



# Barış Salman

**Barış Salman**

**Curriculum Vitae**

**E-mail Phone Website**

**Github Linkedin Biostars Orcid**

## technical skills

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### • Wet lab and Bioinformatics

<b>DNA/RNA/Protein</b>	Isolation, quantative/qualitative PCRs, Blotting, Cytogenetics and Karyotyping	<b>Sequencing</b>	WGS, WES, Panels, RNAseq, scRNAseq, Sanger
<b>Functional Studies</b>	Cell Culture, <i>in vivo</i> models, Microscopy	<b>Micro-arrays</b>	Genotyping, Expression
<b>Genetic Engineering</b>	Recombinant DNA technologies		

### • Computers and Data Analyses

<b>Development</b>	Python, R, Unix Shell	<b>Analysis</b>	Python Scientific Toolkit(numpy, pandas, scipy etc.), R language, Shell Scripting, Bioinformatic tools Machine Learning, PCA, Clustering
<b>Workflow</b>	Linux, Doom Emacs, Git, Jupyter	<b>Statistics and Probability</b>	Hypothesis testing, Bayesian Theorem, Data distributions
<b>Writing</b>	Orgmode, Zotero, Bibtex, $\LaTeX$	<b>Visualization</b>	Matplotlib, Plotly/Dash, $\LaTeX$
<b>Web Development</b>	Django, Javascript		

## education

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<b>Doctorate, 2021–2024</b>	Genetics, Institute of Health Sciences, Istanbul University, Turkey	<b>GPA 3.5</b>
<b>Thesis</b>	Work in progress	
<b>Adviser</b>	Prof. Dr. Sibel Uğur İşeri	
<b>Notable Courses</b>	Advanced Molecular Genetics, Mendelian and non Mendelian Diseases, Chromosomal Diseases, Systems Biology	
<b>Master, 2016–2019</b>	Genetics, Institute of Health Sciences, Istanbul University, Turkey	<b>GPA 2.14</b>
<b>Thesis</b>	Study of Split Hand/Foot Malformation with Genomic Techniques and Bioinformatic Approaches	
<b>Adviser</b>	Assoc Prof. Dr. Sibel Uğur İşeri	
<b>Notable Courses</b>	Bioinformatics, Genetic Counseling, Medical Genetics, Hospital Rotation for Genetic Counseling	
<b>Bachelor, 2011–2016</b>	Molecular Biology and Genetics, Istanbul University, Turkey	<b>GPA 2.44</b>
<b>Thesis</b>	Metabolic Regulation in Prokaryotes with Small Non-Coding RNAs	
<b>Adviser</b>	Asst. Prof. Dr. Semian Karaer Uzuner	
<b>Notable Courses</b>	Methods Used in Molecular Biology(lab), Molecular Genetics, Genetic Engineering	

## jobs

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<b>Company</b>	Refgen Biotechnology	Development of platform for bioinformatics
<b>Date</b>	Mar 2021–Ongoing	Pipeline research and development
<b>Position</b>	Genetics and Bioinformatics Specialist	
<b>Company</b>	Gen-Era Diagnostics	Development of pipelines for NGS
<b>Date</b>	Oct. 2018–Mar 2021	Variant pathogenity classification automatization
<b>Position</b>	Genetics and Bioinformatics Specialist	
<b>Company</b>	Aktif Gen	Development of bioinformatic pipelines
<b>Date</b>	Jan. 2018–Oct 2018	
<b>Position</b>	R&D Personnel	

## • Internships & Rotations

<b>Genetic Counseling</b> rotation	Fall 2018 IU Cerrahpaşa, Department of Pediatric Genetics and Teratology
<b>Genetic Counseling</b> rotation	Spring 2018 IU Cerrahpaşa, Department of Medical Genetics
<b>Cell Culture</b> internship	August 2016 IU Cerrahpasa Department of Medical Biology
<b>Genomics</b> internship	June-July 2015 IU A.S. Institute of Exprmental Medicine Department of Genetics
<b>Cytogenetics</b> internship	June-July 2014 IU Department of Medical Genetics
<b>Molecular Dynamics</b> internship	2012-2013 IU Faculty of Science Quantum Technologies Laboratory

## • Certificates

<b>Use and care of laboratory animals</b>	Mar 21 - Apr 08 IU HADYEK 46. Laboratory Animal Use Training
<b>Liquid handling robot</b>	2-5 November 2021 Hamilton Venus Online Advanced Software Training 9-12 August 2021 Hamilton Venus Online Basic Software Training

## publications

1. Mercan S, Akcakaya NH, **Salman B**, Yapici Z, Ozbek U, Ugur Iseri SA. **Clinical and genetic analyses in syndromic intellectual disability with primary microcephaly reveal biallelic and de novo variants in patients with parental consanguinity.** Genes Genom. Published online November 12, 2022. doi:10.1007/s13258-022-01344-8
2. 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

3. **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
4. 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
5. Akçakaya NH, **Salman B**, Ugur İşeri SA., 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
6. 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**)
- 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

- Epi25 Collaborative

## ● Posters

1. 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

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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

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<b>VVC</b>	Variant Version Control ( <b>in development</b> ) Tool for tracking changes in variant annotation using git.
<b>XΣΔ</b>	Cross-symbol checker This tool checks Ensembl and NCBI annotation files for different genome versions and shows which gene symbol is used. This way a more appropriate gene set can be used to avoid the false negatives in the variant discovery process.
<b>Dove</b>	Downstream VCF Evaluation Tool for annotating VCF files, multi genome analysis and filtering variants.
<b>Pigeon</b>	Pipelining Genomic Operations Tool for pipelining bioinformatics tools written considering NGS applications.
<b>Picus</b>	Pointed Interpretation of Clinical Variant Significance Tool for classifying sequence variants according to ACMG/AMP criteria.

## awards

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

## congresses&courses

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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 and Genetic Diagnosis with Epilepsy Panel** (Poster Presentation)

## general skills

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### • Languages

**Turkish** Mother Tongue

**English** Advanced

Fluent

These are the languages I am still learning

**French** Beginner

Basic words and phrases

**Japanese** Beginner

Basic words and phrases

**Latin** Beginner

Basic words and phrases