

**Supplementary information to:**

**Original article:**

**ASSOCIATION BETWEEN 15 INSERTION/DELETION GENETIC POLYMORPHISMS AND RISK OF SCHIZOPHRENIA USING POOLED SAMPLES**

Maedeh Bordbar<sup>ID</sup>, Mostafa Saadat\*<sup>ID</sup>

Department of Biology, College of Sciences, Shiraz University, Shiraz 71467-13565, Iran

\* **Corresponding author:** Mostafa Saadat, Department of Biology, College of Sciences, Shiraz University, Shiraz 71467-13565, Iran. Fax: +98-71-32280926.  
E-mail: [saadat@shirazu.ac.ir](mailto:saadat@shirazu.ac.ir)

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**Supplementary Table 1:** The specific forward and reverse primers and PCR product length of each indel polymorphism

Loci	Primer sequences	PCR product length (bp)	Reference
<b>APOB</b>	<b>F:</b> 5' ACCGGCCCTGGCGCCCCGCCAGCA 3' <b>R:</b> 5' CAGCTGGCGATGGACCCGCCGA 3'	93/84	Boerwinkle and Chan, 1989
<b>ADRA2B</b>	<b>F:</b> 5' CAAGCTGAGGCCGGAGACACT 3' <b>R:</b> 5' AGGGTGTGTTGTGGGCATCT 3'	112/103	Eldeeb et al., 2021
<b>PDCD6I</b>	<b>F:</b> 5' CCTTAAGGTCTGTGTCAACC 3' <b>R:</b> 5' TATTCCCTCCACTCGAACAAAC 3'	227/212	Liu et al., 2014
<b>LRPAP1</b>	<b>F:</b> 5' AGTGTGCGTGGAGCCTATG 3' <b>R:</b> 5' GGTGTTCTGGACACAAAGGA 3'	222/185	Benes et al., 2000
<b>TLR2</b>	<b>F:</b> 5' CACGGAGGCAGCGAGAAA 3' <b>R:</b> 5' CTGGGCCGTGCAAAGAAG 3'	286/264	Tahara et al., 2007
<b>DHF</b>	<b>F:</b> 5' CGCAAGTCTGGCCCCATC 3' <b>R:</b> 5' TCAGGTATCTGCCGGGCC 3'	119/100	Hayashi et al., 2010
<b>VEGF</b>	<b>F:</b> 5' GCTGAGGATGGGGCTGACTAGGTA 3' <b>R:</b> 5' GTTCTGACCTGGCTATTCCAGG 3'	230/212	Buraczynska et al., 2007
<b>HLA-G</b>	<b>F:</b> 5' GTGATGGGCTTTAAAGTGTCAACC 3' <b>R:</b> 5' GGAAGGAATGCAGTCAGCATGA 3'	224/210	Hviid et al., 1999
<b>TPA</b>	<b>F:</b> 5' GTAAGAGTCCGTAACAGGACAGCT 3' <b>R:</b> 5' CCCCACCCCTAGGAGAACTTCTCTTT 3'	424/113	Valle-Garay et al., 2013
<b>DBH</b>	<b>F:</b> 5' TGCAAAAATCAGGCACATGC 3' <b>R:</b> 5' TCCAATAATTGGCCTCAATC 3'	166/147	Barbanti et al., 2019
<b>UCP2</b>	<b>F:</b> 5' CAGTGAGGGAAGTGGGAGG 3' <b>R:</b> 5' GGGCAGGACGAAGATT 3'	502/457	Wang et al., 2007
<b>FADS2</b>	<b>F:</b> 5' TTTCTCAAAGGCCGTGGTGT 3' <b>R:</b> 5' AGTGCTAACCACTCCTGGAA 3'	629/607	Kothapalli et al., 2016
<b>MDM2</b>	<b>F:</b> 5' TTTCTTTCTGGTAGGCTGG 3' <b>R:</b> 5' CACCTACTTCCCACAGAGA 3'	262/222	Lalonde et al., 2012
<b>TP53</b>	<b>F:</b> 5' TCAAATCATCCATTGCTTGG 3' <b>R:</b> 5' TGGGACTGACTTCTGCTCTT 3'	195/179	Wu et al., 2002
<b>SLC6A4</b>	<b>F:</b> 5' GACATAATCTGTCTTCTGGCCTCTCAAG 3' <b>R:</b> 5' CAATGTCTGGCGCTTCCCACATAT 3'	310/266	Hauser et al., 2003

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