-LAWRIE K.
 CURRENT HIV RESEARCH, vol.4, no.2, pp.131-139, 2006 (SCI-Expanded)
23. **Coordinate enhancement of transgene transcription and translation in a lentiviral vector**
 Yilmaz A., FERNANDEZ S., LAIRMORE M. D., BORIS-LAWRIE K.
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Articles Published in Other Journals

1. **Balıkesir/Antandros Antik Kenti Kazısı Toprak Örneklerinin Metagenomik Analizi**
 Bal D., Balci B., Yilmaz A., Polat G., Arican E.
 Süleyman Demirel Üniversitesi Fen Bilimleri Enstitüsü Dergisi, vol.26, no.1, pp.1-12, 2022 (Peer-Reviewed Journal)
2. **Convolutional Neural Network Approach to Predict Tumor Samples Using Gene Expression Data**
 DARENDELİ KIRAZ B. N., YILMAZ A.
 Zeki sistemler teori ve uygulamaları dergisi (Online), vol.4, no.2, pp.136-141, 2021 (Peer-Reviewed Journal)
3. **Assessment of Mutation Susceptibility in DNA Sequences with Word Vectors**
 YILMAZ A.
 Zeki Sistemler Teori ve Uygulamaları Dergisi, vol.3, no.1, pp.1-6, 2020 (Peer-Reviewed Journal)
4. **POPCORN: An Online resource providing access to distributed and diverse maize project data**
 Cannon E. K., Birkett S. M., Braun B. L., Kodavali S., Jennewein D. M., Yilmaz A., Antonescu V., Antonescu C., Harper L. C., Gardiner J. M., et al.
 International Journal of Plant Genomics, vol.2011, 2011 (Scopus)

Papers Published in Refereed Scientific Meetings

1. **A New Network-Based Tool to Analyse Competing Endogenous RNAs**
 Yuka S., Yilmaz A.

16th International Symposium on Bioinformatics Research and Applications, ISBRA 2020, Moscow, Russia, 1 - 04 December 2020, pp.274-281

2. Constructing Accurate Compound Networks for Reliable Network Analysis

Alpaydin D., Öztürk R., Yilmaz A.

12th International Symposium on Health Informatics and Bioinformatics (HIBIT) 2019, İzmir, Turkey, 17 - 18 October 2019, pp.230

3. Reduction of Depression Symptoms via the Metabolites of Gut Microbiota

Mawaldı I., Yilmaz A.

12th International Symposium on Health Informatics and Bioinformatics (HIBIT) 2019, İzmir, Turkey, 17 - 18 October 2019, pp.218

4. Automation of Genetic Screening and Reporting with Snakemake

Kalyon O., Müslümanoğlu M. H., Yilmaz A.

12th International Symposium on Health Informatics and Bioinformatics (HIBIT) 2019, İzmir, Turkey, 17 - 18 October 2019, pp.219

5. Prediction of Mutation Susceptibility Based on Solely DNA Sequences with Word2Vec

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6. Breast cancer gene expression profile analysis in terms of genes specific to non- breast tissues

Konuk H. B., Yilmaz A.

1st International Conference of Genomics and Bioinformatics (ICGB), Aydin, Turkey, 1 - 04 November 2018, pp.30

7. Batch Analysis of RNA-Seq Data with Docker in AWS Cloud

Yilmaz A.

ICAMLS 2018 International Conference on Applied Mathematics, Modeling and Life Science Problems, İstanbul, Turkey, 3 - 05 October 2018, pp.87

8. What is the relationship between cancer DEGs and tissue specific genes

Konuk H. B., Cesur M. R., Yilmaz A.

Lisansüstü Araştırmalar Sempozyumu ve Tanıtım Günleri, Kocaeli, Turkey, 17 - 18 May 2017

9. Öztürk R., Yilmaz A.

1.Temel Onkoloji Sempozyumu, İzmir, Turkey, 9 - 11 May 2018, pp.30

10. Darendeli B. N., Al-Qabbani H., Yilmaz A.

1.Temel Onkoloji Sempozyumu, İzmir, Turkey, 9 - 11 May 2018, pp.48

11. Pinpointing the importance of compounds in human metabolic network

Öztürk E. R., Yilmaz A.

7th International Molecular Biology and Biotechnology Congress, Konya, Turkey, 25 - 27 April 2018, pp.160

12. Multifactorial Modelling of MicroRNA Associated Repression and Its Subsequent Effects on Gene Expression in MicroRNA:Target Network

Arı S., Yilmaz A.

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13. RIGOROUS IDENTIFICATION OF TISSUE SPECIFIC GENES IN SILICO: REVEALING THE INTERPLAY BETWEEN CANCER SPECIFIC EXPRESSION AND TISSUE SPECIFIC EXPRESSION

KONUK H. B., YILMAZ A.

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14. GENERATING LOSSLESS COMPRESSION OF GENOME SCALE K-MER FREQUENCY TABLE AS RASTER IMAGE

KONUK H. B., YILMAZ A.

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15. IDENTIFICATION of TISSUE SPECIFIC GENES and ASSESMENT of THEIR INTERSECTION WITH DIFFERENTIALLY EXPRESSED GENES in CANCER DATA to UNDERSTAND TUMOR HETEROGENEITY IN

SILICO

Konuk H. B., Cesur M. R., Yilmaz A.

10 th International Symposium on Health Informatics and Bioinformatics, Güzelyurt, Cyprus (Kktc), 28 June 2017

16. SURVEY OF CHIMERIC mRNAs IN HIV INFECTED PATIENT RNA-SEQ DATA

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10th International Symposium on Health Informatics and Bioinformatics, Güzelyurt, Cyprus (Kktc), 28 June 2017

17. Prebiotics and Probiotics: Break the 'Self-Domino Effect' for obesity Treatment

Özel C., Yilmaz A.

IVEK 3rd International Convention of Pharmaceuticals and Pharmacies, Istanbul, Turkey, 26 April 2017, pp.338

18. Robust and Rigorous Classification of Tissue Specific Genes

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INTERNATIONAL DNA DAY AND GENOME CONGRESS (IDDGC) 2017, Kırşehir, Turkey, 24 April 2017

19. Prediction of splice site using AdaBoost with a new sequence encoding approach

Pashaei E., YILMAZ A., ÖZEN M., AYDIN N.

2016 IEEE International Conference onSystems, Man, and Cybernetics, 9 - 12 October 2016

20. A Combined SVM and Markov Model approach for Splice Site Identification

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6th International Conference on Computer and Knowledge Engineering (ICCKE), Mashhad, Iran, 20 - 21 October 2016, pp.200-204

21. A Novel Method for Splice Sites Prediction Using Sequence Component and Hidden Markov Model

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22. Utilization of Multiple Mapped Sequences in Next Generation Sequencing Especially in Chip Seq Experiments

ZOR S., YILMAZ A.

International Health Informatics and Bioinformatics Symposium, Muğla, Turkey, 16 - 17 October 2015

23. Computational Identification of miRNA Targets Using Olive Library

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24. Molecular Analysis of Lipid Transfer Gene Selected from Olive Olea europaea ESTs Expressed Sequence Tags Collection

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25. Construction and Comparison of EST Library for Expressed Genes in Normal and Tumoral Mammary Gland Tissue of the Terrier Dog

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26. Design and synthesis of small peptide sequences to detect concentrations of free transition metal ions

YÜCESAN G., Yılmaz A., Adiyaman R.

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27. Analyses of EST Collection from Prenatal and Postnatal Mammary Tissue of Sheep

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28. Discovery of regulatory networks in plants by linking promoter and transcription factor databases.

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29. Construction of genomic regulatory encyclopedias: Strategies and case studies

Alam M., Bitterman T., Drews F., Ecker K., Elnitski L., Evans S., YILMAZ A., Welch L.

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Supervised Theses

Yılmaz A., PRENATAL TANIDA TRİZOMİ RİSK TAHMİNİ İÇİN QF-PCR SONUÇLARI, DEMOGRAFİK, KLINİK VE GENETİK VERİLERİN ENTEGRASYONUNUN MAKİNE ÖĞRENMESİ TABANLI ANALİZİ, Postgraduate, K.TÜREGÜN(Student), 2025

Yılmaz A., Modeling redistribution of regulators after perturbation on large biological networks, Doctorate, S.ARI(Student), 2024

Yılmaz A., Meme kanseri yeni nesil dizileme verilerini birleştirerek biyokimyasal etki hesaplama yöntemi geliştirilmesi, Doctorate, M.ÇELİK(Student), 2024

Yılmaz A., INSR/IGF1R hibrit reseptörünü inhibe edecek ilaçların moleküler dinamik simülasyon yöntemiyle incelenmesi, Postgraduate, F.TOSUN(Student), 2024

Yılmaz A., Preimplantation genetic diagnosis in balanced rearrangement carriers and investigation of inter chromosomal effect, Doctorate, Ç.OĞUR(Student), 2023

Yılmaz A., Survey of expression profiles of circular RNAs and their parent genes in context of tissue specificity, Postgraduate, E.İREM(Student), 2022

Yılmaz A., Discovery of Indirect Interactions Between Genes by Deep Learning Using Gene Expression Data, Postgraduate, G.ÇELEN(Student), 2022

Yılmaz A., Using gene expression profiles of cancer patients with image-based deep learning approach to develop methods for classification and prediction of cancer while revealing critical genes, Postgraduate, B.NUR(Student), 2021

Yılmaz A., Analysis of interplay between gene expression and miRNA expression in breast cancer next generation sequencing data, Postgraduate, S.ARI(Student), 2018

Yılmaz A., Analysis of biochemical pathway with graph based algorithms, Postgraduate, E.RAVZA(Student), 2018

Yılmaz A., Analysis of Intra-tumoral Heterogeneity in Context of Tissue Specific Gene Expression with Computational Approach, Postgraduate, H.BÜŞRA(Student), 2018

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Peer Reviews in Scientific Publications

Plant Methods dergisi için makale değerlendirilmesi, SCI Journal, June 2013

Türk Biyoloji Dergisi, BIY-1111-23 nolu makalenin değerlendirilmesi, National Scientific Refreed Journal, March 2012