Yaser Mansouri

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EDUCATION

Tehran University of Medical Sciences
Tehran, Iran
PhD. Medical Genetics (2018)

Isfahan University of Medical Sciences
Isfahan, Iran
M.SC. Human Genetics (2010)

EXPERIENCES:

Researcher

- Completed 5 years of research experience at Fasa University of Medical Sciences, Noncommunicable Diseases Research Center, Central Research Laboratory from 2018-2023.
- Worked as a Senior Research Scientist and Lab Manager at Central Research Laboratory, Noncommunicable Diseases Research Center, Fasa University of Medical Sciences for 3 years, from 2020-2023.
- Successfully established a molecular diagnosis laboratory at Valiasr Fasa Hospital for COVID-19 molecular diagnosis during the pandemic, Fasa University of Medical Sciences.
- Acquired funding for cancer research at Fasa University of Medical Sciences from 2019-2023.
- Worked as an NGS (Next-Generation Sequencing) data analysis and interpretation specialist (WGS, WES, RNA-Seq, and diagnostic panels) at Shiraz Genetics Institute, Shiraz, Iran, from 2017-2019.
- Contribute to several grant proposal writing that approved by the Ministry of Health and Medical Education (Iran) from 2017-2023.
- Participated as a collaborator in more than 10 research projects approved by the Ministry of Health and Medical Education (Iran) from 2017-2023.
- Successfully supervised over 8 postgraduate theses (PhD, MSc, and MD) during my tenure.
- Conducted interdisciplinary research in collaboration with other departments (e.g. biotechnology, microbiology and biochemistry) and international institutions.
- Active member of Breast Cancer Association (BrCA), Universal Scientific Education and Research Network (USERN) from 2018-2023.
- Member of the Medical Council Organization from 2018-2023.
- Editorial board of Journal of Advanced Biomedical Sciences (JABS)
- Conducted research in evaluating microarray data and gene expression profiling for breast cancer
- Used an integrative bioinformatics approach for system biology-based cancer research
- Identified cell-free diagnostic and prognostic biomarkers for triple negative breast cancers
- Manipulated, analyzed, and presented a high volume of data with SPSS, Prism, Cytoscape, R Studio, and Excel

Responsibilities:

- Worked as the Principal Investigator (PI) of Fasa PERSIAN cohort study, Noncommunicable Diseases Research Center, Fasa University of Medical Sciences from 2022-2023.

Iran University of Medical Sciences
Tehran, Iran
B.Sc. Clinical Laboratory Technology (2008)

SKILLS

- Proficient in molecular genetic laboratory techniques
- Experienced with analyses big data such NGS, Microarray, and RNA-seq
- Statistical and programming Software such SPSS, Prism and RStudio
- Proficient user of bioinformatics software such as Cytoscape
- Background in Cancer System Biology
- Performed extraction of mRNA-miRNA-circRNA regulatory networks by integrative analysis of multiple microarray datasets for different types of cancer, especially Breast Cancer
- Cell culture, Real-time PCR, Tetra-ARMS, Transfection, MTT assay, Western Blot, Flow Cytometry
- Familiar with CRISPR-Cas9 technology
- Proficient user of databases and in-silico tools for non-coding RNAs (circRNAs, miRNAs, and lncRNAs)

- Currently serving as the Genomics Scientist of the Molecular Diagnostic Laboratory at Valiasr Fasa Hospital from 2021.
- Served as a member of the Research Council at Noncommunicable Diseases Research Center, Fasa University of Medical Sciences for 3 years from 2020-2023.
- Served as a Member of Research Ethics Committee from 2022-2023.

Lecturer

During my PhD course at Tehran University of Medical Sciences (2014-2018):

- Presented journal club presentations and delivered high-impact visual reports on project findings.
- Provided scheduled office hours for academic counseling of students.
- Trained staff in wet lab techniques.

At Fasa University of Medical Sciences (Fasa, Iran) from 2018-2023:

- Presented an oral lecture at the 5th international and 17th Iranian Genetics Congress in March 2023.
- Prepared and delivered lectures on Molecular Biology for master's, bachelor's, and medical students from 2018-2023.
- Prepared and delivered lectures on Medical Genetics for master's, bachelor's, and medical students from 2018-2023.

HONORS and AWARDS

- Ranked 1st among university researchers in 2019 and 2022
- Focal point for COVID-19 Genetic Diagnosis at Fasa University of Medical Sciences from 2021-2023
- Received a letter of appreciation for social accountability during the recent pandemic from the Deputy for Curative Affairs of the Ministry of Health and Medical Education
- Received a letter of appreciation for social responsibility during the recent pandemic from the University President
- Ranked third in the national university entrance exam for Ph.D. in Medical Genetics in the year 2014

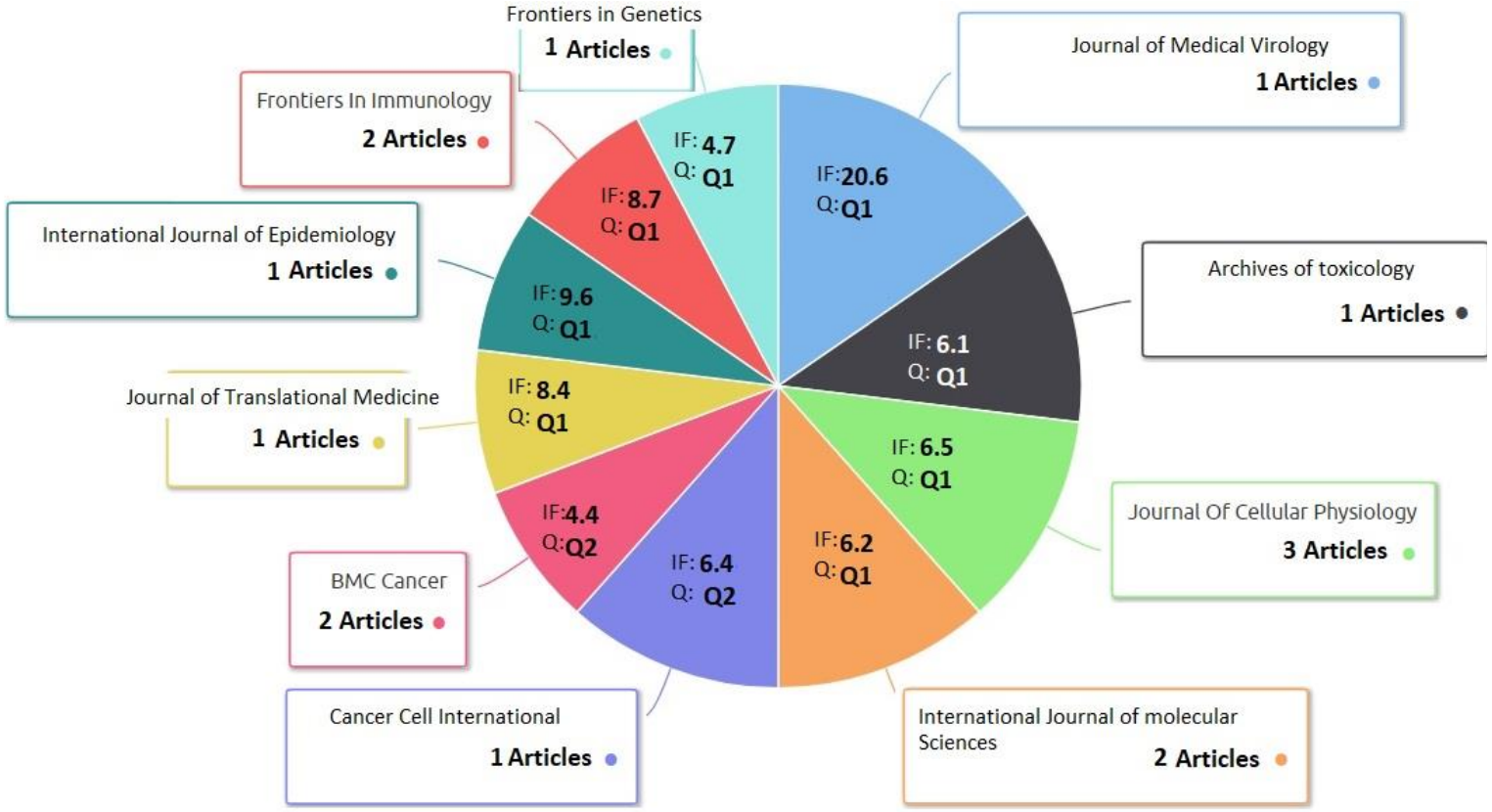
- Team player and comfortable to work independent

RESEARCH INTEREST

- Cancer Molecular signatures
- Personalized Medicine in Cancer
- Cancer Biomarkers
- Cancer Immunotherapy
- Cancer Bioinformatics
- Cancer Epigenetics
- Non-coding RNA network in Cancer

PUBLICATIONS

- Published 13 Articles with International Collaboration (2017-2023)
- Published 37 Original Articles (2017-2023)
- Published 13 Review Articles (2019-2023)



Atlas of My Best Published Articles from 2018 to 2023

1. Evaluation of Circ_0000977-Mediated Regulatory Network in Breast Cancer: A Potential Discriminative Biomarker for Triple-Negative Tumors.

Darbeheshti F, **Mansoori Y**, Azizi-Tabesh G, Zolfaghari F, Kadkhoda S, Rasti A, Rezaei N, Shakoori A. Biochem Genet. 2023 Jan 16. doi: 10.1007/s10528-023-10331-x.

2. Cohort Profile: The Fasa Adults Cohort Study (FACS): a prospective study of non-communicable diseases risks.

Homayounfar R, Farjam M, Bahramali E, Sharafi M, Poustchi H, Malekzadeh R, **Mansoori Y**, Naghizadeh MM, Vakil MK, Dehghan A. Int J Epidemiol. 2023 Jan 2:dyac241. doi: 10.1093/ije/dyac241.

3. Potential roles of NLRP3 inflammasome in the pathogenesis of Kawasaki disease.

Shahi A, Afzali S, Firoozi Z, Mohaghegh P, Moravej A, Hosseinipour A, Bahmanyar M, **Mansoori Y.** J Cell Physiol. 2023 Mar;238(3):513-532. doi: 10.1002/jcp.30948.

4. Investigation of BRCAness associated miRNA-gene axes in breast cancer: cell-free miR-182-5p as a potential expression signature of BRCAness.

Darbeheshti F, Kadkhoda S, Keshavarz-Fathi M, Razi S, Bahramy A, **Mansoori Y**, Rezaei N. BMC Cancer. 2022 Jun 17;22(1):668. doi: 10.1186/s12885-022-09761-4.

5. A Proposed TUSC7/miR-211/Nurr1 ceRNET Might Potentially be Disturbed by a cer-SNP rs2615499 in Breast Cancer.

Abdollahzadeh R, Azarnezhad A, Paknahad S, **Mansoori Y**, Pirhoushiaran M, Kanaani K, Bafandeh N, Jafari D, Tavakkoly-Bazzaz J. Biochem Genet. 2022 Dec;60(6):2200-2225. doi: 10.1007/s10528-022-10216-5.

6. Potential roles of hsa_circ_000839 and hsa_circ_0005986 in breast cancer.

Firoozi Z, Mohammadisoleimani E, Shahi A, Mansoori H, Naghizadeh MM, Bastami M, Nariman-Saleh-Fam Z, Daraei A, Raoofat A, **Mansoori Y.** J Clin Lab Anal. 2022 Mar;36(3):e24263. doi: 10.1002/jcla.24263.

7. Epi-miRNAs: Regulators of the Histone Modification Machinery in Human Cancer.

Mortazavi D, Sohrabi B, Mosallaei M, Nariman-Saleh-Fam Z, Bastami M, **Mansoori Y**, Daraei A, Zununi Vahed S, Navid S, Saadatian Z, Jamialahmadi T, Teng Y, Sahebkar A. J Oncol. 2022 Jan 11;2022:4889807. doi: 10.1155/2022/4889807

8. Purinergic receptors are a key bottleneck in tumor metabolic reprogramming: The prime suspect in cancer therapeutic resistance.

Aria H, Rezaei M, Nazem S, Daraei A, Nikfar G, Mansoori B, Bahmanyar M, Tavassoli A, Vakil MK, **Mansoori Y.** Front Immunol. 2022 Aug 22;13:947885. doi: 10.3389/fimmu.2022.947885.

9. LncRNA-miRNA-mRNA Networks of Gastrointestinal Cancers Representing Common and Specific LncRNAs and mRNAs.

Dastsooz H, Alizadeh A, Habibzadeh P, Nariman A, Hosseini A, **Mansoori Y**, Haghi-Aminjan H. Front Genet. 2022 Jan 24;12:791919. doi: 10.3389/fgene.2021.791919.

10. Dysregulated Expression of miR-146a and Its Associated Immune Effectors in Peripheral Blood Mononuclear Cells of Esophageal Carcinoma Patients.

Nariman-Saleh-Fam Z, **Mansoori Y**, Saadatian Z, Tavakkoly-Bazzaz J, Daraei A, Zununi Vahed S, Mahmoodzadeh H, Bastami M. Immunol Invest. 2022 Feb;51(2):290-300. doi: 10.1080/08820139.2020.1828454.

11. Genetic variations in ATM and H2AX loci contribute to risk of hematological abnormalities in individuals exposed to BTEX chemicals.

Jafari Roshan S, **Mansoori Y**, Hosseini SR, Sabour D, Daraei A.J Clin Lab Anal. 2022 Apr;36(4):e24321. doi: 10.1002/jcla.24321.

12. Anticancer traits of chimeric antigen receptors (CARs)-Natural Killer (NK) cells as novel approaches for melanoma treatment.

Bahmanyar M, Vakil MK, Al-Awsi GRL, Kouhpayeh SA, **Mansoori Y**, Mansoori B, Moravej A, Mazarzaei A, Ghasemian A. BMC Cancer. 2022 Nov 25;22(1):1220. doi: 10.1186/s12885-022-10320-0.

13. Opportunities and obstacles for the melanoma immunotherapy using T cell and chimeric antigen receptor T (CAR-T) applications: a literature review.

Bahmanyar M, Vakil MK, Al-Awsi GRL, Kouhpayeh SA, Mansoori H, **Mansoori Y**, Salahi A, Nikfar G, Tavassoli A, Behmard E, Moravej A, Ghasemian A. Mol Biol Rep. 2022 Nov;49(11):10627-10633. doi: 10.1007/s11033-022-07633-5.

14. Individual genetic variability mainly of Proinflammatory cytokines, cytokine receptors, and toll-like receptors dictates pathophysiology of COVID-19 disease.

Vakil MK, **Mansoori Y**, Al-Awsi GRL, Hosseinipour A, Ahsant S, Ahmadi S, Ekrahi M, Montaseri Z, Pezeshki B, Mohaghegh P, Sohrabpour M, Bahmanyar M, Daraei A, Dadkhah Jouybari T, Tavassoli A, Ghasemian A. J Med Virol. 2022 Sep;94(9):4088-4096. doi: 10.1002/jmv.27849.

15. Upregulation of hsa_circ_0004812 promotes COVID-19 cytokine storm via hsa-miR-1287-5p/IL6R, RIG-I axis.

Mohammadisoleimani E, Firoozi Z, Naghizadeh MM, Ghanbari Asad A, Pezeshki B, Gholampour Y, Mirzaei E, Simin M, Saliani M, Moravej A, Armoon Z, **Mansoori Y.** J Clin Lab Anal. 2022 Oct;36(10):e24666. doi: 10.1002/jcla.24666.

16. Hsa_circ_0000479/Hsa-miR-149-5p/RIG-I, IL-6 Axis: A Potential Novel Pathway to Regulate Immune Response against COVID-19.

Firoozi Z, Mohammadisoleimani E, Shahi A, Naghizadeh MM, Mirzaei E, Asad AG, Salmanpour Z, Javad Nouri SM, **Mansoori Y.** Can J Infect Dis Med Microbiol. 2022 Aug 30;2022:2762582. doi: 10.1155/2022/2762582.

17. Expression analysis of hsa_circ_0020397, hsa_circ_0005986, hsa_circ_0003028, and hsa_circ_0006990 in renal cell carcinoma.

Mohammadisoleimani E, Firoozi Z, Naghizadeh MM, Asad AG, Jafari A, Pourjafarian MH, Ariafar A, Mansoori H, Dastsooz H, Sabaie H, Zeighami S, **Mansoori Y.** Exp Mol Pathol. 2023 Feb;129:104848. doi: 10.1016/j.yexmp.2022.104848.

18. Potential Roles of Long Noncoding RNAs as Therapeutic Targets in Organ Transplantation.
Salehi S, Afzali S, Shahi A, Amirzargar AA, **Mansoori Y.** Front Immunol. 2022 Mar 11;13:835746. doi: 10.3389/fimmu.2022.835746.
19. Applications of Metallic Nanoparticles in the Skin Cancer Treatment.
Marzi M, Osanloo M, Vakil MK, **Mansoori Y**, Ghasemian A, Dehghan A, Zarenezhad E. Biomed Res Int. 2022 Nov 14;2022:2346941. doi: 10.1155/2022/2346941
20. Circular RNA-associated ceRNA network involved in HIF-1 signalling in triple-negative breast cancer: circ_0047303 as a potential key regulator.
Darbeheshti F, Mahdiannasser M, Noroozi Z, Firoozi Z, Mansoori B, Daraei A, Bastami M, Nariman-Saleh-Fam Z, Valipour E, **Mansoori Y.** J Cell Mol Med. 2021 Dec;25(24):11322-11332. doi: 10.1111/jcmm.17066.
21. Expression profiles and functional prediction of long non-coding RNAs LINC01133, ZEB1-AS1 and ABHD11-AS1 in the luminal subtype of breast cancer.
Mehrpour Layeghi S, Arabpour M, Shakoori A, Naghizadeh MM, **Mansoori Y**, Tavakkoly Bazzaz J, Esmaeili R. J Transl Med. 2021 Aug 26;19(1):364. doi: 10.1186/s12967-021-03026-7.
22. Circular RNA hsa_circ_0044234 as distinct molecular signature of triple negative breast cancer: a potential regulator of GATA3.
Darbeheshti F, Zokaei E, **Mansoori Y**, Emadi Allahyari S, Kamaliyan Z, Kadkhoda S, Tavakkoly Bazzaz J, Rezaei N, Shakoori A. Cancer Cell Int. 2021 Jun 14;21(1):312. doi: 10.1186/s12935-021-02015-6.
23. A methylation signature at the CpG island promoter of estrogen receptor beta (ER-β) in breasts of women may be an early footmark of lack of breastfeeding and nulliparity.
Daraei A, Izadi P, Khorasani G, Nafissi N, Naghizadeh MM, Meysamie A, **Mansoori Y**, Nariman-Saleh-Fam Z, Bastami M, Saadatian Z, Roshan SJ, Bayani N, Tavakkoly-Bazzaz J. Pathol Res Pract. 2021 Feb;218:153328. doi: 10.1016/j.prp.2020.153328.
24. Perturbation of miR-146b and relevant inflammatory elements in esophageal carcinoma patients supports an immune downregulatory mechanism.
Bastami M, Mahmoodzadeh H, Saadatian Z, Daraei A, Zununi Vahed S, **Mansoori Y**, Nariman-Saleh-Fam Z. Pathol Res Pract. 2021 Sep;225:153560. doi: 10.1016/j.prp.2021.153560.
25. Critical roles of microRNA-196 in normal physiology and non-malignant diseases: Diagnostic and therapeutic implications.
Bastami M, Masotti A, Saadatian Z, Daraei A, Farjam M, Ghanbariasad A, Vahed SZ, Eyvazi S, **Mansoori Y**, Nariman-Saleh-Fam Z. Exp Mol Pathol. 2021 Oct;122:104664. doi: 10.1016/j.yexmp.2021.104664.
26. Peripheral Blood Mononuclear Cells Expression Levels of miR-196a and miR-100 in Coronary Artery Disease Patients.
Saadatian Z, Nariman-Saleh-Fam Z, Khaheshi I, **Mansoori Y**, Daraei A, Ghaderian SMH, Omrani MD. Immunol Invest. 2021 Nov;50(8):914-924. doi: 10.1080/08820139.2020.1791177.
27. Faculty retention in regional medical schools in Iran: a qualitative content analysis.

Shaterjalali M, Gholampoor Y, Jeihooni AK, **Mansoori Y**, Homayounfar R, Ehrampoush E, Karimi S. BMC Med Educ. 2021 Jan 6;21(1):24. doi: 10.1186/s12909-020-02473-y.

28. Expression and clinicopathological significance of AOC4P, PRNCR1, and PCAT1 lncRNAs in breast cancer.
Abdollahzadeh R, **Mansoori Y**, Azarnezhad A, Daraei A, Paknahad S, Mehrabi S, Tabei MB, Jafari D, Shakoori A, Tavakkoly-Bazzaz J. Pathol Res Pract. 2020 Oct;216(10):153131. doi: 10.1016/j.prp.2020.153131.

29. Competing endogenous RNA (ceRNA) cross talk and language in ceRNA regulatory networks: A new look at hallmarks of breast cancer.
Abdollahzadeh R, Daraei A, **Mansoori Y**, Sepahvand M, Amoli MM, Tavakkoly-Bazzaz J. J Cell Physiol. 2019 Jul;234(7):10080-10100. doi: 10.1002/jcp.27941.

30. Integrative analyses of triple negative dysregulated transcripts compared with non-triple negative tumors and their functional and molecular interactions.
Darbeheshti F, Rezaei N, Amoli MM, **Mansoori Y**, Tavakkoly Bazzaz J. J Cell Physiol. 2019 Dec;234(12):22386-22399. doi: 10.1002/jcp.28804.

31. Breast cancer-linked lncRNA u-Eleanor is upregulated in breast of healthy women with lack or short duration of breastfeeding.
Mansoori Y, Zendehbad Z, Askari A, Kouhpayeh A, Tavakkoly-Bazzaz J, Nariman-Saleh-Fam Z, Bastami M, Saadatian Z, Mansoori B, Yousefvand A, Mansoori H, Daraei A. J Cell Biochem. 2019 Jun;120(6):9869-9876. doi: 10.1002/jcb.28269.

32. Methylation of progesterone receptor isoform A promoter in normal breast tissue: An epigenetic link between early age at menarche and risk of breast cancer?
Daraei A, Izadi P, Khorasani G, Nafissi N, Naghizadeh MM, Younosi N, Meysamie A, **Mansoori Y**, Nariman-Saleh-Fam Z, Bastami M, Saadatian Z, Zendehbad Z, Tavakkoly-Bazzaz J. J Cell Biochem. 2019 Aug;120(8):12393-12401. doi: 10.1002/jcb.28505.

33. Evidences from a systematic review and meta-analysis unveil the role of miRNA polymorphisms in the predisposition to female neoplasms
Bastami M, Choupani J, Saadatian Z, Zununi Vahed S, Ouladsahebmadarek E, **Mansoori Y**, Daraei A, Samadi Kafil H, Yousefi B, Mahdipour M, Masotti A, Nariman-Saleh-Fam Z. Int. J. Mol. Sci. 2019 Oct, 20(20), 5088; https://www.mdpi.com/1422-0067/20/20/5088

34. The intricate role of miR-155 in carcinogenesis: potential implications for esophageal cancer research
Nariman-Saleh-Fam Z, Saadatian Z, Daraei A, **Mansoori Y**, Bastami M, Tavakkoli-Bazzaz J. Biomark Med. 2019 Feb;13(2):147-159. doi: 10.2217/bmm-2018-0127

35. Dysregulated expression of STAT1, miR-150, and miR-223 in peripheral blood mononuclear cells of coronary artery disease patients with significant or insignificant stenosis

Zahra Saadatian, Nariman-Saleh-Fam Z, Bastami M, **Mansoori Y**, Khaheshi I, Alipour Parsa S, Daraei A, Zununi Vahed S, Yousefi B, Samadi Kafil H, Eyvazi S, Ghaderian SMH, Omrani MD. Journal of cellular biochemistry. 2019 Jul;13(2):147-159. doi.org/10.1002/jcb.29286

36. Expression pattern of miR-21, miR-25 and PTEN in peripheral blood mononuclear cells of patients with significant or insignificant coronary stenosis.

Nariman-Saleh-Fam Z, Vahed SZ, Aghaee-Bakhtiari SH, Daraei A, Saadatian Z, Kafil HS, Yousefi B, Eyvazi S, Khaheshi I, Parsa SA, Moravej A, Mousavi N, Bastami M, **Mansoori Y.** Gene. 2019 May 25;698:170-178. doi: 10.1016/j.gene.2019.02.074.

37. miRNA Polymorphisms and Risk of Cardio-Cerebrovascular Diseases: A Systematic Review and Meta-Analysis.

Bastami M, Choupani J, Saadatian Z, Zununi Vahed S, **Mansoori Y**, Daraei A, Samadi Kafil H, Masotti A, Nariman-Saleh-Fam Z. Int J Mol Sci. 2019 Jan 12;20(2):293. doi: 10.3390/ijms20020293.

38. The Effect of Mesenchymal Stem Cells on the Expression of IDO and Qa2 Molecules in Dendritic Cells.

Moravej A, Kouhpayeh A, Geramizadeh B, Azarpira N, Yaghobi R, **Mansoori Y**, Karimi MH. Adv Pharm Bull. 2019 Feb;9(1):56-63. doi: 10.15171/apb.2019.007.

39. Expression levels of breast cancer-related GAS5 and LSINCT5 lncRNAs in cancer-free breast tissue: Molecular associations with age at menarche and obesity.

Mansoori Y, Tabei MB, Askari A, Izadi P, Daraei A, Bastami M, Naghizadeh MM, Nariman-Saleh-Fam Z, Mansoori B, Tavakkoly-Bazzaz J. Breast J. 2018 Nov;24(6):876-882. doi: 10.1111/tbj.13067.

40. A link between expression level of long-non-coding RNA *ZFAS1* in breast tissue of healthy women and obesity.

Mansoori Y, Tabei MB, Askari A, Izadi P, Daraei A, Naghizadeh MM, Zendehbad Z, Bastami M, Nariman-Saleh-Fam Z, Mansoori H, Tavakkoly-Bazzaz J. Int J Biol Markers. 2018 Nov;33(4):500-506. doi: 10.1177/1724600818762258.

41. Association between polymorphism of GSTP1, GSTT1, GSTM1 and CYP2E1 genes and susceptibility to benzene-induced hematotoxicity.

Nourozi MA, Neghab M, Bazzaz JT, Nejat S, **Mansoori Y**, Shahtaheri SJ. Arch Toxicol. 2018 Jun;92(6):1983-1990. doi: 10.1007/s00204-017-2104-9.

42. Association of VDR gene polymorphisms with risk of relapsing-remitting multiple sclerosis in an Iranian Kurdish population.

Abdollahzadeh R, Moradi Pordanjani P, Rahmani F, Mashayekhi F, Azarnezhad A, **Mansoori Y.** Int J Neurosci. 2018 Jun;128(6):505-511. doi: 10.1080/00207454.2017.1398158.

43. Effects of Genetic Polymorphism on Susceptibility to Nephrotoxic Properties of BTEXs Compounds.

Neghab M, Nourozi MA, Shahtaheri SJ, **Mansoori Y**, Bazzaz JT, Nedjat S. J Occup Environ Med. 2018 Aug;60(8):e377-e382. doi: 10.1097/JOM.0000000000001364.

44. Epigenetic Changes of the ESR1 Gene in Breast Tissue of Healthy Women: A Missing Link with Breast Cancer Risk Factors?

Daraei A, Izadi P, Khorasani G, Nafissi N, Naghizadeh MM, Younosi N, Meysamie A, **Mansoori Y**, Bastami M, Tavakkoly-Bazzaz J. Genet Test Mol Biomarkers. 2017 Aug;21(8):464-470. doi: 10.1089/gtmb.2017.0028.

45. miRNA-Related Polymorphisms in miR-423 (rs6505162) and PEX6 (rs1129186) and Risk of Esophageal Squamous Cell Carcinoma in an Iranian Cohort.
Nariman-Saleh-Fam Z, Bastami M, Somi MH, Behjati F, **Mansoori Y**, Daraei A, Saadatian Z, Nariman-Saleh-Fam L, Mahmoodzadeh H, Makhdoumi Y, Tabrizi FV, Ebrahimi-Sharif B, Hezarian A, Naghashi S, Abbaszadegan MR, Tavakkoly-Bazzaz J. Genet Test Mol Biomarkers. 2017 Jun;21(6):382-390. doi: 10.1089/gtmb.2016.0346.
46. The SDF1 A/G Gene Variant: A Susceptibility Variant for Myocardial Infarction.
Mansoori Y, Daraei A, Zendebad Z, Madadizadeh F, Mansoori B, Naghizadeh MM, Darbeheshti F. Genet Test Mol Biomarkers. 2017 Aug;21(8):506-511. doi: 10.1089/gtmb.2017.0023.
47. Influences of IL-1b-3953 C>T and MMP-9-1562C >T Gene Variants on Myocardial Infarction Susceptibility in a Subset of the Iranian Population.
Daraei A, **Mansoori Y**, Zendebad Z, Tavakkoly-Bazzaz J, Madadizadeh F, Naghizadeh MM, Arghavani A, Mansoori B. Genet Test Mol Biomarkers. 2017 Jan;21(1):33-38. doi: 10.1089/gtmb.2016.0240.