

Supplementary table S1

The GWAS data about associations of the studied candidate genes polymorphisms with blood pressure/hypertension, some cardiovascular diseases and anthropometric characteristics

SNP of gene (position (hg38))	Phenotype	Association (significance) (affected allele)	Reference
rs1173771 <i>AC026703.1</i> (5: 32814922)	SBP	$\beta = 0.50 (p = 1.8 \times 10^{-16})$ (G)	International Consortium et al., 2011
	DBP	$\beta = 0.26 (p = 9.1 \times 10^{-12})$ (G)	
	Hypertension	$\beta = 0.06 (p = 3.2 \times 10^{-10})$ (G)	
	MAP	$\beta = 0.28 (p = 3.51 \times 10^{-9})$ (G)	
	PP	$\beta = 0.28 (p = 4.56 \times 10^{-9})$ (G)	Wain L. V., et al., 2011
	MAP	$\beta = -0.35 (p = 3.19 \times 10^{-8})$ (A)	
	Hip circumference (sex-combined)	$\beta = 0.03 (p = 6.13 \times 10^{-13})$ (A)	Shungin D., et al., 2015
	Hip circumference (men)	$\beta = 0.03 (p = 6.19 \times 10^{-8})$ (A)	
	SBP	$\beta = -0.43 (p = 2.0 \times 10^{-28})$ (A)	Hoffmann T. J. et al., 2017
	PP	$\beta = -0.25 (p = 1.0 \times 10^{-21})$ (A)	
	DBP	$\beta = -0.19 (p = 8.0 \times 10^{-16})$ (A)	
	Height	$\beta = 0.04 (p = 9.0 \times 10^{-8})$ (A)	
	SBP	$\beta = 0.51 (p = 3.20 \times 10^{-12})$ (G)	Wain L. V., et al., 2017
	PP	$\beta = 0.28 (p = 2.36 \times 10^{-9})$ (G)	
	SBP (smoking interaction) (European)	$p = 1.74 \times 10^{-38}$	Sung Y. J., et al., 2018
	DBP (smoking interaction) (European)	$p = 3.62 \times 10^{-22}$	
	SBP (smoking interaction) (trans-ethnic data)	$p = 8.25 \times 10^{-44}$	
	DBP (smoking interaction) (trans-ethnic data)	$p = 2.75 \times 10^{-28}$	
	SBP	$\beta = -0.42 (p = 4.19 \times 10^{-8})$ (A)	Takeuchi F., et al., 2018
	MAP	$\beta = -0.29 (p = 5.39 \times 10^{-8})$ (A)	
rs1799945 <i>HFE</i> (6: 26090951)	DBP	$\beta = 0.46 (p = 1.5 \times 10^{-15})$ (G)	International Consortium et al., 2011
	SBP	$\beta = 0.62 (p = 7.7 \times 10^{-12})$ (G)	
	Hypertension	$\beta = 0.10 (p = 1.8 \times 10^{-10})$ (G)	
	DBP	$\beta = -0.43 (p = 3.1 \times 10^{-16})$ (C)	Ehret G. B. et al., 2016
	SBP	$\beta = -0.60 (p = 3.28 \times 10^{-12})$ (C)	
	DBP	$\beta = -0.29 (p = 1.0 \times 10^{-18})$ (C)	Hoffmann T. J. et al., 2017

	SBP	$\beta = -0.36 (p = 4.0 \times 10^{-11})$ (C)	Liu C. et al., 2016
	DBP	$\beta = 0.52 (p = 9.2 \times 10^{-18})$ (G)	
	MAP	$\beta = 0.58 (p = 1.0 \times 10^{-17})$ (G)	
	SBP	$\beta = 0.70 (p = 6.2 \times 10^{-13})$ (G)	
	Hypertension	Z score = 0.039 ($p = 3.7 \times 10^{-9}$) (G)	Surendran P. et al., 2016
	DBP (trans-ethnic data)	$\beta = -0.03 (p = 1.26 \times 10^{-19})$ (C)	
	DBP (European)	$\beta = -0.03 (p = 9.82 \times 10^{-10})$ (C)	
	DBP	$\beta = 0.47 (p = 8.87 \times 10^{-14})$ (G)	Wain L. V. et al., 2017
	SBP	$\beta = 0.63 (p = 7.63 \times 10^{-10})$ (G)	
	DBP (smoking interaction) (European)	$p = 1.51 \times 10^{-31}$	Sung Y. J. et al., 2018
	SBP (smoking interaction) (European)	$p = 8.46 \times 10^{-17}$	
	DBP (smoking interaction) (trans-ethnic data)	$p = 3.77 \times 10^{-33}$	
	SBP (smoking interaction) (trans-ethnic data)	$p = 7.43 \times 10^{-19}$	
	SBP	$\beta = 0.37 (p = 1.5 \times 10^{-11})$ (G)	International Consortium et al., 2011
rs805303 <i>BAG6/BAT2-BAT5</i> (6: 31648589)	DBP	$\beta = 0.23 (p = 3.0 \times 10^{-11})$ (G)	
	Hypertension	$\beta = 0.05 (p = 1.1 \times 10^{-10})$ (G)	
	SBP (European)	$\beta = -0.02 (p = 3.02 \times 10^{-6})$ (A)	Surendran P. et al., 2016
	SBP	$\beta = 0.48 (p = 7.1 \times 10^{-16})$ (G)	International Consortium et al., 2011
rs932764 <i>PLCE1</i> (10: 94136183)	Hypertension	$\beta = 0.06 (p = 9.4 \times 10^{-9})$ (G)	
	SBP	$\beta = -0.50 (p = 6.88 \times 10^{-17})$ (A)	Ehret G. B., et al., 2016
	DBP	$\beta = -0.22 (p = 6.28 \times 10^{-10})$ (A)	
	SBP	$\beta = -0.30 (p = 1.0 \times 10^{-14})$ (A)	Hoffmann T. J. et al., 2016
	PP	$\beta = -0.16 (p = 3.0 \times 10^{-10})$ (A)	
	PP	$\beta = 0.26 (p = 5.67 \times 10^{-8})$ (G)	Wain L. V., et al., 2017
	DBP (European)	$\beta = 0.22 (p = 5.55 \times 10^{-8})$ (A)	Surendran P. et al., 2016
rs4387287 <i>OBFC1</i> (10:103918139)	SBP (European)	$\beta = 0.34 (p = 2.21 \times 10^{-7})$ (A)	
	DBP (trans-ethnic data)	$\beta = 0.22 (p = 4.21 \times 10^{-10})$ (A)	
	SBP (trans-ethnic data)	$\beta = 0.36 (p = 9.12 \times 10^{-10})$ (A)	
	Hypertension (trans-ethnic data)	Z score = 5.58 ($p = 2.37 \times 10^{-8}$) (A)	
	SBP	$\beta = -0.57 (p = 1.2 \times 10^{-17})$ (G)	International Consortium et al., 2011
rs633185 <i>ARHGAP42</i>	DBP	$\beta = -0.33 (p = 2.0 \times 10^{-15})$ (G)	

(11:100722807)	Hypertension	$\beta = -0.07 (p = 6.4 \times 10^{-11})$ (G)	Ehret G. B., et al., 2016
	SBP	$\beta = 0.52 (p = 6.97 \times 10^{-15})$ (C)	
	DBP	$\beta = 0.29 (p = 2.38 \times 10^{-12})$ (C)	
	SBP	$\beta = -0.49 (p = 1.0 \times 10^{-31})$ (G)	
	DBP	$\beta = -0.27 (p = 1.0 \times 10^{-26})$ (G)	Hoffmann T. J. et al., 2016
	PP	$\beta = -0.22 (p = 1.0 \times 10^{-15})$ (G)	
	SBP	$\beta = 0.52 (p = 8.43 \times 10^{-11})$ (C)	
	DBP	$\beta = 0.27 (p = 2.33 \times 10^{-8})$ (C)	
	SBP x alcohol consumption interaction	$\beta = 0.54 (p = 2.24 \times 10^{-29})$ (C)	Feitosa M. F. et al., 2018
	MAP x alcohol consumption interaction	$\beta = 0.22 (p = 2 \times 10^{-12})$ (C)	
	MAP	$\beta = 0.39 (p = 2.56 \times 10^{-13})$ (C)	
	SBP	$\beta = 0.51 (p = 7.16 \times 10^{-12})$ (C)	
	DBP	$\beta = 0.33 (p = 8.93 \times 10^{-12})$ (C)	Takeuchi F., et al., 2018
	Hypertension	$\beta = 0.08 (p = 5.13 \times 10^{-10})$ (C)	
	Coronary artery disease	$(p = 8.81 \times 10^{-9})$ (C)	
	High blood pressure and chronic obstructive pulmonary disease	$(p = 1.18 \times 10^{-47})$	
	SBP (smoking interaction) (European)	$p = 8.44 \times 10^{-30}$	Sung Y. J., et al., 2018
	DBP (smoking interaction) (European)	$p = 2.68 \times 10^{-30}$	
	SBP (smoking interaction) (trans-ethnic data)	$p = 1.80 \times 10^{-40}$	
	DBP (smoking interaction) (trans-ethnic data)	$p = 1.18 \times 10^{-40}$	
	MAP	$\beta = 0.03 (p = 3.05 \times 10^{-50})$ (C)	Sakaue S. et al., 2021
rs7302981 <i>CERS5/AC074032.1</i> (12:50144032)	Hypertension	$Z \text{ score} = 6.23 (p = 4.8 \times 10^{-10})$ (A)	Liu C. et al., 2016
	SBP	$\beta = 0.37 (p = 9.4 \times 10^{-15})$ (A)	
	DBP	$\beta = 0.25 (p = 9.4 \times 10^{-19})$ (A)	
	DBP (European)	$\beta = 0.25 (p = 1.38 \times 10^{-17})$ (A)	Surendran P. et al., 2016
	SBP (European)	$\beta = 0.34 (p = 6.06 \times 10^{-13})$ (A)	
	Hypertension (European)	$Z \text{ score} = 6.07 (p = 1.28 \times 10^{-9})$ (A)	
	DBP (trans-ethnic data)	$\beta = 0.25 (p = 2.60 \times 10^{-19})$ (A)	
	SBP (trans-ethnic data)	$\beta = 0.35 (p = 9.94 \times 10^{-19})$ (A)	
	Hypertension (trans-ethnic data)	$Z \text{ score} = 6.17 (p = 6.82 \times 10^{-10})$ (A)	
rs2681472 <i>ATP2B1</i>	DBP	$\beta = 0.50 (p = 1.47 \times 10^{-9})$ (A)	Levy D. et al., 2009
	Hypertension	$\beta = 0.15 (p = 1.75 \times 10^{-11})$ (A)	

(12: 89615182)	Coronary artery disease	OR = 1.08 ($p = 6.17 \times 10^{-11}$) (G)	Nikpay M., et al., 2015
	Myocardial infarction	OR = 1.08 ($p = 6.03 \times 10^{-9}$) (G)	
	MAP	$\beta = -0.59$ ($p = 1.1 \times 10^{-21}$) (G)	
	SBP	$\beta = -0.85$ ($p = 1.3 \times 10^{-21}$) (G)	
	DBP	$\beta = -0.47$ ($p = 3.7 \times 10^{-17}$) (G)	Liu C. et al., 2016
	Hypertension	$\beta = -0.033$ ($p = 3.5 \times 10^{-8}$) (G)	
	Coronary artery disease	OR = 1.07 ($p = 1 \times 10^{-21}$) (G)	
	SBP	$\beta = 0.72$ ($p = 1.06 \times 10^{-20}$) (A)	
	DBP	$\beta = 0.33$ ($p = 3.77 \times 10^{-11}$) (A)	
	MAP	$\beta = 0.46$ ($p = 5.05 \times 10^{-17}$) (A)	
	PP	$\beta = 0.40$ ($p = 5.49 \times 10^{-14}$) (A)	
	Hypertension	$\beta = 0.07$ ($p = 1.41 \times 10^{-6}$) (A)	
	Coronary artery disease (trans-ethnic data)	$\beta = 0.060$ ($p = 6.8 \times 10^{-25}$) (G)	Koyama S., et al., 2020
	Coronary artery disease (Japanese)	$\beta = 0.068$ ($p = 2.6 \times 10^{-11}$) (G)	
	Myocardial infarction	OR = 1.07 ($p = 1.3 \times 10^{-12}$) (G)	
	Myocardial infarction	$\beta = 0.071$ ($p = 1.17 \times 10^{-11}$) (G)	
rs8068318 <i>TBX2/</i> <i>TBX2-AS1</i> (17:61406405)	MAP	$\beta = -0.28$ ($p = 2.0 \times 10^{-8}$) (C)	Liu C. et al., 2016
	SBP	$\beta = -0.42$ ($p = 3.9 \times 10^{-17}$) (C)	
	Hypertension	Z score = -6.96 ($p = 3.0 \times 10^{-12}$) (C)	
	DBP	$\beta = -0.26$ ($p = 3.0 \times 10^{-18}$) (C)	
	SBP (European)	$\beta = 0.42$ ($p = 1.3 \times 10^{-15}$) (T)	Surendran P. et al., 2016
	DBP (European)	$\beta = 0.26$ ($p = 1.95 \times 10^{-16}$) (T)	
	Hypertension (European)	Z score = 6.97 ($p = 3.21 \times 10^{-12}$) (T)	
	DBP (trans-ethnic data)	$\beta = 0.25$ ($p = 2.75 \times 10^{-18}$) (T)	
	SBP (trans-ethnic data)	$\beta = 0.41$ ($p = 2.3 \times 10^{-17}$) (T)	
	Hypertension (trans-ethnic data)	Z score = 6.96 ($p = 3.43 \times 10^{-12}$) (T)	
rs167479 <i>RGL3</i> (19:11416089)	SBP	$\beta = 0.41$ ($p = 1.6 \times 10^{-21}$) (G)	Hoffmann T. J. et al., 2017
	DBP	$\beta = 0.25$ ($p = 4.3 \times 10^{-22}$) (G)	
	PP	$\beta = 0.18$ ($p = 3.2 \times 10^{-8}$) (G)	
	MAP	$\beta = -0.30$ ($p = 7.3 \times 10^{-11}$) (T)	Liu C. et al., 2016
	Hypertension	Z score = -7.72 ($p = 1.2 \times 10^{-14}$) (T)	
	SBP	$\beta = -0.45$ ($p = 1.0 \times 10^{-23}$) (T)	
	DBP	$\beta = -0.30$ ($p = 4.2 \times 10^{-28}$) (T)	
	DBP (European)	$\beta = -0.33$ ($p = 1.99 \times 10^{-31}$) (T)	Surendran P. et al., 2016
	SBP (European)	$\beta = -0.50$ ($p = 1.49 \times 10^{-26}$) (T)	

	Hypertension (European)	Z score = -7.86 ($p = 4.01 \times 10^{-15}$) (T)	
	DBP (trans-ethnic data)	$\beta = -0.31$ ($p = 2.76 \times 10^{-32}$) (T)	
	SBP (trans-ethnic data)	$\beta = -0.47$ ($p = 8.64 \times 10^{-27}$) (T)	
	Hypertension (trans-ethnic data)	Z score = -7.88 ($p = 3.37 \times 10^{-15}$) (T)	
	SBP	$\beta = -0.41$ ($p = 4.32 \times 10^{-36}$) (T)	Giri A. et el. 2018
	Hypertension	$\beta = 0.05$ ($p = 8.0 \times 10^{-16}$) (G)	German C. A., 2019
	Hypertension	OR = $0.84 - 0.92$ ($p = 2.37 \times 10^{-8}$) (T)	Jeong H. et al., 2020
	SBP	$\beta = -0.027$ ($p = 3.24 \times 10^{-46}$) (T)	Sakaue S. et al., 2021
	DBP	$\beta = -0.026$ ($p = 2.24 \times 10^{-48}$) (T)	
	PP	$\beta = -0.016$ ($p = 7.72 \times 10^{-18}$) (T)	
	MAP	$\beta = -0.028$ ($p = 3.64 \times 10^{-48}$) (T)	

Notes: SBP - systolic blood pressure; DBP - diastolic blood pressure; MAP - mean arterial pressure; PP - pulse blood pressure; z-score - standard scores, β – effect, OR – odds ratio, p – significance level.

References

1. Ehret GB, Ferreira T, Chasman DI, et al. The genetics of blood pressure regulation and its target organs from association studies in 342,415 individuals. *Nat Genet.* 2016;48(10):1171-1184. doi:10.1038/ng.3667
2. Feitosa MF, Kraja AT, Chasman DI, et al. Novel genetic associations for blood pressure identified via gene-alcohol interaction in up to 570K individuals across multiple ancestries. *PLoS One.* 2018;13(6):e0198166. doi:10.1371/journal.pone.0198166
3. German CA, Sinsheimer JS, Klimentidis YC, Zhou H, Zhou JJ. Ordered multinomial regression for genetic association analysis of ordinal phenotypes at Biobank scale. *Genet Epidemiol.* 2020;44(3):248-260. doi:10.1002/gepi.22276
4. Giri A, Hellwege JN, Keaton JM, et al. Trans-ethnic association study of blood pressure determinants in over 750,000 individuals. *Nat Genet.* 2019;51(1):51-62. doi:10.1038/s41588-018-0303-9
5. Hartiala JA, Han Y, Jia Q, et al. Genome-wide analysis identifies novel susceptibility loci for myocardial infarction. *Eur Heart J.* 2021;42(9):919-933. doi:10.1093/eurheartj/ehaa1040
6. Hoffmann TJ, Ehret GB, Nandakumar P, et al. Genome-wide association analyses using electronic health records identify new loci influencing blood pressure variation. *Nat Genet.* 2017;49(1):54-64. doi:10.1038/ng.3715
7. International Consortium for Blood Pressure Genome-Wide Association Studies, Ehret GB, Munroe PB, et al. Genetic variants in novel pathways influence blood pressure and cardiovascular disease risk. *Nature.* 2011;478(7367):103-109. doi:10.1038/nature10405
8. Jeong H, Jin HS, Kim SS, Shin D. Identifying Interactions between Dietary Sodium, Potassium, Sodium-Potassium Ratios, and FGF5 rs16998073 Variants and Their Associated Risk for Hypertension in Korean Adults. *Nutrients.* 2020;12(7):2121. doi:10.3390/nu12072121
9. Kato N, Loh M, Takeuchi F, et al. Trans-ancestry genome-wide association study identifies 12 genetic loci influencing blood pressure and implicates a role for DNA methylation. *Nat Genet.* 2015;47(11):1282-1293. doi:10.1038/ng.3405
10. Koyama S, Ito K, Terao C, et al. Population-specific and trans-ancestry genome-wide analyses identify distinct and shared genetic risk loci for coronary artery disease. *Nat Genet.* 2020;52(11):1169-1177. doi:10.1038/s41588-020-0705-3
11. Levy D, Ehret GB, Rice K, et al. Genome-wide association study of blood pressure and hypertension. *Nat Genet.* 2009;41(6):677-687. doi:10.1038/ng.384

12. Liu C, Kraja AT, Smith JA, et al. Meta-analysis identifies common and rare variants influencing blood pressure and overlapping with metabolic trait loci. *Nat Genet.* 2016;48(10):1162-1170. doi:10.1038/ng.3660
13. Nelson CP, Goel A, Butterworth AS, et al. Association analyses based on false discovery rate implicate new loci for coronary artery disease. *Nat Genet.* 2017;49(9):1385-1391. doi:10.1038/ng.3913
14. Nikpay M, Goel A, Won HH, et al. A comprehensive 1,000 Genomes-based genome-wide association meta-analysis of coronary artery disease. *Nat Genet.* 2015;47(10):1121-1130. doi:10.1038/ng.3396
15. Sakaue S, Kanai M, Tanigawa Y, et al. A cross-population atlas of genetic associations for 220 human phenotypes. *Nat Genet.* 2021;53(10):1415-1424. doi:10.1038/s41588-021-00931-x
16. Shungin D, Winkler TW, Croteau-Chonka DC, et al. New genetic loci link adipose and insulin biology to body fat distribution. *Nature.* 2015;518(7538):187-196. doi:10.1038/nature14132
17. Sung YJ, Winkler TW, de Las Fuentes L, et al. A Large-Scale Multi-ancestry Genome-wide Study Accounting for Smoking Behavior Identifies Multiple Significant Loci for Blood Pressure. *Am J Hum Genet.* 2018;102(3):375-400. doi:10.1016/j.ajhg.2018.01.015
18. Surendran P, Drenos F, Young R, et al. Trans-ancestry meta-analyses identify rare and common variants associated with blood pressure and hypertension. *Nat Genet.* 2016;48(10):1151-1161. doi:10.1038/ng.3654
19. Tachmazidou I, Süveges D, Min JL, et al. Whole-Genome Sequencing Coupled to Imputation Discovers Genetic Signals for Anthropometric Traits. *Am J Hum Genet.* 2017;100(6):865-884. doi:10.1016/j.ajhg.2017.04.014
20. Takeuchi F, Akiyama M, Matoba N, et al. Interethnic analyses of blood pressure loci in populations of East Asian and European descent. *Nat Commun.* 2018;9(1):5052. doi:10.1038/s41467-018-07345-0
21. Wain LV, Vaez A, Jansen R, et al. Novel Blood Pressure Locus and Gene Discovery Using Genome-Wide Association Study and Expression Data Sets From Blood and the Kidney. *Hypertension.* 2017; HYPERTENSIONAHA.117.09438. doi:10.1161/HYPERTENSIONAHA.117.09438
22. Wain LV, Verwoert GC, O'Reilly PF, et al. Genome-wide association study identifies six new loci influencing pulse pressure and mean arterial pressure. *Nat Genet.* 2011;43(10):1005-1011. doi:10.1038/ng.922
23. Zhou W, Nielsen JB, Fritzsche LG, et al. Efficiently controlling for case-control imbalance and sample relatedness in large-scale genetic association studies. *Nat Genet.* 2018;50(9):1335-1341. doi:10.1038/s41588-018-0184-y

24. Zhu Z, Wang X, Li X, et al. Genetic overlap of chronic obstructive pulmonary disease and cardiovascular disease-related traits: a large-scale genome-wide cross-trait analysis. *Respir Res*. 2019;20(1):64. doi:10.1186/s12931-019-1036-8