

Barış Salman

Curriculum Vitae

■ barslmn@gmail.com□ +90 534 387 84 20⑤ barissalman.xyz (Under Construction)

technical skills

DNA/RNA/Protein

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Isolation,

quantative/qualitative

PCRs, Blotting, Cytogenetics and

Karyotyping

Functional Studies Cell Culture, in vivo

models, Microscopy

Genetic Engineering Recombinant DNA

technologies

Computers

Development Python, R, Unix Shell

Workflow Linux, Doom Emacs,

Git, Jupyter

Writing Orgmode, Zotero,

Bibtex, LAT⊨X

Web Development Django, Javascript

• Bioinformatics

Sequencing WGS, WES, Panels, RNAseq, scRNAseq,

Sanger

Micro-arrays Genotyping,

Expression

• Data Analyses

Analysis Python Scientific

Toolkit(numpy, pandas,

scipy etc.), R language, Shell Scripting,

Scripting,
Bioinformatic tools

Machine Learning, PCA, Clustering

Statistics and Probability

Hypothesis testing, Bayesian Theorem, Data distributions

Visualization Matplotlib, Plotly/Dash,

EX

education

Doctorate, 2021-2024 Genetics, Institute of Health Sciences, İstanbul University, Turkey **GPA 3.5**

Thesis Work in progress

Adviser Prof. Dr. Sibel Uğur İşeri

Notable Courses Advanced Molecular Genetics, Mendelian and non Mendelian

Diseases, Chromosomal Diseases, Systems Biology

Master, 2016-2019 Genetics, Institute of Health Sciences, İstanbul University, Turkey GPA 2.14

Thesis Study of Split Hand/Foot Malformation with Genomic Techniques

and Bioinformatic Approaches

Adviser Assoc Prof. Dr. Sibel Uğur İşeri

Notable Courses Bioinformatics, Genetic Counseling, Medical Genetics, Hospital

Rotation for Genetic Counseling

Bachelor, 2011-2016 Molecular Biology and Genetics, İstanbul University, Turkey

Thesis Metabolic Regulation in Prokaryotes with Small Non-Coding

RNAs

Adviser Asst. Prof. Dr. Semian Karaer Uzuner

Notable Courses Methods Used in Molecular Biology(lab), Molecular Genetics,

Genetic Engineering

Internships

Certificates

robot

Liquid handling

Cell Culture Use and care of August 2016 IU Cerrahpasa laboratory animals **IU HADYEK 46.**

Department of Medical

Biology

Genomics June-July 2015

IU A.S. Inst. of Expr.

Medicine Department

of Genetics

Cytogenetics June-July 2014

IU Department of

Medical Genetics

Molecular Dynamics 2012-2013

IU Faculty of Science

Quantum **Technologies** Laboratory

Mar 21 - Apr 08

Laboratory Animal Use

GPA 2.44

Training

2-5 November 2021

Hamilton Venus Online Advanced Software

Training

9-12 August 2021

Hamilton Venus Online

Basic Software

Training

jobs

CompanyRefgen BiotechnologyDevelopment of pipelines for NGSDateMar 2021–OngoingVariant classification automatization

Position Genetics and Bioinformatics Specialist

CompanyGen-Era DiagnosticsDateOct. 2018-Mar 2021

Position Genetics and Bioinformatics Specialist

Company Aktif Gen

Date Jan. 2018–Oct 2018

Position R&D Personnel

publications

Oguz-Akarsu E, Salman B, Ugur-Iseri S, Baykan B. An Extraordinary EEG Phenomenon Misdiagnosed as Nonconvulsive Status Epilepticus: Frequent Subclinical Periodic Discharges Terminated by Sudden Auditory Stimuli. Clin EEG Neurosci. Published online October 5, 2022:15500594221129964. doi:10.1177/15500594221129965

- Salman B., Yucesan E., Samanci B., Bilgic B., Hanagasi H., Gurvit H., Ozbek U., Ugur Iseri S. Combined Analysis Of Linkage And Whole Exome Sequencing Reveals CIC As A Candidate Gene For Isolated Dystonia. Journal of Istanbul Faculty of Medicine. 2021;84(4):457-463. doi:10.26650/IUITFD.2021.913346
- 3. Haryanyan G, Ozdemir O, Tutkavul K, Salman B et al. The rare rs769301934 variant in NHLRC1 is a common cause of Lafora disease in Turkey. J Hum Genet. 2021;66(12):1145-1151. doi:10.1038/s10038-021-00944-8
- 4. Akçakaya NH, Salman B, Görmez Z, et al. A Novel and Mosaic WDR45 Nonsense Variant Causes Beta-Propeller Protein-Associated Neurodegeneration Identified Through Whole Exome Sequencing and X chromosome Heterozygosity Analysis. Neuromolecular Med. 2019;21(1):54-59. doi:10.1007/s12017-018-08522-6
- 5. Bekdik Şirinocak P, Salman B, Kesim FY, Bebek N, Baykan B, Ugur İşeri SA. **Susceptibility to Juvenile Myoclonic Epilepsy Associated with the EFHC1 Gene: First Case Report in Turkey**. tnd. 2019;25(4):233-236. doi:10.4274/tnd.galenos.2019.61214

• in pipeline

- Exome sequencing reveals a treatable neurometabolic origin in two siblings with an undiagnosed form of neurodevelopmental disorder: From exon to neuron (in review)
- Clinical and genetic analyses in syndromic intellectual disability with primary microcephaly reveal biallelic and de novo variants in patients with parental consanguinity (in review)

- ELP2 Mutation Leads Developmental and Epileptic Encephalopathy: Case Report Consanguineous Family from Turkey and Review of Literature (in review)
- Variant Version Control: A Git Framework For Keeping Track Of Variant Annotation Changes (in prep.)

Collaborative

- 1. Niestroj LM, Perez-Palma E, Howrigan DP, et al. Epilepsy subtype-specific copy number burden observed in a genome-wide study of 17,458 subjects. Brain. 2020;143(7):2106-2118. doi:10.1093/brain/awaa171
- 2. Stevelink R, Luykx JJ, Lin BD, et al. Shared genetic basis between genetic generalized epilepsy and background electroencephalographic oscillations. Epilepsia. 2021;62(7):1518-1527. doi:10.1111/epi.16922
- 3. Leu C, Stevelink R, Smith AW, et al. Polygenic burden in focal and generalized epilepsies. Brain. 2019;142(11):3473-3481. doi:10.1093/brain/awz292
- 4. Koko M, Krause R, Sander T, et al. Distinct gene-set burden patterns underlie common generalized and focal epilepsies. EBioMedicine. 2021;72:103588. doi:10.1016/j.ebiom.2021.103588
- Epi25 Collaborative. Electronic address: s.berkovic@unimelb.edu.au, Epi25 Collaborative. Ultra-Rare Genetic Variation in the Epilepsies: A Whole-Exome Sequencing Study of 17,606 Individuals. Am J Hum Genet. 2019;105(2):267-282. doi:10.1016/j.ajhg.2019.05.020
- Epi25 Collaborative. Electronic address: jm4279@cumc.columbia.edu, Epi25 Collaborative. Subgenic intolerance, ClinVar, and the epilepsies: A whole-exome sequencing study of 29,165 individuals. Am J Hum Genet. 2021;108(10):2024. doi:10.1016/j.ajhg.2021.08.008

Posters

- Susgun S., Kesim Y., Salman B., Yucesan E., Khalilov D., Sirin G., Baykan B, Bebek N., Iseri Ugur S., Two candidates bi-allelic variant to neurodevelopmental disorder in a consanguineous family from Turkey, ESHG 2022
- 2. Acar A., Say M., Salman B., Dulger M.V., Comparison of Established Microsatellite Instability Detection Tools in Next Generation Sequencing, ESHG 2021
- 3. Ugur Iseri S., Akçakaya N. H., Salman B., et al., **Exome Sequencing Identifies a FBXO38 Variant**Inherited from a Mosaic Mother to cause Distal Hereditary Neuronopathy Type IID with Distinct
 Features, ASHG 2017

Congresses

1. Investigation of Gene Variants Associated with Syndromic Intellectual Disability, 7. International Congress of Cerebral Palsy and Developmental Disorders, February 23-25 2018, Istanbul, Turkey

projects

1. (in review) TUBITAK 1001 Project, Advanced Molecular Autopsy Protocol for Sudden Deaths under 40 Years Research Fellow

- 2. 2021-2022 TUBITAK 1501 Project, 3210420, Modeling and Automatic Analysis of Genotype-Phenotype Related Copy Number Variations Researcher
- 3. 2021-2022 Research Fund of Istanbul University, ONAP-37862, **Transcriptomics Approaches To Biomarker Potential Of B Cell Expression Analyses In Anti-Neuronal Antibody Related Autoimmune Epilepsy** Researcher
- 4. 2018-2019 Research Fund of Istanbul University, TYL-2018-30315, **Study of Split Hand/Foot Malformation with Genomic Techniques and Bioinformatics Approaches** Researcher
- 5. 2018–2018 TUBITAK 1512 Project, 3210420 **Development of Cloud Based Software for Next Generation Amplicon Sequencing Technologies**, Jan 2018- Oct 2018 Research Fellow
- 6. 2017-2018 Research Fund of Istanbul University, TDP-2017-25510, **Genetic Analyses of Progressive Myoclonic Epilepsy** Researcher
- 7. 2016–2018 TUBITAK 1001 Project, **Investigation of Epileptogenesis After Febrile Seizures**Research Fellow

software

VVC	Variant Version Control (in development) Tool for tracking changes in variant annotation using git.
Dove	Downstream VCF Evaluation Tool for annotating VCF files, multi genome analysis and filtering variants.
Pigeon	Pipelining Genomic Operations Tool for pipelining bioinformatics tools written considering NGS applications.
Picus	Pointed Interpretation of Clinical Variant Significance Tool for classifying sequence variants according to ACMG/AMP criteria.

awards

III. Rare Neurological Diseases Symposium and Neurogenetics Course, **Second place project award**, Istanbul, Turkey, 2017

congresses&courses

1. 2022 Turkish Society of Medical Genetics, Nov 9-13, **Genome-Level Evaluation of Copy Number Changes, One of the Important Risk Factors for Genetic Epilepsy** (Oral Presentation)

- 2. 2020 ESHG 2020, June 6-9, Virtual Conference, **GenerAVI: Variant Interpreter and Genetic Analysis Summary Generator** (Poster Presentation)
- 3. 2019 7th International Congress of the Molecular Biology Association of Turkey, Sep 27-29, Istanbul, Turkey, **PICUS: Pointed Interpretation of Clinical Variant Significance** (Poster Presentation)
- 4. 2019 Bioinformatics Days III, Gazi University Faculty of Medicine, May 25, Ankara, Turkey, **NGS Pipelines with Python** (Oral Presentation)
- 5. 2019 Erciyes University Faculty of Medicine Genetics Days, February 21-23, Kayseri, Turkey, **Making**Sense of Human Genome with Databases and Bioinformatic Tools (Oral Presentation)
- 6. 2018 IV. Rare Neurological Diseases Symposium and Neurogenetics Course, May 31-July 1, Istanbul, Turkey, **Analysis of Neurogenetics Data Interactive Training and Practice** (Instructor)
- 7. 2018 11. National Epilepsy Congress, May 3–6, 2018, Dalaman, Turkey, **Meta Analysis of SCN1A Gene Variants of 114 Patients from Epi25 Exome Data** (Oral Presentation)
- 8. 2018 7. International Congress of Cerebral Palsy and Developmental Disorders, February 23-25, Istanbul, Turkey, Investigation of Gene Variants Associated with Syndromic Intellectual Disability
- 9. 2017 53. National Neurology Congress, November 24–30, Antalya, Turkey, **Genomic Approaches to Intellectual Disability Case with Epilepsy** (Oral Presentation)
- 10. 2017 III. Rare Neurological Diseases Symposium and Neurogenetics Course, July 1–2, Istanbul, Turkey, **Profiling Anti-Epileptic Drug Resistance an Genetic Diagnosis with Epilepsy Panel** (Poster Presentation)

general skills

Languages

Turkish Mother Tongue

English Advanced Fluent

These are the languages I am still learning

French Beginner Basic words and phrases

Japanese BeginnerBasic words and phrases

Latin Beginner Basic words and phrases