

**Supplemental Table 3: Genomic regions selected for analysis**

Genomic region	GWAS Study	Chromosome	Start Position <sup>a</sup>	Stop Position <sup>a</sup>
<i>ARHGAP42</i>	ICBP, 2011 [1]	11	100553384	100867668
<i>ATP2B1</i>	ICBP, 2011 [1]	12	89976828	90108077
<i>BAG6</i>	ICBP, 2011 [1]	6	31606805	31620953
<i>C10orf107</i>	ICBP, 2011 [1]	10	63417719	63531524
<i>CACNB2</i>	ICBP, 2011 [1]	10	18424606	18835798
<i>CSK</i>	ICBP, 2011 [1]	15	75074425	75095539
<i>FES</i>	ICBP, 2011 [1]	15	91421925	91444006
<i>GOSR2</i>	ICBP, 2011 [1]	17	44995483	45110003
<i>GUCY1A3</i>	ICBP, 2011 [1]	4	156582863	156658501
<i>HFE</i>	ICBP, 2011 [1]	6	26082509	26103571
<i>MECOM</i>	ICBP, 2011 [1]	3	168796287	169386406
<i>MTHFR</i>	ICBP, 2011 [1]	1	11840780	11871977
<i>NT5C2</i>	ICBP, 2011 [1]	10	104840940	104958056
<i>PLCE1</i>	ICBP, 2011 [1]	10	95748746	96097580
<i>PLEKHA7</i>	Franceschini, 2013 [2]	11	16794842	17040990
<i>PLEKHG1</i>	ICBP, 2011 [1]	6	150915999	151169799
rs10850411	ICBP, 2011 [1]	12	115337796	115437796
rs1173771	ICBP, 2011 [1]	5	32765028	32865028
rs11953630	ICBP, 2011 [1]	5	157795402	157895402
rs13082711	ICBP, 2011 [1]	3	27487909	27587909
rs13209747	Franceschini, 2013 [2]	6	127065454	127165454
rs1327235	ICBP, 2011 [1]	20	10919030	11019030
rs1458038	ICBP, 2011 [1]	4	81114723	81214723
rs17428471	Franceschini, 2013 [2]	7	27287867	27387867
rs2932538	ICBP, 2011 [1]	1	113196543	113296543
rs4373814	ICBP, 2011 [1]	10	18369972	18469972
rs7129220	ICBP, 2011 [1]	11	10300538	10400538
<i>SH2B3</i>	ICBP, 2011 [1]	12	111838752	111894427
<i>SLC39A8</i>	ICBP, 2011 [1]	4	103167198	103357415
<i>SOX6</i>	Franceschini, 2013 [2]	11	15982995	16766138
<i>ULK4</i>	ICBP, 2011 [1]; Franceschini, 2013 [2]	3	41283090	42008922
<i>ZNF652</i>	ICBP, 2011 [1]	17	47361568	47444835
<i>ZNF831</i>	ICBP, 2011 [1]	20	57735895	57834168

<sup>a</sup> Start and stop positions for the genomic region include buffers (+/-5kb for gene regions; +/-50kb for non-gene regions).

## References

1. International Consortium for Blood Pressure Genome-Wide Association Studies; Ehret, G.B.; Munroe, P.B.; Rice, K.M.; Bochud, M.; Johnson, A.D.; Chasman, D.I.; Smith, A.V.; Tobin, M.D.; Verwoert, G.C. *et al.* Genetic Variants in Novel Pathways Influence Blood Pressure and Cardiovascular Disease Risk. *Nature* **2011**, *478*, 103-109.
2. Franceschini, N.; Fox, E.; Zhang, Z.; Edwards, T.L.; Nalls, M.A.; Sung, Y.J.; Tayo, B.O.; Sun, Y.V.; Gottesman, O.; Adeyemo, A. *et al.* Genome-Wide Association Analysis of Blood-Pressure Traits in African-Ancestry Individuals Reveals Common Associated Genes in African and Non-African Populations. *Am. J. Hum. Genet.* **2013**, *93*, 545-554.