

# FaceSNPs: Identifying Face-Related SNPs from the Human Genome - Supplementary Materials

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## I. SUPPLEMENT A

Table I shows a sample of the genes we found through our automated literature search strategy. For each gene, we show its function as reported by the relevant literature and provide references.

### REFERENCES

- [1] S. Shen *et al.*, "Seven-cpg-based prognostic signature coupled with gene expression predicts survival of oral squamous cell carcinoma," *Clinical Epigenetics*, vol. 9, no. 1, p. 88, 2017.
- [2] L. Tosca *et al.*, "Genotype-phenotype correlation in 13q13.3-q21.3 deletion," *European Journal of Medical Genetics*, vol. 54, no. 5, e489-e494, 2011.
- [3] N. F. Pedro *et al.*, "Candidate biomarkers for oral squamous cell carcinoma: Differential expression of oxidative stress-related genes," *Asian Pacific journal of cancer prevention : APJCP*, vol. 19, no. 5, pp. 1343-1349, 2018, PMC6031819[pmcid].
- [4] N. Han, Z. Chen, and Q. Zhang, "Expression of klf5 in odontoblastic differentiation of dental pulp cells during in vitro odontoblastic induction and in vivo dental repair," *International Endodontic Journal*, vol. 50, no. 7, pp. 676-684, 2017.
- [5] C. Chen *et al.*, "Genetic variants of mgmt, rhpn2, and fam49a contributed to susceptibility of nonsyndromic orofacial clefts in a chinese population," *Journal of Oral Pathology & Medicine*, vol. 47, no. 8, pp. 796-801, 2018.
- [6] R. W. J. Collin *et al.*, "Mutations of esrrb encoding estrogen-related receptor beta cause autosomal-recessive nonsyndromic hearing impairment dfnb35," *American journal of human genetics*, vol. 82, no. 1, pp. 125-138, 2008, S0002-9297(07)00014-6[PII].
- [7] A. L. Fletcher *et al.*, "Observations regarding retinopathy in mitochondrial trifunctional protein deficiencies," *Molecular Genetics and Metabolism*, vol. 106, no. 1, pp. 18-24, 2012.
- [8] W. D'Souza, S. Pradhan, and D. Saranath, "Multiple single nucleotide polymorphism analysis and association of specific genotypes in fhit, samd4a, and ankrd17 in indian patients with oral cancer," *Head & Neck*, vol. 39, no. 8, pp. 1586-1595, 2017.
- [9] M. Crompton *et al.*, "A mutation in nischarin causes otitis media via limk1 and nf- $\kappa$ b pathways," *PLOS Genetics*, vol. 13, no. 8, pp. 1-28, 2017.
- [10] R. L. Shah *et al.*, "A genome-wide association study of corneal astigmatism: The cream consortium," *Molecular vision*, vol. 24, pp. 127-142, 2018, PMC5800430[pmcid].
- [11] L. Huang *et al.*, "Association of coding and utr variants in the known regions with wet age-related macular degeneration in han chinese population," *Journal of Human Genetics*, vol. 63, no. 10, pp. 1055-1070, 2018.
- [12] N. Urraca *et al.*, "Significant transcriptional changes in 15q duplication but not angelman syndrome deletion stem cell-derived neurons," *Molecular Autism*, vol. 9, no. 1, p. 6, 2018.
- [13] X. Li *et al.*, "Genome-wide linkage study suggests a susceptibility locus for isolated bilateral microtia on 4p15.32-4p16.2," *PLOS ONE*, vol. 9, no. 7, pp. 1-8, 2014.
- [14] J. S. Bae *et al.*, "Genetic association analysis of ciita variations with nasal polyp pathogenesis in asthmatic patients," *Molecular Medicine Reports*, vol. 7, no. 3, pp. 927-934, 2013.
- [15] J. L. Wiggs and L. R. Pasquale, "Genetics of glaucoma," *Human Molecular Genetics*, vol. 26, no. R1, R21-R27, 2017.
- [16] M.-J. An *et al.*, "Transcriptome analysis for UVB-induced phototoxicity in mouse retina," *Environmental Toxicology*, vol. 33, no. 1, pp. 52-62, 2018.
- [17] M. Hayashi and T. Suzuki, "Oculocutaneous albinism type 4," in *GeneReviews® [Internet]*, M. Adam *et al.*, Eds., Updated 2017 Sep 7, Seattle (WA): University of Washington, Seattle, pp. 1993-2019.
- [18] Y. Peng *et al.*, "Downregulation and aberrant localization of forkhead box j1 in allergic nasal mucosa," *International Archives of Allergy and Immunology*, vol. 176, no. 2, pp. 115-123, 2018.
- [19] Deciphering Developmental Disorders Study, "Large-scale discovery of novel genetic causes of developmental disorders," *Nature*, vol. 519, no. 7542, pp. 223-228, 2015, nature14135[PII].
- [20] H. Kasama *et al.*, "Adenosine a2b receptor promotes progression of human oral cancer," *BMC Cancer*, vol. 15, no. 1, p. 563, 2015.
- [21] C. Halgren *et al.*, "Haploinsufficiency of celf4 at 18q12.2 is associated with developmental and behavioral disorders, seizures, eye manifestations, and obesity," *European Journal Of Human Genetics*, vol. 20, 1315 EP -, 2012, Short Report.
- [22] K. Ma *et al.*, "Clinical features and linkage analysis for a chinese family with autosomal dominant central areolar choroidal dystrophy," *Chinese Medical Journal*, vol. 122, no. 22, 2009.
- [23] B. Coste *et al.*, "Gain-of-function mutations in the mechanically activated ion channel piezo2 cause a subtype of distal arthrogyrosis," *Proceedings of the National Academy of Sciences*, vol. 110, no. 12, pp. 4667-4672, 2013.
- [24] M. Rodríguez Pulido *et al.*, "Foot-and-mouth disease virus infection induces proteolytic cleavage of ptb, eif3a,b, and pabp rna-binding proteins," *Virology*, vol. 364, no. 2, pp. 466-474, 2007.
- [25] A. Sultana *et al.*, "Olfactomedin 2: Expression in the Eye and Interaction with Other Olfactomedin Domain-Containing Proteins," *Investigative Ophthalmology & Visual Science*, vol. 52, no. 5, pp. 2584-2592, 2011.
- [26] K. Szczaluba *et al.*, "High-resolution array comparative genomic hybridization utility in polish newborns with isolated cleft lip and palate," *Neonatology*, vol. 107, no. 3, pp. 173-178, 2015.
- [27] Y. Chen, P. Sternberg, and J. Cai, "Characterization of a Bcl-XL-Interacting Protein FKBP8 and Its Splice Variant in Human RPE Cells," *Investigative Ophthalmology & Visual Science*, vol. 49, no. 4, pp. 1721-1727, 2008.
- [28] S. G. Younkin *et al.*, "A genome-wide study of inherited deletions identified two regions associated with nonsyndromic isolated oral clefts," *Birth Defects Research Part A: Clinical and Molecular Teratology*, vol. 103, no. 4, pp. 276-283, 2015.
- [29] G. Jedraszak *et al.*, "Clinical and molecular characterization of the 20q11.2 microdeletion syndrome: Six new patients," *American Journal of Medical Genetics Part A*, vol. 167, no. 3, pp. 504-511, 2015.
- [30] Z. W. E. Yong *et al.*, "Genetic alterations of chromosome 8 genes in oral cancer," *Scientific Reports*, vol. 4, 6073 EP -, 2014, Article.
- [31] G. A. Stamatou and K. M. Stankovic, "A comprehensive network and pathway analysis of human deafness genes," *Otology & Neurotology*, vol. 34, no. 5, 2013.
- [32] C. Luna *et al.*, "Regulation of trabecular meshwork cell contraction and intraocular pressure by mir-200c," *PLOS ONE*, vol. 7, no. 12, pp. 1-9, 2012.

- [33] H. Huang *et al.*, "Expression of microRNA-10a, microRNA-342-3p and their predicted target gene *tiam1* in extranodal nk/t-cell lymphoma, nasal type," *Oncology letters*, vol. 11, no. 1, pp. 345–351, 2016, OL-0-0-3831[PII].
- [34] G. Giroto *et al.*, "Hearing function and thresholds: A genome-wide association study in european isolated populations identifies new loci and pathways," *Journal of Medical Genetics*, vol. 48, no. 6, pp. 369–374, 2011.
- [35] C.-M. Chung *et al.*, "Combined genetic biomarkers and betel quid chewing for identifying high-risk group for oral cancer occurrence," *Cancer Prevention Research*, vol. 10, no. 6, pp. 355–362, 2017.
- [36] D. Fine *et al.*, "A syndrome of congenital microcephaly, intellectual disability and dysmorphism with a homozygous mutation in *frmd4a*," *European Journal Of Human Genetics*, vol. 23, 1729 EP -, 2014, Short Report.
- [37] Y.-Z. Kuo *et al.*, "Mir-99a exerts anti-metastasis through inhibiting myotubularin-related protein 3 expression in oral cancer," *Oral Diseases*, vol. 20, no. 3, e65–e75, 2014.
- [38] N. Okamoto *et al.*, "Deletion at chromosome 10p11.23-p12.1 defines characteristic phenotypes with marked midface retrusion," *Journal Of Human Genetics*, vol. 57, 191 EP -, 2012, Original Article.
- [39] V. R. M. Chavali *et al.*, "Association of oct derived drusen measurements with amd associated-genotypic snps in amish population," *Journal of clinical medicine*, vol. 4, no. 2, pp. 304–317, 2015, PMC4398021[pmcid].
- [40] A. T. Fung *et al.*, "New *best1* mutations in autosomal recessive bestrophinopathy," *Retina (Philadelphia, Pa.)*, vol. 35, no. 4, pp. 773–782, 2015, PMC4425987[pmcid].
- [41] J.-D. Cha, H. J. Kim, and I.-H. Cha, "Genetic alterations in oral squamous cell carcinoma progression detected by combining array-based comparative genomic hybridization and multiplex ligation-dependent probe amplification," *Oral Surgery, Oral Medicine, Oral Pathology, Oral Radiology and Endodontics*, vol. 111, no. 5, pp. 594–607, 2011.
- [42] M. L. Kwee *et al.*, "An autosomal dominant high bone mass phenotype in association with craniosynostosis in an extended family is caused by an *Irp5* missense mutation," *Journal of Bone and Mineral Research*, vol. 20, no. 7, pp. 1254–1260, 2005.
- [43] S. David *et al.*, "Hypotonia and delayed motor development as an early presentation of lowe syndrome: Case report and literature review," *Acta Clinica Belgica*, vol. 0, no. 0, pp. 1–5, 2018, PMID: 30501482.
- [44] K. Duś-Szachniewicz *et al.*, "Protein tyrosine phosphatase receptor r and z1 expression as independent prognostic indicators in oral squamous cell carcinoma," *Head & Neck*, vol. 37, no. 12, pp. 1816–1822, 2015.
- [45] A. G. Fincham *et al.*, "Isolation and partial characterization of a human amelogenin from a single fetal dentition using HPLC techniques," *Calcified tissue international*, vol. 47, no. 2, pp. 105–111, 1990.
- [46] A. M. Heikkinen *et al.*, "Pilot study on the genetic background of an active matrix metalloproteinase-8 test in finnish adolescents," *Journal of Periodontology*, vol. 88, no. 5, pp. 464–472, 2017.
- [47] J. L. Llorente *et al.*, "Nasosinusal adenocarcinoma: Molecular and genetic analysis by MLPA," *Acta Otorrinolaringologica (English Edition)*, vol. 59, no. 4, pp. 151–158, 2008.
- [48] National Center for Biotechnology Information, *Gene*, <https://www.ncbi.nlm.nih.gov/gene>.

TABLE I  
MANUAL LITERATURE VALIDATION OF SELECTED GENES

Chr	Genes	Function (from literature)	Region	Ref	Chr	Genes	Function (from literature)	Region	Ref
1	AJAPI	Associated with Oral Squamous Cell Carcinoma (OSCC)	mouth	[1]	13	DCLK1	Associated with macrocephaly, high forehead, hypertelorism, large nose, large and malformed ears and retrognathia	all	[2]
	DHCR24	Down-regulated in Oral Squamous Cell Carcinoma cases	mouth	[3]		KLF5	Associated with odontoblastic differentiation of dental pulp cells	mouth	[4]
2	FAM49A	Associated with nonsyndromic orofacial clefts	mouth	[5]	14	ESRRB	Essential for inner-ear development and function	ear	[6]
	HADHB	Associated with retinopathy	eye	[7]		SAMD4A	Associated with oral cancer	mouth	[8]
3	NISCH	Mutation associated with chronic otitis media	ear	[9]	15	TNFAIP8L3	Candidate gene for corneal astigmatism	eye	[10]
	BBX	Associated with age-related macular degeneration	eye	[11]		HERC2	Associated with Angelman Syndrome	face shape, forehead	[12]
4	EVC2	Association with microtia, Ellis-van Creveld syndrome or Weyers acrofacial dysostosis	ear, mouth, face shape	[13]	16	CIITA	Associated with nasal polyposis	nose	[14]
	AFAP1	Associated with primary-open-angle glaucoma	eye	[15]		RNF40	May affect how the retina absorbs UV radiations	eye	[16]
5	SLC45A2	Associated with Oculocutaneous Albinism Type 4	eye	[17]	17	DNAH9	Associated with allergic rhinitis	nose	[18]
	COL4A3BP	Associated with several developmental disabilities	face shape	[19]		ADORA2B	May be a key regulator of tumoral progression in OSCCs	mouth	[20]
6	GMDS	Associated with primary-open-angle glaucoma	eye	[15]	18	CELF4	An important role in eye development	eye	[21]
	RIMS1	Associated with Central Areolar Choroidal Dystrophy	eye	[22]		PIEZO2	Mutation can cause a subtype of Distal Arthrogyposis Type 5	eye	[23]
7	EIF3B	Plays a role in Foot-and-mouth disease virus infection	mouth	[24]	19	OLFM2	Associated with glaucoma	eye	[25]
	CHN2	Candidate gene for cleft lip and palate	mouth	[26]		FKBP8	Associated with loss of retinal pigment epithelial cells	eye	[27]
8	ADAM3A	May be involved in the etiology of oral clefts.	mouth	[28]	20	GDF5	Associated with craniofacial dysmorphism	forehead	[29]
	CSGALNACT1	Associated with Oral Squamous Cell Carcinoma	mouth	[30]		HNF4A	Associated with genetic hearing loss	ear	[31]
9	LPAR1	Affects regulation of trabecular meshwork (TM) cell networks	eye	[32]	21	TIAM1	May contribute to the pathogenesis of extranodal natural killer (NK)/T-cell lymphoma, nasal type (ENKTCL)	nose	[33]
	PTPRD	Affects hearing functions	ear	[34]		HSPA13	Associated with OSCC occurrence	mouth	[35]
10	FRMD4A	Associated with congenital microcephaly	face shape	[36]	22	MTMR3	Possible therapeutic target for oral cancer treatment	mouth	[37]
	MPP7	Potentially associated with maxillofacial abnormalities	mouth	[38]		SYN3	Associated with larger area perifoveal ring and larger drusen volume	eye	[39]
11	BEST1	Mutations cause autosomal recessive bestrophinopathy	eye	[40]	X	CHRD1	Associated with Oral Squamous Cell Carcinoma	mouth	[41]
	LRP5	Missense mutation cause macrocephaly, craniocystosis	all	[42]		OCRL	Associated with Lowe syndrome	eye	[43]
12	PTPRR	Potentially associated with oral cancer	mouth	[44]	Y	AMELY	Associated with tooth enamel development	mouth	[45]
	VDR	Polymorphisms linked to initial periodontitis	mouth	[46]		UTY	Associated with nasosinus adenocarcinoma	nose	[47]

NB: The functions are summaries from *NCBI Gene* [48]. The references our system relied on to associate the genes with the face regions are not listed due to space limitation, but are available from [http://community.wvu.edu/~daadjero/~/projects/faceSNPs/FaceSNPs\\_BIBM2019suppl.pdf](http://community.wvu.edu/~daadjero/~/projects/faceSNPs/FaceSNPs_BIBM2019suppl.pdf)