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SCI, SSCI ve AHCI İndekslerine Giren Dergilerde Yayınlanan Makaleler

- I. **Identification of novel *MYH14* variants in families with autosomal dominant sensorineural hearing loss**
DUMAN D., Ramzan M., Subasioglu A., MUTLU A., Peart L., Seyhan S., Guo S., Ila K., Balta B., KALCIOĞLU M. T., et al.
AMERICAN JOURNAL OF MEDICAL GENETICS PART A, 2024 (SCI-Expanded)
- II. **Genetic heterogeneity in hereditary hearing loss: Potential role of kinociliary protein TOGARAM2**
Ramzan M., Zafeer M. F., Abad C., Guo S., Owrang D., Alper O., MUTLU A., Atik T., DUMAN D., Bademci G., et al.
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- III. **Novel GPR156 variants confirm its role in moderate sensorineural hearing loss**
Ramzan M., Bozan N., Seyhan S., Zafeer M. F., Ayril A., Duman D., Bademci G., Tekin M.
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- IV. **Dispersed DNA variants underlie hearing loss in South Florida's minority population**
Peart L., Gonzalez J., Morel Swols D., Duman D., Saridogan T., Ramzan M., Zafeer M. F., Liu X. Z., Eshraghi A. A., Hoffer M. E., et al.
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- V. **Genome sequencing identifies coding and non-coding variants for non-syndromic hearing loss**
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- VI. **Biallelic KITLG variants lead to a distinct spectrum of hypomelanosis and sensorineural hearing loss**
Vona B., Schwartzbaum D. A., Rodriguez A. A., Lewis S. S., Toosi M. B., Radhakrishnan P., Bozan N., Akın R., Doosti M., Manju R., et al.
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- VII. **Mutations in MINAR2 encoding membrane integral NOTCH2-associated receptor 2 cause deafness in humans and mice**
Bademci G., Lachgar-Ruiz M., Deokar M., Zafeer M. F., Abad C., Baylan M. Y., Ingham N. J., Chen J., Sineni C. J., Vadgama N., et al.
Proceedings of the National Academy of Sciences of the United States of America, cilt.119, sa.26, 2022 (SCI-Expanded)
- VIII. **Whole Mitochondrial Genome Analysis in Turkish Patients with Mitochondrial Diseases**
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BALKAN MEDICAL JOURNAL, cilt.39, sa.2, ss.96-106, 2022 (SCI-Expanded)
- IX. **Long-range cis-regulatory elements controlling GDF6 expression are essential for ear development**

Bademci G., Abad C., Cengiz F. B., Seyhan S., İNCESULU Ş. A., Guo S., Fitoz S., ATLI E. İ., Gosstola N. C., DEMİR S., et al. JOURNAL OF CLINICAL INVESTIGATION, cilt.130, sa.8, ss.4213-4217, 2020 (SCI-Expanded)

- X. **Novel variant p.E269K confirms causative role of PLS1 mutations in autosomal dominant hearing loss**
Diaz-Horta O., Bademci G., TOKGÖZ YILMAZ S., Guo S., Zafeer F., Sineni C. J., DUMAN D., Farooq A., Tekin M. CLINICAL GENETICS, cilt.96, sa.6, ss.575-578, 2019 (SCI-Expanded)
- XI. **Adams-Oliver syndrome caused by mutations of the EOGT gene**
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- XII. **A truncating CLDN9 variant is associated with autosomal recessive nonsyndromic hearing loss**
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- XIII. **Genetic Causes of Inner Ear Anomalies: a Review from the Turkish Study Group for Inner Ear Anomalies**
OCAK E., DUMAN D., Tekin M. BALKAN MEDICAL JOURNAL, cilt.36, sa.4, ss.206-211, 2019 (SCI-Expanded)
- XIV. **FOXF2 is required for cochlear development in humans and mice**
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- XV. **Dysfunction of GRAP, encoding the GRB2-related adaptor protein, is linked to sensorineural hearing loss**
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- XVI. **Identification of candidate gene FAM183A and novel pathogenic variants in known genes: High genetic heterogeneity for autosomal recessive intellectual disability**
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- XVII. **Elucidating the genetic architecture of Adams-Oliver syndrome in a large European cohort**
Meester J. A. N., Sukalo M., Schroeder K. C., Schanze D., Baynam G., Borck G., Bramswig N. C., DUMAN D., Gilbert-Dussardier B., Holder-Espinasse M., et al. HUMAN MUTATION, cilt.39, sa.9, ss.1246-1261, 2018 (SCI-Expanded)
- XVIII. **MPZL2 is a novel gene associated with autosomal recessive nonsyndromic moderate hearing loss**
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- XIX. **Variants in CIB2 cause DFNB48 and not USH1J**
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- XX. **Novel pathogenic variants underlie SLC26A4-related hearing loss in a multiethnic cohort**
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- XXI. **Novel EYA1 variants causing Branchio-oto-renal syndrome**
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- XXII. **Variations in Multiple Syndromic Deafness Genes Mimic Non-syndromic Hearing Loss**
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- XXIII. **Spectrum of DNA variants for non-syndromic deafness in a large cohort from multiple continents**
Yan D., Tekin D., Bademci G., Foster J., Cengiz F. B., Kannan-Sundhari A., Guo S., Mittal R., Zou B., Grati M., et al.
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- XXIV. **ROR1 is essential for proper innervation of auditory hair cells and hearing in humans and mice**
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- XXV. **Comprehensive analysis via exome sequencing uncovers genetic etiology in autosomal recessive nonsyndromic deafness in a large multiethnic cohort**
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- XXVI. **Novel mutations confirm that COL11A2 is responsible for autosomal recessive non-syndromic hearing loss DFNB53**
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- XXVII. **HPSE2 Mutations in Urofacial Syndrome, Non-Neurogenic Neurogenic Bladder and Lower Urinary Tract Dysfunction**
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- XXVIII. **Identification of Copy Number Variants Through Whole-Exome Sequencing in Autosomal Recessive Nonsyndromic Hearing Loss**
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- XXIX. **Analysis Of Hpse2 Gene Mutations In Children With Non-neurogenic Neurogenic Bladder And Urofacial (ochoa) Syndrome**
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- XXX. **FAM65B is a membrane-associated protein of hair cell stereocilia required for hearing**
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- XXXI. **Evidence for genotype-phenotype correlation for OTOF mutations**
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- XXXII. **BRANCHIO-OCULO-FACIAL SYNDROME IN A NEWBORN CAUSED BY A NOVEL TFAP2A MUTATION**
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- XXXIII. **High frequency of kidney and urinary tract anomalies in asymptomatic first-degree relatives of patients with ÇAKUT**
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- XXXIV. **SLITRK6 mutations cause myopia and deafness in humans and mice**
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- XXXV. **Whole-Exome Sequencing Efficiently Detects Rare Mutations in Autosomal Recessive Nonsyndromic Hearing Loss**
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- XXXVI. **MASP-1 is crucial for lectin pathway activation in human serum, while neither MASP-1 nor MASP-3 are required for alternative pathway function**
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- XXXVII. **Mutations in OTOGL, Encoding the Inner Ear Protein Otogelin-like, Cause Moderate Sensorineural Hearing Loss**
Yariz K. O., DUMAN D., Seco C. Z., Dallman J., Huang M., Peters T. A., Sirmaci A., Lu N., Schraders M., Skromne I., et al.
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- XXXVIII. **A truncating mutation in GPSM2 is associated with recessive non-syndromic hearing loss**
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- XXXIX. **Cardiomyopathy with alopecia and palmoplantar keratoderma (CAPK) is caused by a JUP mutation**
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- XL. **A founder TMIE mutation is a frequent cause of hearing loss in southeastern Anatolia**
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- XLI. **Homozygous FGF3 mutations result in congenital deafness with inner ear agenesis, microtia, and microdontia**
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- XLII. **Paternal X could relate to arithmetic function; study of cognitive function and parental origin of X chromosome in Turner syndrome**
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- XLIII. **The c.IVS1+1G > A mutation in the GJB2 gene is prevalent and large deletions involving the GJB6 gene are not present in the Turkish population**
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- XLIV. **A novel missense mutation in a C2 domain of OTOF results in autosomal recessive auditory neuropathy**
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- XLV. **657del5 mutation of the Nijmegen breakage syndrome gene (NBS1) in the Turkish population**
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- XLVI. **Screening the SLC26A4 gene in probands with deafness and goiter (Pendred syndrome) ascertained from a large group of students of the schools for the deaf in Turkey**
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Diğer Dergilerde Yayınlanan Makaleler

- I. **Research of genetic bases of hereditary non-syndromic hearing loss**
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- I. **Molecular diagnosis of lethal multiple pterygium syndrome in a fetus presenting with fetal akinesia deformation sequence**
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- II. **Genome sequencing identifies coding and non-coding variants in hereditary deafness missed by exome sequencing**
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- III. **Biallelic variants in KITLG cause Waardenburg syndrome type 2, albinism-deafness syndrome and oculocutaneous albinism**
Girisha K., Vona B., Schwartzbaum D., Rodriguez A. A., Lewis S. S., Toosi M. B., Radhakrishnan P., Bozan N., Akın R., Doosti M., et al.
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- IV. **Characterization of MINAR2 as a Novel Autosomal Recessive Deafness Gene**
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- V. **Expanding KITLG to Include Autosomal Recessive Inheritance, Oculocutaneous Albinism and Waardenburg/Albinism-Deafness Syndromes**
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- VI. **Leigh Sendromu (LS) Ön Tanılı Bir Hastada Olası Genetik Etmenlerin Araştırılması**
gencer öncül e. b., DUMAN D., EMİNOĞLU F. T., AKTUNA S., DUMAN M. T.
VI. Tıbbi Biyoloji ve Genetik Kongresi, Muğla, Türkiye, 27 - 30 Ekim 2019
- VII. **İşitme kaybının genetiği**
DUMAN D.
V. Uluslararası Katılımlı Odyoloji Kongresi, Ankara, Türkiye, 11 - 12 Mayıs 2019
- VIII. **Whole Mitochondrial Genome Screening in Children with Suspected Mitochondrial Disease**
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Uluslararası Metabolik Hastalıklar ve Beslenme Kongresi, 10 - 14 Nisan 2019
- IX. **A founder mutation uncovers MPZL2 (DFNB111) as a novel autosomal recessive non-syndromic moderate hearing loss gene**
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- X. **Using Whole Genome Sequencing to Identify Novel Variants in Branchio-Oto-Renal Syndrome**
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- XI. **İŞİTME KAYBI GENETİĞİ**

DUMAN D.

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- XII. **Mitokondriyal kalıtım gösteren tanı konulamamış nadir nöroloji, psikiyatri, metabolizma hastalarında mitokondriyal DNA (mtDNA)'nın taranması**
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- XIII. **Association of TOMM40 (Rs1160985 and Rs157581) Polymorphisms with Alzheimer Disease**
RAJAB M. F., gencer oncül e. b., TAŞTAN H., DUMAN D., DİRİK E. B., DUMAN M. T.
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- XIV. **Three novel hearing loss genes reveal previously unrecognized roles of their protein products in the perception of sound.**
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- XV. **Sendromik olmayan işitme kayıplı Türk ailelerde sorumlu genlerin araştırılması**
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- XVI. **A recessive variant in forkhead box domain of FOXF2 is associated with profound hearing loss and inner ear anomaly**
Bademci G., Diaz-Horta O., İNCESULU Ş. A., Cengiz F. B., Abad C., Sineni C., Foster J., DUMAN D., FİTOZ Ö. S., Blanton S., et al.
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- XVII. **Non Sendromik Ailesel İşitme Kayıplı Hastalarda Genetik Analiz**
YILDIRIM H., YILDIRIM BAYLAN M., DUMAN D., Bademci G.
5. Ulusal Otoloji Nöroloji Kongresi, Antalya, Türkiye, 4 - 07 Mayıs 2017
- XVIII. **Comprehensive Search for Deafness Genes Associated with Inner Ear Anomalies**
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- XIX. **Tedavi Edilebilir Bir Otizm Nedeni Olarak BCKD-Kinaz Eksikliği: İki Olgu Sunumu**
ÖZTÜRK HİŞMİ B., AKAY H., ÖZBEK M. N., BACANLI A., KUMRU B., DUMAN D., HİŞMİ A., Bademci G., Tekin M.
XIII. Ulusal Metabolik Hastalıklar ve Beslenme Kongresi, Adana, Türkiye, 14 - 18 Nisan 2015
- XX. **BCKD kinase deficiency Expanding the disease phenotypewith sensorineural hearing loss and transient acrodermatitisenteropathica like dermatitis**
HİŞMİ B., ÖZTÜRKMEN AKAY H., BADEMCİ G., DUMAN D., BACANLI A., HİŞMİ A., KUMRU B., TEKİN M.
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