# Trans-ethnic meta-analysis of white blood cell phenotypes 

 Andrew D. Johnson ${ }^{20,22}$, Ming-Huei Chen ${ }^{21,22}$, Albert V. Smith ${ }^{23,24}$, Andrew P. Morris ${ }^{25,26}$, Toshiko Tanaka ${ }^{3}$, Luigi Ferrucci${ }^{3}$, Alan B. Zonderman ${ }^{4}$, Guillaume Lettre ${ }^{27,28}$, Tamara Harris ${ }^{2}$, Melissa Garcia ${ }^{2}$, Stefania Bandinelli ${ }^{29}$, Rehan Qayyum ${ }^{30}$, Lisa R. Yanek ${ }^{30}$, Diane M. Becker ${ }^{30}$, Lewis C. Becker ${ }^{30,31}$, Charles Kooperberg ${ }^{12}$, Brendan Keating ${ }^{32,33}$, Jared Reis ${ }^{34}$, Hua Tang ${ }^{35}$, Eric Boerwinkle ${ }^{36}$, Yoichiro Kamatani ${ }^{13}$, Koichi Matsuda ${ }^{37}$, Naoyuki Kamatani ${ }^{13}$, Yusuke  Albert Hofman ${ }^{16,19}$, André G. Uitterlinden ${ }^{16,18,19}$, Cornelia M. van Duijn ${ }^{16,19}$, Oscar H. Franco ${ }^{16,17,19,}$ Dan L. Longo ${ }^{5}$, Andrew B. Singleton ${ }^{1}$, Bruce M. Psaty ${ }^{9,10,11,42, ~ M i c h e l l e ~ K . ~ E v a n s ~}{ }^{6}$, L. Adrienne Cupples ${ }^{22,43}$, Jerome I. Rotter ${ }^{44,45}$, Christopher J. O’Donnell ${ }^{20,22}$, Atsushi Takahashi ${ }^{13, \ddagger}$, James G. Wilson ${ }^{46, \ddagger}$, Santhi K. Ganesh ${ }^{47,48, \ddagger, *}$ and Mike A. Nalls ${ }^{1, \downarrow, *}$ for the CHARGE Hematology, COGENT, and BioBank Japan Project (RIKEN) Working Groups ${ }^{\bullet}$<br>${ }^{1}$ Laboratory of Neurogenetics, ${ }^{2}$ Laboratory of Epidemiology and Population Sciences, National Institute on Aging, National Institutes of Health, Bethesda, MD, USA, ${ }^{3}$ Longitudinal Studies Section, Clinical Research Branch, ${ }^{4}$ Behavioral Epidemiology Section, Laboratory of Epidemiology \& Population Sciences, National Institute on Aging Intramural Research Program, National Institutes of Health, Baltimore, MD, USA, ${ }^{5}$ Laboratory of Genetics, ${ }^{6}$ Health Disparities Research Section, Clinical Research Branch, National Institute on Aging, National Institutes of Health, Baltimore, MD, USA, ${ }^{7}$ Department of Biological Anthropology, Temple University, Philadelphia, PA, USA, ${ }^{8}$ Department of Epidemiology, ${ }^{9}$ Cardiovascular Health Research Unit, ${ }^{10}$ Department of Medicine, ${ }^{11}$ Department of Epidemiology and Health Services, University of Washington, Seattle, WA, USA, ${ }^{12}$ Division of Public Health Sciences, Fred Hutchinson Cancer Research Center, Seattle, WA, USA, ${ }^{13}$ Laboratory for Statistical Analysis, ${ }^{14}$ Laboratory for Genotyping Development, RIKEN Center for Integrative Medical Sciences, Yokohama, Japan, ${ }^{15}$ Department of Human Genetics and Disease Diversity, Graduate School of Medical and Dental Sciences, Tokyo Medical and Dental University, Tokyo, Japan, ${ }^{16}$ Department of Epidemiology, ${ }^{17}$ ErasmusAGE, Department of Epidemiology, ${ }^{18}$ Department of Internal Medicine, Erasmus Medical Center, Rotterdam, The Netherlands, ${ }^{19}$ Consortium for Healthy Aging (NGI-NCHA), The Netherlands Genomics Initiative, Leiden, The Netherlands, ${ }^{20}$ Cardiovascular Epidemiology and Human Genomics Branch, NHLBI Division of Intramural Research, Bethesda, MD, USA, ${ }^{21}$ Department of Neurology, Boston University School of Medicine, Boston, MA, USA, ${ }^{22}$ NHLBI Framingham Heart Study, Bethesda, MD, USA, ${ }^{23}$ Icelandic Heart Association, Kopavogur, Iceland, ${ }^{24}$ University of Iceland, Reykjavik, Iceland, ${ }^{25}$ Genetic and Genomic Epidemiology Unit, Wellcome Trust Centre for Human Genetics, University of Oxford, Oxford, UK, ${ }^{26}$ Department of Biostatistics, University of Liverpool, Liverpool, UK, ${ }^{27}$ Montreal Heart Institute, Montréal, Canada, ${ }^{28}$ Département de Médecine, Université de Montréal, Montréal, Canada, ${ }^{29}$ Geriatric Rehabilitation Unit, Azienda Sanitaria Firenze (ASF), Florence, Italy, ${ }^{30}$ GeneSTAR Research Program, Division of General Internal Medicine, ${ }^{31}$ Division of Cardiology, Johns Hopkins School of Medicine, Baltimore, MD, USA, ${ }^{32}$ Center for Applied Genomics, Children's Hospital of Philadelphia, PA, USA, ${ }^{33}$ Department of Pediatrics, University of Pennsylvania,

[^0]PA, USA, ${ }^{34}$ Division of Cardiovascular Sciences, National Heart, Lung, and Blood Institute, Bethesda, MD, USA, ${ }^{35}$ Stanford University School of Medicine, Stanford, CA 94305, USA, ${ }^{36}$ The Brown Foundation, Institute of Molecular Medicine for the Prevention of Human Diseases, University of Texas, Houston, TX, USA, ${ }^{37}$ Laboratory of Molecular Medicine, Human Genome Center, Institute of Medical Science, The University of Tokyo, Tokyo, Japan, ${ }^{38}$ Department of Medicine, ${ }^{39}$ Department of Surgery, Center for Personalized Therapeutics, The University of Chicago, Chicago, IL, USA, ${ }^{40}$ Department of Epidemiology, ${ }^{41}$ Department of Medicine, Brown University, Providence, RI, USA, ${ }^{42}$ Group Health Research Institute, Group Health Cooperative, Seattle, WA, USA, ${ }^{43}$ Boston University Department of Statistics, Boston, MA, USA, ${ }^{44}$ Institute for Translational Genomics and Population Sciences, Los Angeles BioMedical Research Institute at Harbor-UCLA Medical Center, Torrance, CA, USA, ${ }^{45}$ Division of Genetic Outcomes, Department of Pediatrics, HarborUCLA Medical Center, Torrance, CA, USA, ${ }^{46}$ Department of Physiology and Biophysics, University of Mississippi Medical Center, Jackson, MS, USA ${ }^{47}$ Division of Cardiovascular Medicine, Department of Internal Medicine, University of Michigan, Ann Arbor, MI, USA and ${ }^{48}$ Department of Human Genetics, University of Michigan, Ann Arbor, MI, USA

Received April 25, 2014; Revised and Accepted July 31, 2014


#### Abstract

White blood cell (WBC) count is a common clinical measure used as a predictor of certain aspects of human health, including immunity and infection status. WBC count is also a complex trait that varies among individuals and ancestry groups. Differences in linkage disequilibrium structure and heterogeneity in allelic effects are expected to play a role in the associations observed between populations. Prior genome-wide association study (GWAS) meta-analyses have identified genomic loci associated with WBC and its subtypes, but much of the heritability of these phenotypes remains unexplained. Using GWAS summary statistics for over 50000 individuals from three diverse populations (Japanese, African-American and European ancestry), a Bayesian model methodology was employed to account for heterogeneity between ancestry groups. This approach was used to perform a trans-ethnic meta-analysis of total WBC, neutrophil and monocyte counts. Ten previously known associations were replicated and six new loci were identified, including several regions harboring genes related to inflammation and immune cell function. Ninety-five percent credible interval regions were calculated to narrow the association signals and fine-map the putatively causal variants within loci. Finally, a conditional analysis was performed on the most significant SNPs identified by the trans-ethnic meta-analysis (MA), and nine secondary signals within loci previously associated with WBC or its subtypes were identified. This work illustrates the potential of trans-ethnic analysis and ascribes a critical role to multi-ethnic cohorts and consortia in exploring complex phenotypes with respect to variants that lie outside the European-biased GWAS pool.


## INTRODUCTION

White blood cells (WBCs) are critically involved in the body's immune system, serving as a primary defense mechanism against foreign pathogens. WBC count is used as a clinical marker of inflammation status, and higher WBC count has been associated with a risk of cardiovascular disease, cancer mortality and all-cause mortality $(1-3)$. Elevated WBC count is also associated with disease risk factors including increasing age, high blood pressure, cigarette smoking, adiposity and increasing plasma inflammatory markers (4).

WBCs are classified into five subtypes according to their morphology and functions, including neutrophils, basophils, eosinophils, lymphocytes and monocytes. Total WBC count is highly variable even among healthy individuals of the same population (5). WBC count is a moderately heritable phenotype, with $h^{2}$ estimates ranging from 0.14 to 0.40 across the WBC subtypes (6). Additionally, between 25 and $50 \%$ of individuals of African descent exhibit benign ethnic neutropenia, characterized by low neutrophil counts, due to a regulatory variant in the Duffy antigen receptor for chemokines ( $D A R C$ ) gene
$(5,7,8)$. Given the importance of WBC in both host defense and, potentially, pathologic inflammation, elucidation of additional genetic mechanisms responsible for regulating while blood cell count could have a substantial medical impact.

Admixture mapping and genome-wide association studies (GWAS) performed on cohorts of differing continental ancestry, including European, Japanese and African-American, have been successful in identifying multiple loci associated with WBC phenotypes $(5,8-14)$. The joint effects of these loci generally explain only a small portion of the overall heritability of either total WBC or WBC subtypes. Some prior GWAS have not defined the subtypes of WBC that are driving their observed associations; however, neutrophils in particular are often implicated $(9,11)$. Furthermore, the loci identified by GWAS generally encompass large genomic regions, often containing many genes and variants with comparable association signals. Thus, fine-mapping methods aimed at pinpointing association signals more precisely are needed (15).

Recently, the 1000 Genomes Project and phase three of the HapMap project released comprehensive reference panels for a number of ethnic groups, including African, Asian and
additional European populations $(16,17)$. Imputation using these higher density reference panels allows inference of genotypes not captured by genotyping arrays, markedly increasing the breadth of genetic variation that can be included in association tests. This has provided new opportunities both to detect novel loci and to refine the localization of association signals for a number of phenotypes, including WBC phenotypes.

Trans-ethnic meta-analysis (MA) potentially offers a more comprehensive view of the genetic variation that is associated with a trait, but traditional fixed-effects MA methods do not adequately address heterogeneity in allelic effects, allele frequencies or differences in linkage disequilibrium between ethnicities (18). For example, in a previous fixed-effects analysis of the cohorts included in the current study, only 152 of 161 single nucleotide polymorphisms (SNPs) that had been associated with WBC phenotypes in earlier analyses were replicated at a Bonferroni-corrected significance threshold of $P<3.57 \mathrm{E}-3$, and no novel associations were observed (9). Whereas random-effects-based methods of MA do account for inter-study heterogeneity, they lose statistical power in the setting of high levels of heterogeneity that may result from experimental or statistical differences in study design $(15,19)$.

These shortcomings have been addressed in the software package, MANTRA (Meta-Analysis of Trans-ethnic Association studies), which allows for heterogeneity between diverse ethnic groups and provides increased power and mapping resolution compared with random-effects-based methods (15). In the current study, we used MANTRA to combine summary results of ancestryspecific GWAS of WBC traits in three distinct populations. We identify novel loci associated with WBC count, assess heterogeneity in allelic effects between ancestry groups and improve finemapping resolution of some previously identified regions.

## RESULTS

Descriptive statistics for each cohort are found in Table 1. In the trans-ethnic MANTRA analyses, we observed strong evidence of association, defined by a $\log _{10}$ Bayes factor (BF) of $>6$, at 10 previously identified loci and six novel loci, and detected nine secondary signals within 500 kb of a previously identified locus. The population-specific and trans-ethnic results for the
established and novel loci associated with each WBC phenotype (total WBC, neutrophil and monocyte count) are summarized in Table 2. Cohort-level Manhattan plots are shown in Figure 1. Of the 15 previously identified variants (at 10 loci), six of the six monocyte associations, two of the four neutrophil associations and two of the five WBC count associations were initially identified in the original GWAS papers from which the data employed by this analysis are drawn.

Regions previously identified by single-ethnicity GWAS reappeared in the MANTRA trans-ethnic analyses, but in some instances, the index SNP from the original publication was not the most significant. These include rs4065321 and rs17609240 on 17q21.1 (WBC count and neutrophil count), rs2517524 on 6p21.33 (WBC count) and rs10956483 on 8q24.21 (monocyte count). Additionally, rs2814778, a marker identifying the Duffy null blood group antigen and located on the DARC gene of chromosome 1 at position (b37) 159174 683, was available only in the COGENT data. This marker accounts for $20 \%$ of population variance in the WBC of African ancestry populations and is monomorphic in non-African populations.

## Novel associations

In addition to replicating known variants, the trans-ethnic analysis identified six novel trait-locus associations (Table 2). For neutrophil counts, novel findings include rs6936204 in region 6 p 21.32 , located nearest to $A K 123889$. This region is very near known locus 6 p 21.33 , which was previously associated with WBC and lymphocyte counts (9). These loci are near the $H L A$ region; thus, it is possible that population stratification is driving this association (20). These variants are not in linkage disequilibrium (LD) with any of the known HLA markers, but as meiotic crossovers are known to cluster around $H L A$, it is possible that these variants are separated from this region by a recombination hotspot. Novel association for WBC count includes rs 10932765 in region $2 q 35$, located near ARPC2, which has previously been associated with monocyte count and inflammatory bowel disease $(12,21)$. This region is notable because of its proximity to $I L 8 R A$, which encodes CXCR1. CXCR1, the receptor for the chemokine $I L-8$, is a mediator of inflammatory responses; interestingly, the Duffy antigen

Table 1. Descriptive statistics

| Study | RIKEN | CHARGE | COGENT |
| :---: | :---: | :---: | :---: |
| Population ancestry | Japanese | European | African-American |
| Total WBC |  |  |  |
| WBC count: mean (SD) in cells $\times 10^{3} / \mathrm{ml}$ | 6.20 (1.67), $n=16843$ | 5.72 (1.24), $n=19509$ | 5.94 (1.88), $n \sim 16388$ |
| Neutrophils: mean (SD) in cells $\times 10^{3} / \mathrm{ml}$ | 3.80 (1.40), $n=9802$ | 3.52 (1.06), $n=16550$ | 3.57 (1.57), $n \sim 7391$ |
| Monocytes: mean (SD) in cells $\times 10^{3} / \mathrm{ml}$ | 0.36 (0.22), $n=9810$ | 0.43 (0.14), $n=16550$ | 0.36 (0.17), $n \sim 7369$ |
| Covariates |  |  |  |
| Age in years: mean (SD) | 63.5 (10.6) | 63.8 (8.9) | 50.9 (7.6) |
| BMI: mean (SD) $\mathrm{kg} / \mathrm{m}^{2}$ | 23.1 (3.6) | NA | NA |
| \% Female | 39.1 | 53.5 | 65.6 |
| \% Current smoker | 52.6 | 17.6 | NA |
| Sample size |  |  |  |
| Total $N$ | 17218 | 19509 | 16388 |

Counts are reported in thousands of cells per milliliter of blood (cells $\left.\times 10^{9} / l\right)$.
SD, standard deviation; NA, not available.

Table 2. Loci defined in MANTRA analysis

| Lead SNP | Chr | base pair <br> (BP) (bd 37) | Effect | Other | MANTRA trans-ethnic meta-analysis |  |  |  | European ancestry GWAS meta-analysis |  |  |  |  |  | Japanese ancestry GWAS meta-analysis |  |  |  |  |  | African-American ancestry GWAS meta-analysis |  |  |  |  |  |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  |  |  |  |  | $\log _{10} \mathrm{BF}$ (bayes factor) assocatio | PPA | Sample size | Gene(s) | EAF | Beta | SE | $P$-value | Sample size | $R^{2 a}$ | EAF | Beta | SE | $P$-value | Sample size | $R^{2}$ | EAF | Beta | SE | $P$-value | Sample size | $R^{2 a}$ |
| WBC count-established loci |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| rs2518564* | 1 | 159062436 | A | G | 341.59 | 1 | 51768 | DARC | 0.293 | 0.007 | 0.003 | $2.81 \mathrm{E}-02$ | 19509 | 0.883 | 0.7159 | 0 | 0.003 | 0.9936 | 16843 | 0.991 | 0.1725 | 0.1905 | 0.0048 | $1.39 \mathrm{E}-149$ | 15416 | 1 |
| rs1371799 | 4 | 74977837 | C | T | 13.5 | 1 | 52740 | CXCL2 | 0.572 | -0.012 | 0.003 | $2.14 \mathrm{E}-05$ | 19509 | 0.985 | 0.461 | -0.012 | 0.003 | $3.63 \mathrm{E}-05$ | 16843 | 0.921 | 0.761 | 0.023 | 0.0041 | $5.21 \mathrm{E}-04$ | 16388 | 1 |
| rs4895441 | 6 | 135426573 | A | G | 9.64 | 0.165 | 52703 | Intergenic | 0.347 | 0.011 | 0.003 | $3.42 \mathrm{E}-04$ | 19509 | 1 | 0.628 | 0.018 | 0.003 | $8.99 \mathrm{E}-11$ | 16843 | 1 | 0.902 | 0.0002 | 0.0059 | $9.90 \mathrm{E}-01$ | 16351 | 1 |
| rs445 | 7 | 92408370 | C | T | 16.8 | 1 | 52740 | CDK6 | 0.776 | 0.019 | 0.005 | $4.51 \mathrm{E}-04$ | 19509 | 1 | 0.685 | 0.019 | 0.003 | $8.99 \mathrm{E}-11$ | 16843 | 1 | 0.193 | -0.0249 | 0.0049 | $9.90 \mathrm{E}-01$ | 16388 | 1 |
| rs4794822 | 17 | 38156712 | C | T | 29.64 | 1 | 52740 | PSMD3 | 0.586 | -0.028 | 0.003 | $3.23 \mathrm{E}-23$ | 19509 | 0.999 | 0.481 | -0.019 | 0.003 | $2.90 \mathrm{E}-12$ | 16843 | 0.999 | 0.325 | 0.0128 | 0.0038 | $4.33 \mathrm{E}-02$ | 16388 | 0.918 |
| WBC count-novel loci |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| rs10932765 | 2 | 219099484 | C | T | 6.95 | 0.025 | 52740 | ARPC2 | 0.503 | 0.011 | 0.003 | $3.36 \mathrm{E}-05$ | 19509 | 1 | 0.58 | 0.012 | 0.003 | $1.69 \mathrm{E}-05$ | 16843 | 0.986 | 0.877 | 0.0039 | 0.0056 | $7.26 \mathrm{E}-01$ | 16388 | 1 |
| rs6734238 | 2 | 113841030 | A | G | 6.62 | 0.028 | 52691 | IL1F10 | 0.451 | -0.009 | 0.003 | $1.46 \mathrm{E}-03$ | 19509 | 1 | 0.967 | -0.018 | 0.008 | $2.49 \mathrm{E}-02$ | 16843 | 0.907 | 0.553 | -0.0158 | 0.0035 | $5.92 \mathrm{E}-03$ | 16339 | 1 |
| rs2853946 | 6 | 31247203 | A | T | 12.16 | 0.009 | 52740 | HLA-B | 0.348 | -0.017 | 0.003 | $1.28 \mathrm{E}-08$ | 19509 | 1 | 0.717 | -0.013 | 0.003 | $8.19 \mathrm{E}-06$ | 16843 | 0.976 | 0.512 | -0.0106 | 0.0035 | $7.65 \mathrm{E}-02$ | 16388 | 1 |
| rs2163950 | 8 | 130597585 | A | C | 7.75 | 0.065 | 52740 | Intergenic | 0.054 | -0.017 | 0.006 | $5.91 \mathrm{E}-03$ | 19509 | 0.861 | 0.174 | -0.021 | 0.004 | $5.36 \mathrm{E}-09$ | 16843 | 0.995 | 0.124 | -0.0053 | 0.0058 | $6.44 \mathrm{E}-01$ | 16388 | 0.788 |
| Monocyte count-established loci |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| rs1449263 | 2 | 182319301 | C | T | 19.24 | , | 33729 | ITGA4 | 0.541 | -0.036 | 0.005 | 6.70E-14 | 16550 | 0.942 | 0.626 | -0.033 | 0.006 | $8.32 \mathrm{E}-08$ | 9,810 | 0.994 | 0.58 | 0.0087 | 0.0031 | $4.68 \mathrm{E}-03$ | 7369 | 0.965 |
| rs12988934 | 2 | 182323665 | C | T | 7.75 | 0.022 | 26360 | ITGA4 | 0.789 | -0.034 | 0.01 | $1.01 \mathrm{E}-03$ | 16550 | 0.511 | 0.732 | -0.041 | 0.008 | $1.42 \mathrm{E}-07$ | 9810 | 0.73 | 0.018 | -0.0165 | 0.0854 | $8.47 \mathrm{E}-01$ | 1803 | 0.418 |
| rs9880192 | 3 | 128297569 | C | G | 8.3 | , | 33745 | C3orf27 | 0.441 | -0.028 | 0.005 | $1.35 \mathrm{E}-08$ | 16550 | 0.867 | 0.091 | $-0.037$ | 0.012 | $1.21 \mathrm{E}-03$ | 9810 | 0.804 | 0.188 | 0.0037 | 0.0047 | $4.31 \mathrm{E}-01$ | 7385 | 0.674 |
| rs3095254 | 6 | 31221668 | C | G | 6.81 | 1 | 33745 | MHC | 0.377 | 0.008 | 0.005 | $1.19 \mathrm{E}-01$ | 16550 | 0.976 | 0.461 | 0.035 | 0.006 | $8.27 \mathrm{E}-09$ | 9810 | 0.896 | 0.562 | 0.0031 | 0.0031 | $3.31 \mathrm{E}-01$ | 7385 | 0.935 |
| rs1991866 | 8 | 130624105 | C | G | 15.3 | 1 | 33598 | Intergenic | 0.451 | $-0.032$ | 0.005 | $4.58 \mathrm{E}-11$ | 16550 | 0.96 | 0.598 | -0.034 | 0.006 | $7.38 \mathrm{E}-08$ | 9810 | 0.892 | 0.52 | -0.0046 | 0.0031 | $1.32 \mathrm{E}-01$ | 7238 | 1 |
| rs10980800 | 9 | 113915905 | C | T | 11.63 | 1 | 33692 | Intergenic | 0.769 | 0.044 | 0.006 | $1.10 \mathrm{E}-14$ | 16550 | 0.988 | 0.056 | 0.004 | 0.013 | $7.61 \mathrm{E}-01$ | 9810 | 0.941 | 0.787 | -0.0049 | 0.0037 | $1.85 \mathrm{E}-01$ | 7332 | 1 |
| Monocyte count-novel loci |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| rs2047076 | 5 | 76058509 | C | T | 6.03 | 1 | 33729 | Intergenic | 0.714 | 0.005 | 0.006 | $3.80 \mathrm{E}-01$ | 16550 | 0.999 | 0.9998 | 2.678 | 0.474 | $1.64 \mathrm{E}-08$ | 9,810 | 1 | 0.037 | 0.0023 | 0.0091 | $7.97 \mathrm{E}-01$ | 7369 | 0.789 |
| Neutrophil count-established loci |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| rs7667376 | 4 | 74967890 | C | T | 11.13 | 1 | 33743 | CXCL2 | 0.596 | 0.017 | 0.004 | $1.47 \mathrm{E}-04$ | 16550 | 0.972 | 0.542 | 0.017 | 0.005 | $5.96 \mathrm{E}-04$ | 9802 | 1 | 0.221 | -0.0428 | 0.008 | $8.50 \mathrm{E}-04$ | 7391 | , |
| rs445 | 7 | 92408370 | C | T | 10.52 | 1 | 33744 | CDK6 | 0.776 | 0.022 | 0.009 | $1.21 \mathrm{E}-02$ | 16550 |  | 0.685 | 0.024 | 0.005 | 8.96E-06 | 9802 | 1 | 0.203 | -0.0479 | 0.0092 | $1.15 \mathrm{E}-03$ | 7392 | 1 |
| rs8078723 | 17 | 38166879 | C | T | 28.06 | 1 | 33693 | CSF3, <br> MED24 | 0.626 | 0.043 | 0.004 | $2.84 \mathrm{E}-23$ | 16550 | 0.997 | 0.519 | 0.032 | 0.005 | $1.39 \mathrm{E}-10$ | 9802 | 0.995 | 0.678 | -0.0071 | 0.0073 | 6.19E-01 | 7341 | 1 |
| rs4794822 | 17 | 38156712 | C | T | 28.92 | 1 | 33753 | PSMD3 | 0.586 | -0.043 | 0.004 | $3.64 \mathrm{E}-23$ | 16550 | 0.999 | 0.481 | -0.032 | 0.005 | $7.09 \mathrm{E}-11$ | 9802 | 0.999 | 0.325 | 0.0048 | 0.0073 | $7.41 \mathrm{E}-01$ | 7401 | 0.918 |
| Neutrophil count-novel loci |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| rs6936204 | 6 | 32217092 | C | T | 6.45 | 0.048 | 33706 | AK123889 | 0.585 | 0.02 | 0.004 | $5.83 \mathrm{E}-06$ | 16550 | 0.991 | 0.912 | 0.032 | 0.009 | $3.97 \mathrm{E}-04$ | 9802 | 0.983 | 0.279 | 0.01 | 0.0074 | $4.68 \mathrm{E}-01$ | 7354 | 1 |

Strong evidence for association is defined as a BF of $>6$. Differences in effect alleles between cohorts were corrected before the MANTRA analysis.
PPA, posterior probability of association; SE, standard error; EAF, effect allele frequency.
Consortia summary of imputation qualtiy score.
*Proxy for rs2814778.


Figure 1. Manhattan plots subset by WBC subtype and cohort. Horizontal axis indicates the chromosomal position. Vertical axis for the blue Manhattan plots indicates $-\log _{10} P$-values from fixed-effects meta-analysis, and vertical axis for the black Manhattan plots indicates BFs from trans-ethnic meta-analysis. Significance values are truncated at 30 on the $y$-axis for clarity and scaling of images. A 50 Mb region of apparent significance surrounds the centromere of chromosome 1 and suggests two distinct peaks; however, this results from the lack of genotyped or imputed SNPs in the region and is a spatial inflation of a truly causal variant in the nearby DARC gene [Nalls et al. (9) and Reiner et al. (12)].
is also a receptor for $I L-8$. Additional novel associations for WBC include rs2163950 at 8 q 24.21 , located in an intergenic region, and rs6734238 in region 2 q13, located near IL1F10 and ILIRN. Top SNP rs6734238 tags the $I L-1$ gene family locus near 2q13 and has also been associated with C-reactive protein levels in a European ancestry population (22). Notably, a large region of the implicated chromosome 6 associations for WBC, tagged by rs2853946, contains an apparently bimodal signal of association, between 310.00 and 313.00 Mb (Fig. 2). A single novel association for monocyte count was identified by rs2047076 in region 5q13.3, also within an intergenic region of the genome. This region falls between $F 2 R$, associated with platelet count (23), and F2RL1 which encodes PAR-2, a monocyte receptor (24).

Variants showing strong evidence for association were examined for heterogeneity in their allelic effects across ancestries, indicated by a posterior probability of heterogeneity (PPH) of $>0.5$ (Table 3). For the novel monocyte-associated locus on chromosome 5 , the RIKEN cohort has a large posterior mean allelic effect (PMAE), whereas the COGENT and CHARGE cohorts have PMAE that are much smaller. This suggests that the association may be specific to the Japanese population, or that the variant tagging this region, rs2047076, may not be a good proxy among European or African-American individuals because of differing LD structure between these populations. The associations reported on chromosome 4 for both WBC and neutrophils exhibit allelic effects in opposite directions between the ancestry groups, which could reflect multiple risk variants, or differing LD structure (1). The novel associations reported on chromosome 2 for WBC have a posterior probability of association (PPA) of $\sim 0.03$, and the PMAE of these variants are similar across the three ancestries $(\sim 0.01)$. PPA is estimated from the weighted average of the alternative
models and accounts for the differences in likelihood and statistical power between tests (25). Locus plots for novel associations are shown in Figures 2-7, and locus plots for known associations are shown in Supplementary Material, Figures S1-S11.

To determine how consistent the MANTRA results are across studies, a random-effects MA was also performed using METAL. The results are largely supportive of the known and novel loci reported here, with the exception of those variants with high levels of heterogeneity between populations, such as the novel variant associated with monocytes and located on chromosome 5, rs2047076. The results of this additional MA are found in Supplementary Material, Table S1.

Cohort-level data were imputed before the trans-ethnic analysis was performed. In instances where a novel variant was both imputed across multiple cohorts and exhibits low allele frequencies within those cohorts, additional replication is needed to validate the associations made here. In particular, these include rs2047076, associated with monocytes, and rs2163950, associated with WBC count (Table 2). Without replication, these novel results should be viewed cautiously. We are optimistic that in time, a similarly sized, ancestry-matched cohort will be available to replicate these analyses. In the meantime, we have provided genome-wide summary statistics for the primary MANTRA analyses of monocyte, neutrophil and WBC count traits in Supplementary Material, Table S2.

## Fine mapping

Credible sets were defined to assess the extent to which the transethnic analysis improved fine-mapping resolution of known associations. Credible region summary data for associated loci are presented in Supplementary Material, Table S3.


Figure 2. Locus plot for Monocyte association on chromosome 5. Vertical axis indicates BF, and horizontal axes indicate both chromosomal position and gene location.

The major locus affecting WBC levels in African-Americans is located in the $D A R C$ gene of chromosome 1, spanning 900000 bp between 158724683 and 159624683 (3). The main variant associated with this signal is rs2814778, located at position (b37) 159174683 . Our analyses replicated this finding, identifying a number of significant hits surrounding rs2814778 (chr1: 159174683 ). The surrounding variants are the product of a well-established selective sweep, and proximity to the Duffy null mutations predisposed them to association (26). While rs2814778 was not included in all three cohorts due to the removal of monomorphic SNPs during quality control, credible region analyses of this region identify a single nearby variant, rs2518564 (chr1: 159062 436), as encompassing 99\% of the signal. As the functional variant in the $D A R C$ locus is present exclusively in the African-American population consortium, we did not expect the this region to show meaningful fine mapping. The use of a proxy variant in high LD with the other two cohorts
provides additional evidence of what is already known, this region is highly associated with WBC.

Other previously identified associations with WBC subtypes, however, were substantially narrowed. For example, the 7q21.2 region tagged soley by rs 445 in both WBC and neutrophil counts was found to encompass $99 \%$ of the association signal for these traits. This variant is located within an intronic section of $C D K 6$, a gene in the cyclin-dependent protein kinase family. At another previously identified locus for WBC and neutrophil count, located on chromosome 17, the association signal could be limited with $99 \%$ confidence to seven variants across a $\sim 15 \mathrm{~kb}$ region associated with WBC count and two variants across a $\sim 10 \mathrm{~kb}$ region associated with neutrophil count. The previously identified variant tagging this region, rs4794822, is within both of these SNP sets, but individually reaches only $72 \%$ confidence in WBC count and $88 \%$ confidence in neutrophil count.

Table 3. Heterogeneity of allelic effects

| Subtype | Variant details |  |  | MANTRA MA |  | CHARGE |  | RIKEN |  | COGENT |  |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  | SNP | Chr | Position (b37) | $\log _{10}(\mathrm{BF})$ | PPH | PMAE | PSD | PMAE | PSD | PMAE | PSD |
| Novel |  |  |  |  |  |  |  |  |  |  |  |
| WBC | rs10932765 | 2 | 219099484 | 6.95 | 0.025 | 0.01068 | 0.00182 | 0.01068 | 0.00182 | 0.01057 | 0.00205 |
| WBC | rs6734238 | 2 | 113841030 | 6.62 | 0.028 | -0.01193 | 0.00244 | -0.01181 | 0.00213 | -0.01195 | 0.00227 |
| WBC | rs1371799 | 4 | 74977837 | 13.5 | 1 | -0.01164 | 0.00207 | -0.01163 | 0.00207 | 0.02273 | 0.00408 |
| WBC | rs2853946 | 6 | 31247203 | 12.16 | 0.009 | -0.01379 | 0.00179 | -0.01379 | 0.00179 | -0.01378 | 0.00181 |
| WBC | rs2163950 | 8 | 130597585 | 7.75 | 0.065 | -0.01688 | 0.00287 | -0.01655 | 0.00304 | -0.01614 | 0.00346 |
| MONO | rs2047076 | 5 | 76058509 | 6.03 | 1 | 2.50643 | 0.47953 | 0.00458 | 0.00498 | 0.00457 | 0.00495 |
| NEU | rs7667376 | 4 | 74967890 | 11.13 | 1 | 0.011 | 0.00199 | 0.01104 | 0.00199 | -0.02275 | 0.00405 |
| NEU | rs6936204 | 6 | 32217092 | 6.45 | 0.048 | 0.01028 | 0.00356 | 0.0096 | 0.00259 | 0.00681 | 0.00477 |
| Known |  |  |  |  |  |  |  |  |  |  |  |
| WBC | rs2518564* | 1 | 159062436 | 341.59 | 1 | 0.00318 | 0.00241 | 0.00363 | 0.00258 | 0.19017 | 0.00469 |
| WBC | rs4895441 | 6 | 135426573 | 9.64 | 0.165 | 0.01368 | 0.00248 | 0.01282 | 0.00239 | 0.01169 | 0.00446 |
| WBC | rs445 | 7 | 92408370 | 16.8 | 1 | 0.01905 | 0.00254 | 0.01899 | 0.00297 | -0.02476 | 0.00476 |
| WBC | rs4794822 | 17 | 38156712 | 29.64 | 1 | -0.02088 | 0.00309 | -0.02491 | 0.00337 | 0.01249 | 0.00391 |
| MONO | rs1449263 | 2 | 182319301 | 19.24 | 1 | -0.03449 | 0.00399 | -0.03484 | 0.00378 | 0.00857 | 0.00316 |
| MONO | rs12988934 | 2 | 182323665 | 7.75 | 0.022 | -0.03872 | 0.00621 | -0.03853 | 0.00627 | NA | NA |
| MONO | rs9880192 | 3 | 128297569 | 8.3 | 1 | -0.03049 | 0.00687 | -0.02917 | 0.0046 | 0.00315 | 0.00483 |
| MONO | rs3095254 | 6 | 31221668 | 6.81 | 1 | 0.03449 | 0.00622 | 0.00473 | 0.00306 | 0.00439 | 0.00267 |
| MONO | rs1991866 | 8 | 130624105 | 15.3 | 1 | -0.0322 | 0.00418 | $-0.03217$ | 0.00392 | -0.00466 | 0.00316 |
| MONO | rs10980800 | 9 | 113915905 | 11.63 | 1 | 0.0087 | 0.01581 | 0.043 | 0.00581 | -0.00475 | 0.00379 |
| NEU | rs445 | 7 | 92408370 | 10.52 | 1 | 0.01905 | 0.00254 | 0.01899 | 0.00297 | -0.02476 | 0.00476 |
| NEU | rs8078723 | 17 | 38166879 | 28.06 | 1 | 0.01851 | 0.00278 | NA | NA | -0.01105 | 0.00378 |
| NEU | rs4794822 | 17 | 38156712 | 28.92 | 1 | -0.02088 | 0.00309 | -0.02491 | 0.00337 | 0.01249 | 0.00391 |

Variants encompassing the $95 \%$ credible region of an associated region, as identified using the top hits from the MANTRA analysis, are presented for each subtype. Ancestry-specific posterior mean allelic effects (PMAE) are reported. PSD is the Bayesian equivalent of standard error and characterizes the variance of the effect. PMAE is the posterior mean allelic effect; when these values are similar between ancestry cohorts, it suggests that simiar variants are responsible for the effect. When values are in opposite directions, it suggests multiple risk variants, or differing LD structure.
PPH, a posterior probability of heterogeneity; PPA, posterior probability of association; the probability that an SNP is truly associated with a phenotype.
*Proxy for rs2814778.

## Conditional analysis

Conditional analysis adjusting for the effect of the most significantly associated SNP at each locus was performed to assess the independence of possible novel variants and to detect the presence of any secondary association signals within known regions. Secondary signals were defined as additional associated variants within 500 kb of previously known loci. Of the primary (known and novel) loci, approximately half contained secondary signals; these include four signals associated with monocytes, three associated with neutrophils and six associated with WBC count. In some instances, the top signal identified by the secondary analysis was stronger than that observed in the primary analysis. This could occur, for example, when the allele frequencies of the initial, index SNP are similar across ancestries, but the conditional signal(s) more accurately tag a functional variant in the respective populations. In order to verify the authenticity of our conditional analysis results, we performed reverse conditioning on our secondary signals and found the signals reported here to remain significant, suggesting an independent effect on WBC subtypes. The top association signals from these conditional analyses are found in Table 4.

## Expression quantitative trait loci analysis

All known, conditional and novel loci were assessed as potential expression quantitative trait loci (eQTLs) in leukocyte-derived tissues in order to identify any correlations between association signals and gene expression, as such correlations may account
for functional relationships that are not captured by LD. Two known loci associated with monocyte count, and one novel locus newly associated with both WBC and neutrophil count, represented significant $(P<5 \mathrm{E}-05)$ eQTLs when assessing either the European ancestry sentinel SNPs or their proxies in the YRI (African) and ASN (Asian) populations (1). Three of the four index SNPs at these loci are located within chromosome 6 21.3; each has been associated with a different blood cell trait. The monocyte eQTLs on chromosome six are defined by the transcription factor gene TCF19; the neutrophil eQTLs relate primarily to the $H L A$ transcripts, but depending on the tissue type are also associated with expression of ATP6V1G2, which encodes an enzyme involved in eukaryotic cell compartment acidification. The WBC count eQTLs also relate to various transcripts of this region. Additionally, a previously described locus associated with monocyte count on chromosome 2 is an eQTL for the ITGA4 transcript at 2 q 31.3 , which encodes the integrin alpha- 4 subunit of the very late antigen- 4 receptor on monocytes and other mononuclear cells $(27,28)$. In instances where a proxy eQTL SNP has been used to represent a WBC or subtype-associated SNP, variants with greater concordance to the index SNP are viewed with more confidence than those with lower $R^{2}$ values. Only two eQTL SNPs were found to be concordant or in very high LD: rs3130320 $\left(R^{2}=1\right)$ for neutrophils and rs6740847 ( $R^{2}=0.983$ ) for monocytes. We have reported other SNPs as potential eQTLs if their $R^{2}$ value with the proxy eQTL SNP is $>0.5$; these SNPs are of course likely to be near the blood cell SNPs' physical location. All the proxy eQTL variants identified here are located in Supplementary Material, Table S4.


Figure 3. Locus plot for Neutrophil association on chromosome 6. Vertical axis indicates BF, and horizontal axes indicate both chromosomal position and gene location.

## DISCUSSION

We applied trans-ethnic MA to summary data from Japanese, African-American and European-Americans populations and identified six new regions that contain biologically plausible genetic loci associated with WBC traits. Many of the novel and secondary association signals we observed involve genomic regions that contain several inflammatory and immune cellrelated genes.

Particularly interesting, novel regions include the two loci on chromosome 2 associated with WBC count. The first, identified by rs6734238, falls within an inflammatory gene region of the interleukin- 1 cytokine gene family (29). This region has been associated with several inflammation-related biomarkers, including C-reactive protein and fibrinogen. rs6734238 is located downstream of ILIF10 and upstream of ILIRN. ILIRN encodes IL-1 receptor antagonist (IL-1RA), which regulates a
variety of interleukin-1-related immune and inflammatory responses, including inhibition of interleukin 1, alpha (IL1A) and interleukin 1, beta (IL1B). IL1F10 encodes IL-38, which regulates Th17 immune responses and stimulates IL-6 cytokine production from dendritic cells in vitro (30). The second chromosome 2 region, identified by rs10932765, is near $A P R C 2$ and $C X C R 1$ (also known as IL8RA). CXCR1 is of particular interest as this is a chemokine receptor involved in leukocyte chemotaxis and trafficking (31).

The novel chromosome 5 q 13 region associated with monocyte count lies within a family of protease-activated receptor genes, F2RL2-F2RL1-F2R. The F2RL1 gene (proteaseactivated receptor- 2 or PAR-2) has previously been related to some inflammatory and autoimmune diseases, and is a known receptor on monocytes (32). PAR-2 is a G protein-coupled receptor on monocyte/macrophages and other cell types that appear to have a direct role in the regulation of innate immune


Figure 4. Locus plot for White Blood Cell count association on chromosome 2 (first region). Vertical axis indicates BF, and horizontal axes indicate both chromosomal position and gene location.
function. Specifically, PAR-2 can be activated by a number of endogenous inflammation-associated proteinases (e.g. mast cell tryptase, trypsin and neutrophil proteinase 3) or exogenous patho-gen-derived proteinases (33). Notably, another member of the PAR gene family located on $5 q 13, F 2 R$, encodes PAR-1, the platelet thrombin receptor. Common variants of $F 2 R$ were recently associated with circulating platelet count in a European GWAS (23).

A single region of chromosome 6, associated with neutrophil and WBC count, is located near the HBS1L and MYB genes, which are known to be associated with fetal hemoglobin levels and monocyte counts. This locus has also been reported to be associated with red cell and platelet traits, but not previously with white cell traits. Although there is a single signal for neutrophils at this locus, the signal for WBC count appears bimodal. One of these two regions is captured by a single SNP with $99 \%$ confidence, while the second requires a regional span of nearly 1 Mb to reach the same level of confidence. This locus was
also significant in our eQTL analysis across all ancestry types for both neutrophil and WBC count, in lymphoblastoid cell lines (LCLs) and whole blood.

The region of chromosome 8 newly associated with WBC count lies near a gene of unknown function, GSDMC, which encodes gasdermin C. Other genes belonging to the gasdermin family have been associated with immune-mediated phenotypes such as asthma and alopecia $(34,35)$, suggesting a role for this gene family in inflammatory disorders (36).

A signal was observed on chromosome 16 for WBC count and neutrophils; however, the top associated variant is located within the intronic region of HYDIN, an mRNA transcript sequence involved in cilia motility. As previously reported, it is a likely homolog to the $D A R C$ region of chromosome 1 and represents a spurious signal (12).

The replicated region on chromosome 4, associated with both WBC and neutrophil counts, is located near a chemokine family


Figure 5. Locus plot for White Blood Cell count association on chromosome 2 (second region). Vertical axis indicates BF, and horizontal axes indicate both chromosomal position and gene location.
gene cluster, CXCL5-CXCL3-CXCL2. CXCL2 interacts with another chemokine receptor, $C X C R 2$, to control migration of leukocytes from the bone marrow (37). Taken together with the $D A R C$ locus, these findings extend the importance of common genetic variants of chemokine ligands and receptors in the regulation of WBC counts.

Using the proper reference panel is critical to the dependability and accuracy of this analysis. For this reason, ancestry-matched subsets of the 1000 Genomes were used as the reference panel from which LD was calculated, which are presumed to be drawn from the same general populations as those used here. However, the relatively small sample sizes available through 1000 Genomes increase the possibility of error in LD estimations (38). While the localization and resolution of functional variants may improve with the additional genomic variation measured in newer reference panels, we show that even without these newer
panels, the associations identified using MANTRA provide plausible candidates for functionality.

In conclusion, trans-ethnic meta-analyses allow for an examination of disease traits within a large population of individuals and provide the opportunity to localize previously known regions and detect novel ones, while considering the heterogeneity of allelic effects that may exist between contentially distinct populations. Additionally, our results illustrate the utility of trans-ethnic fine mapping for narrowing regions of association. Well-established loci replicated in the present study show credible intervals that flank the known index variant. For example, the previously known monocyte associations on chromosomes 8 and 9, shown in Supplementary Material, Table S3, have $99 \%$ credible intervals of only a few thousand base-pairs, located in close proximity to the originally identified variants. Our analysis of the DARC region is complicated by the causal


Figure 6. Locus plot for White Blood Cell count association on chromosome 6. Vertical axis indicates BF, and horizontal axes indicate both chromosomal position and gene location.

SNP being monomorphic in two of the three populations employed. When the credible interval analysis is applied to this region, the proxy variant (rs2518564) tagging the known functional variant (rs2814778) $(5,8)$ is identified as accounting for the entire signal. However, prior association studies and evidence of biological function allow confident identification of rs2814778 as the functional variant. By calculating credible intervals across test statistics from analyses of combined ancestries, we were able to narrow expansive loci to smaller regions. Further work is necessary to identify what functional variants may lie within these regions.

The increasing availability of GWAS summary data for many phenotypic traits of interest, from many ethnically diverse populations, suggests that the trans-ethnic GWAS MA approach can yield additional association signals, thereby explaining some of the missing heritability and genetic architecture for other complex traits. In addition, this work is relevant for future
targeted sequencing follow-up studies, as we have narrowed the scope of follow-up sequencing efforts for functional variants. By increasing the mapping resolution of the causal variants within these loci, we hope that these results guide nextgeneration targeted deep sequencing studies, which may disentangle the heterogeneity of effect across ethnicities (39). Future work will discern which functional variants are the same across ethnicities and which tag nearby regions, through LD, that harbor the true functional variant or variants.

## MATERIALS AND METHODS

The trans-ethnic GWA strategy was applied to three consortia containing WBC phenotypes. These include a Japanese population represented by RIKEN (Rikagaku Kenkyusho, Institute of


Figure 7. Locus plot for White Blood Cell count association on chromosome 8. Vertical axis indicates BF, and horizontal axes indicate both chromosomal position and gene location.

Physical and Chemical Research, Japan), a European ancestry population represented by the Cohorts for Heart and Aging Research in Genomic Epidemiology (CHARGE) Consortium and an African-American population represented by the Continental Origins and Genetic Epidemiology Network (COGENT) Consortium ( $9,11,12$ ). All three consortia contain measurements for total WBC count, neutrophil count and monocyte count, using the same scale of transformation and similar analytic paradigms. Clinical information of the subjects includes age, gender and smoking history, and was collected by self-report. Subject BMI was also collected as a measure. The laboratory data include total WBC count and subtypes, as determined using automated hematology cell counters according to the standardized protocol. The WBC phenotypes were natural logtransformed prior to analysis to provide a normal distribution. Samples $>2$ SD outside of the ethnicity specific mean for the given phenotype were excluded. This was done to ensure
normality of the included samples, and to exclude any subclinical inflammation such as the common cold. If a sample was identified as an outlier for one or more subtype, it was excluded entirely from the study.

The RIKEN study comprises over 17000 individuals from The BioBank Japan Project, which is made of up over 300000 subjects (http://biobankjp.org) (11). Samples determined to be of non-Japanese origin by either self-report or by principal components analyses (PCAs) in GWAS were excluded from further analyses. For the GWAS, 592232 SNPs were genotyped using Illumina HumanHap610-Quad Genotyping BeadChip. Subjects with call rates $<0.98$ were removed, as were SNPs with call rates $<0.99$. First- and second-degree relatives were excluded based on identity-by-descent analyses, as were SNPs with minor allele frequency (MAF) $<0.01$ or with Hardy-Weinberg equilibrium (HWE) $P$-values $<1.00 \mathrm{E}-7$. After quality control, genotypes were imputed using MACH 1.0 in a two-step procedure,

Table 4. Loci identified by conditional analysis

| Subtype | Primary top hits ${ }^{\text {a }}$ |  |  | Secondary top hits |  |  |  |  | $\log _{10} \mathrm{BF}$ | PPH | Samplesize | Effect direction |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  | SNP | Chr | Position | SNP | Position | Effect allele | Other allele | $N$ - <br> studies |  |  |  |  |
| MONO | rs1449263 | 2 | 182319301 | rs711801 | 182334873 | C | T | 3 | 10.46982 | 0.579 | 33711 | + + - |
| MONO | rs3095254 | 6 | 31221668 | rs2517774 | 29893982 | C | T | 2 | 72.9753 | 1 | 26360 | ? - - |
| MONO | rs2163952 | 8 | 130610389 | rs1457475 | 134988329 | A | G | 3 | 6.48727 | 1 | 28317 | - - + |
| MONO | rs12350763 | 9 | 113923723 | rs4401938 | 121179496 | C | T | 3 | 68.41786 | 0.99 | 33729 | +++ |
| NEU | rs7667376 | 4 | 74967890 | rs1440404 | 74944449 | C | G | 3 | 54.50984 | 1 | 33713 | + + - |
| NEU | rs445 | 7 | 92408370 | rs3731326 | 92327026 | A | G | 3 | 6.86032 | 1 | 33753 | + - - |
| NEU | rs4794822 | 17 | 38156712 | rs4794321 | 46028844 | C | T | 3 | 7.89538 | 1 | 33753 | - - + |
| WBC | rs6734238 | 2 | 113841030 | rs11899198 | 113840539 | G | T | 3 | 7.90504 | 1 | 52694 | $-++$ |
| WBC | rs1371799 | 4 | 74977837 | rs1440404 | 74944449 | C | G | 3 | 53.18811 | 1 | 52697 | + - - |
| WBC | rs9402686 | 6 | 135427817 | rs1890428 | 140253023 | C | T | 2 | 22.53076 | 1 | 22472 | +?- |
| WBC | rs445 | 7 | 92408370 | rs42626 | 89961237 | C | T | 3 | 11.88853 | 1 | 52740 | $-++$ |
| WBC | rs2163950 | 8 | 130597585 | rs10505542 | 130547253 | C | T | 3 | 19.59631 | 1 | 52686 | - - + |
| WBC | rs2241245 | 17 | 38151014 | rs8070454 | 38160754 | C | T | 2 | 22.03367 | 1 | 33231 | +? - |

Top hits from the GCTA conditional analyses are reported.
PPH, posterior probability of association; or, the probability that an SNP is truly associated with a phenotype. Effect direction order of studies: COGENT, CHARGE, RIKEN.
${ }^{\text {a }}$ Used as covariates in conditional analysis.
described in detail elsewhere $(40,41)$. HapMap Phase II Japanese individuals from Tokyo (JPT) and Han Chinese individiuals from Beijing (CHB) individuals were adopted as references. SNPs with imputation qualities $<0.30$ were excluded prior to analyses, and genomic control was applied to the cohort-level data. All participants provided written informed consent as approved by the ethical committees of the Center for Genomic Medicine, RIKEN and the Institute of Medical Science, the University of Tokyo (11).

The CHARGE consortium dataset is comprised of over 19000 individuals from seven discovery cohorts, including: the Rotterdam Study (RS), Framingham Heart Study (FHS), the NHLBI's Atherosclerosis Risk in Communities (ARIC) Study, the Age, Gene/Environment Susceptibility—Reykjavik Study (AGES), Health Aging and Body Composition study (HABC), the Baltimore Longitudinal Study of Aging (BLSA) and the Invecchaire in Chianti Study (inChianti) (9). Each of these studies, with the exception of the Framingham Heart Study, is comprised of unrelated individuals of confirmed European ancestry, based on PCAs. Prior to MA, SNPs with MAF $<0.01$, missingness $>5 \%$ or $\mathrm{HWE}<1.00 \mathrm{E}-7$ were excluded. Individuals with call rates $<95 \%$ were also excluded. After quality control, genotypes were imputed using the CEU reference panel of the HapMap Phase II haplotype data. The CHARGE consortium is comprised of MA data resulting from the summary statistics of these individual studies. Prior to the meta- analyses, study results were adjusted for genomic inflation factors, and SNPs with imputation quality $<0.30$ were excluded. Meta-analyses were performed using a fixed-effects model in METAL (42).

The COGENT consortium is comprised of over 16000 selfidentified African-Americans from seven discovery cohorts, including: Atherosclerosis Risk in Communities (ARIC), Coronary Artery Risk Development in Young Adults (CARDIA), Johns Hopkins Genetic Study of Atherosclerosis Risk (GeneSTAR), HealthyAging in Neighborhoods of Diversity across the Life Span (HANDLS), Health, Aging, and Body Composition (Health ABC), Jackson Heart Study (JHS) and the Women's Health Initiative (WHI) (12). SNPs were excluded from
cohort-level GWAS if MAF $<1 \%$ or missingness $>5 \%$. Monomorphic SNPs and ambiguously mapped SNPs were also removed. Individual samples exhibiting gender mismatch or genotype missingness $>10 \%$ were excluded. After quality control, genotypes were imputed with HapMap Phase II, using a $1: 1$ mixture of the CEU and YRI reference populations. Prior to MA, SNPs with imputation quality $<0.30$ were excluded. Study-specific GWA results were corrected for genomic inflation factors, and MA was performed using a fixed-effects model in METAL (42).

## Statistical analysis

Summary statistics for the RIKEN GWAS and the CHARGE and COGENT meta-analyses were collected and stratified by ethnicity and WBC subtype availability. Data were input into the trans-ethnic MA software package, MANTRA (15), which makes use of a prior model of relatedness between studies corresponding to Fst, or mean effect allele frequency differences between populations. Relatedness is determined by differences in allele frequency between studies. MANTRA estimates the BF in favor of association for each SNP using a Markov chain Monte Carlo (MCMC) algorithm. Results are reported as $\log _{10}$ (BF), and associations of 6 or greater have the highest posterior odds of being truly present $(43,44)$. Posterior probability of heterogeneity is also reported to examine levels of variation in allelic effects across the populations used in the analysis. Combining results across studies using a Bayesian approach is advantageous, as the evidence produced by this study is directly comparable to future studies performed in the same way. Simulations of distinct ancestry populations show that when MANTRA is compared with random-effects and fixed-effects meta-analyses, MANTRA shows increased performance and produces the highest-powered results in the detection of novel associations ( $15,22,45-47$ ).

Since the initial imputation of the datasets used here, a number of more comprehensive reference panels have been released including the latest HapMap release and the samples available
through the 1000 Genomes Project. As these reference panels contain more individuals and greater genome coverage, more genotypes are predicted with greater confidence than when using prior HapMap releases. Imputing raw data to the latest release of 1000 Genomes would be ideal; however, due to the data-sharing requirements of the cohorts included in this analysis, only summary statistics were available for these datasets. However, MANTRA is still expected to outperform a traditional MA in this case, as MANTRA accounts for heterogeneity while making no assumptions about differences or similarities in allelic effect.

To quantify uncertainty surrounding the top hits from the trans-ethnic MANTRA analysis, we calculated 95 and $99 \%$ credible regions (48). We estimated credible sets of SNPs by first defining a 1 Mb genomic region surrounding lead SNPs ( $\pm 500 \mathrm{~kb}$ ), then ranking the regional SNPs within this region according to their BF and then combining the cumulative posterior probabilities of these ranked SNPs until 95 and $99 \%$ confidence was reached.

In addition, the top hits from MANTRA were input into a conditional analysis in order to identify additional association signals at nearby susceptibility loci and to determine independence of these secondary signals from the index SNP association. As complex diseases are assumed to be influenced by two or more genes acting in concert, it is possible that prior GWAS aimed at identifying single loci have not detected secondary signals. Thus, when large sample sizes are available, conditional approaches can be useful in detecting secondary association signals with loci that initially appear to contribute a negligible risk to disease susceptibility. In addition, as evidence of association is predicated on a given conditioned SNP, it is possible that, in some instances, the secondary associations are stronger than what was observed in the primary single-ethnicity GWAS analysis. This is possible when the allele frequencies of a given SNP are similar across ancestries, as conditional hits may be closer to a functional variant than the original, single-population hit.

We used the software program Genome-wide Complex Trait Analysis (GCTA) v1.13 to perform conditional association analysis for each ancestry-specific set of summary results (49). As individual genotype data were unavailable, this was performed separately for each cohort using summary statistics and incorporating LD information from ancestry-matched reference samples containing individual-level genotype data. When original genotype data are not available, it is essential that the reference samples be from the same population as the original data, so that the LD structure estimated from the reference population is not biased. It is also critical that the reference sample is not affected by cryptic relatedness or population stratification. This is particularly relevant to admixed populations, such as COGENT, which is comprised of African-American individuals. In order to avoid confounding the genetic relationship matrix (GRM) produced by GCTA, eigenvectors are included in the model as covariates, to capture and account for any variance that is present due to population structure (49). In this analysis, we used ancestry-matched subsets from the 1000 Genomes Project to estimate LD structure within our samples (17). Using these samples as LD proxies, the GCTA association analyses were conditioned on the top hits from MANTRA, specific to each locus of interest. Independently associated SNPs were selected using a stepwise model selection procedure. Analyses were performed separately for each ancestry cohort.

The results from each cohort were then meta-analyzed across ethnicities.

In addition to the trans-ethnic association analysis, a secondary analysis exploring eQTLs was performed using SNAP (50). SNAP is a web server that identifies and annotates nearby proxy SNPs in LD (according to HapMap) to those queried. Using ancestry-specific tissues, SNAP was identified alias SNPs for significant index SNPs, and proxy SNPs in high linkage disequilibrium $\left(R^{2}>0.5\right)$. Sentinel, alias and proxy SNPs were queried within a collected database of expression SNP (eSNP) results, drawn from the following leukocyte-derived tissues: fresh lymphocytes (51), fresh leukocytes (52), leukocyte samples in individuals with celiac disease (53), whole blood samples (54-56), LCLs derived from asthmatic children $(57,58)$, HapMap LCL from three populations (59), a separate study on HapMap CEU LCL (60) and additional LCL population samples [(61-63); Mangravite et al., unpublished], CD19+ B cells (64), primary PHA (phytohaemagglutinin)-stimulated T cells (61), CD4 +T cells (65), peripheral blood monocytes $(27,64,66)$, CD11 + dendritic cells before and after Mycobacterium tuberculosis infection (67) and micro-RNA QTLs queried for LCL (68). The collected eSNP results met criteria for statistical thresholds of association with gene transcript levels, as described in the original cited papers. In cases where an SNP was associated with a transcript, we further examined the strongest eSNP for the transcript within that dataset and the LD between the strongest eSNP and blood count-selected eSNPs. This was done to assess the concordance of the blood count and expression signals.

## SUPPLEMENTARY MATERIAL

Supplementary Material is available at $H M G$ online.

## ACKNOWLEDGEMENTS

This study utilized the high-performance computational capabilities of the Biowulf Linux cluster at the National Institutes of Health, Bethesda, MD (http://biowulf.nih.gov).

Conflict of Interest statement. None declared.

## FUNDING

Y.O. is supported by a grant from the Japan Society of the Promotion of Science (JSPS). The development of the software package MANTRA was supported by Wellcome Trust (grant nos WT098017, WT090532 and WT064890). S.K.G. was supported in part by P30HL101290. Funding for CHARGE was made possible by NIA/NIH contract AG000932-2 (2009) Characterization of Normal Genomic Variability. The Age, Gene/ Environment Susceptibility Reykjavik Study is funded by NIH contract N01-AG-12100, the NIA Intramural Research Program, Hjartavernd (the Icelandic Heart Association) and the Althingi (the Icelandic Parliament). The Atherosclerosis Risk in the Communities Study is carried out as a collaborative study supported by National Heart, Lung, and Blood Institute contracts (HHSN268201100005C, HHSN268201100006C, HHSN2682 01100007C, HHSN268201100008C, HHSN268201100009C, HHSN268201100010C, HHSN268201100011C and HHSN268

201100012C), R01HL087641, R01HL59367 and R01HL086694; National Human Genome Research Institute contract (U01HG 004402) and National Institutes of Health contract (HHSN 268200625226 C ). The authors thank the staff and participants of the ARIC study for their important contributions. Infrastructure was partly supported by grant number UL1RR025005, a component of the National Institutes of Health and NIH Roadmap for Medical Research. The National Heart, Lung, and Blood Institute's Framingham Heart Study is a joint project of the National Institutes of Health and Boston University School of Medicine and was supported by the National Heart, Lung, and Blood Institute's Framingham Heart Study (contract No. N01-HC-25195) and its contract with Affymetrix for genotyping services (contract No. N02-HL-6-4278). Analyses reflect the efforts and resource development from the Framingham Heart Study investigators participating in the SNP Health Association Resource (SHARe) project. A portion of this research was conducted using the Linux Cluster for Genetic Analysis (LinGA-II) funded by the Robert Dawson Evans Endowment of the Department of Medicine at the Boston University School of Medicine and Boston Medical Center. The Health ABC Study was supported in part by the Intramural Research Program of the NIH, National Institute on Aging, NIA contracts N01AG62101, N01AG62103 and N01AG 62106. The GWAS was funded by NIA grant 1R01AG03209801A1 to Wake Forest University Health Sciences and genotyping services were provided by the Center for Inherited Disease Research (CIDR). CIDR is fully funded through a federal contract from the National Institutes of Health to The Johns Hopkins University (contract number HHSN268200782096C). The InChianti Study was supported as a 'targeted project' (ICS 110.1RS97.71) by the Italian Ministry of Health, by the US National Institute on Aging (contracts N01-AG-916413, N01-AG-821336, 263 MD 916413 and 263 MD 821336) and in part by the Intramural Research Program, National Institute on Aging, National Institutes of Health, USA. The generation and management of GWAS genotype data for the Rotterdam Study is supported by the Netherlands Organization of Scientific Research NWO Investments (no. 175.010.2005.011, 911-03-012), the Research Institute for Diseases in the Elderly (014-93-015; RIDE2), EUROSPAN (European Special Populations Research Network; LSHG-CT-2006-01947), the Netherlands Organization for Scientific Research (Pionier, 047.016.009, 047.017. 043;050-060-810), Erasmus Medical Center and the Centre for Medical Systems Biology (CMSB I and II and Grand; National Genomics Initiative) of the Netherlands Genomics Initiative (NGI); The Rotterdam Study is further funded by Erasmus Medical Center and Erasmus University, Rotterdam, Netherlands Organization for the Health Research and Development (ZonMw), the Research Institute for Diseases in the Elderly (RIDE), the Ministry of Education, Culture and Science, the Ministry for Health, Welfare and Sports, the European Commission (DG XII) and the Municipality of Rotterdam. Funding for COGENT was obtained through the Broad Institute (N01-HC65226) to create this genotype/phenotype database for wide dissemination to the biomedical research community. The Atherosclerosis Risk in Communities Study is carried out as a collaborative study supported by National Heart, Lung, and Blood Institute contracts (HHSN268201100005C, HHSN268 201100006C, HHSN268201100007C, HHSN268201100008C,

HHSN268201100009C, HHSN268201100010C, HHSN26820 1100011 C and HHSN268201100012C), R01HL087641, R01HL 59367 and R01HL086694; National Human Genome Research Institute (contract U01HG004402) and National Institutes of Health (contract HHSN268200625226C). The authors thank the staff and participants of the ARIC study for their important contributions. Infrastructure was partly supported by grant number UL1RR025005, a component of the National Institutes of Health and NIH Roadmap for Medical Research. ARIC is also supported by University of North Carolina at Chapel Hill (funded by N01-HC-55015), Baylor Medical College (N01-HC-55016), University of Mississippi Medical Center (N01-HC55021), University of Minnesota (N01-HC-55019), Johns Hopkins University (N01-HC-55020), University of Texas Houston (N01-HC-55017) and University of North Carolina (N01-HC-55018). Coronary Artery Risk in Young Adults (CARDIA): University of Alabama at Birmingham (N01-HC48047), University of Minnesota (N01-HC-48048), Northwestern University (N01-HC-48049), Kaiser Foundation Research Institute (N01-HC-48050), University of Alabama at Birmingham (N01-HC-95095), Tufts-New England Medical Center (N01-HC-45204), Wake Forest University (N01-HC45205), Harbor-UCLA Research and Education Institute (N01-HC-05187), University of California, Irvine (N01-HC-45134 and N01-HC-95100). Jackson Heart Study (JHS): Jackson State University (N01-HC-95170), University of Mississippi (N01-HC-95171), Tougaloo College (N01-HC-95172). Healthy Aging in Neighborhoods of Diversity across the Life Span Study (HANDLS): This research was supported by the Intramural Research Program of the NIH, National Institute on Aging and the National Center on Minority Health and Health Disparities (intramural project \# Z01-AG000513 and human subjects protocol \# 2009-149). Health ABC: This research was supported by NIA contracts N01AG62101, N01AG62103 and N01AG62106. The GWAS was funded by NIA grant 1R01AG032098-01A1 to Wake Forest University Health Sciences and genotyping services were provided by the Center for Inherited Disease Research (CIDR). CIDR is fully funded through a federal contract from the National Institutes of Health to The Johns Hopkins University (contract number HHSN268200782096C). This research was supported in part by the Intramural Research Program of the NIH, National Institute on Aging. GeneSTAR Acknowledgement: This research was supported by the National Heart, Lung, and Blood Institute (NHLBI) through the PROGENI (U01 HL72518) and STAMPEED (R01 HL087698-01) consortia. Additional support was provided by grants from the NIH/National Institute of Nursing Research (R01 NR08153) and the NIH/National Center for Research Resources (M01-RR000052) to the Johns Hopkins General Clinical Research Center. WHI: The WHI program is funded by the National Heart, Lung, and Blood Institute, National Institutes of Health, US Department of Health and Human Services, through contracts N01WH22110, 24152, 32100-2, 32105-6, 32108-9, 32111-13, 32115, 32118-32119, 32122, 42107-26, 42129-32 and 44221. The Cardiovascular Health Study (CHS) is supported from contracts N01-HC-35129, N01-HC-45133, N01-HC-75150, N01-HC-85079 through N01-HC-85086, N01 HC-15103, N01 HC-55222 and U01 HL080295 from the National Heart, Lung, and Blood Institute, with additional contribution from the National Institute of Neurological Disorders and Stroke. Additional support for this
work was provided by NIH R01 HL71862-06 and ARRA N000949304 (to A.P.R.). Some of the results of this paper were obtained by using the program package S.A.G.E., which is supported by a US Public Health Service Resource Grant (RR03655) from the National Center for Research Resources. The funders had no role in study design, data collection and analysis, decision to publish or preparation of the manuscript. Funding for RIKEN and the BioBank Japan Project was supported by Ministry of Education, Culture, Sports, Science and Technology, Japan. The funders had no role in study design, data collection and analysis, decision to publish or preparation of the manuscript.

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## Supplemental Materials <br> Working Groups

CHARGE (Cohorts for Heart and Aging Research in Genome Epidemiology)
Sampath Arepalli, Stefania Bandinelli, Alessandro Biffi, Joshua C. Bis, Eric Boerwinkle, Aravinda Chakravarti, Ming-Huei Chen, Sean Chong, Josef Coresh, David J. Couper, L. Adrienne Cupples, Abbas Dehghan, Angela Do'ring, Gudny Eiriksdottir, Janine F. Felix, Luigi Ferrucci, Aaron R. Folsom, Caroline S. Fox, Timothy M. Frayling, Santhi K. Ganesh, Melissa Garcia, Stephen F. Garner, Paolo Gasparini, Christian Gieger, Nicole L. Glazer, Natalia A. Gouskova, Andreas Greinacher, Vilmundur Gudnason, Tamara B. Harris, Dena G. Hernandez, Albert Hofman, Thomas Illig, Yoichiro Kamatani, Naoyuki Kamatani, Michiaki Kubo, Brigitte Kuhnel, Vasiliki Lagou, Guillaume Lettre, Daniel Levi, JingPing Lin, Yongmei Liu, Dan L. Longo, Thomas Lumley, Massimo Mangino, Koichi Matsuda, Christa Meisinger, David Melzer, Stephan Menzel, Matt Moore, Yusuke Nakamura, Michael A. Nalls, Matthias Nauck, Christopher J. O'Donnell, Yukinori Okada, Ben A. Oostra, Willem H. Ouwehand, Kushang V. Patel, Nicola Pirastu, Giorgio Pistis, Holger Prokisch, Inga Prokopenko, Bruce M. Psaty, Alex P. Reiner, Augusto Rendon, Jennifer Sambrook, Andrew B. Singleton, Albert V. Smith, Nicole Soranzo, Tim D. Spector, Jonathan Stephens, Michael Stumvoll, Atsushi Takahashi, Toshiko Tanaka, Toshihiro Tanaka, Kent Taylor, Alexander Teumer, Swee Lay Thein, Anke To'njes, Daniela Toniolo, Tatsuhiko Tsunoda, Andre' G. Uitterlinden, Cornelia M. van Duijn, Frank J. A. van Rooij, Uwe Vo'lker, Henry Vo'lzke, H.-Erich Wichmann, Kerri L. Wiggins, James G. Wilson, Jacqueline C. M. Witteman, Andrew R. Wood, Kazuhiko Yamamoto, Qiong Yang, Neil A. Zakai

## CHARGE genotyping platforms by study:

- AGES-Reykjavik Study - Illumina 370CNV BeadChip array
- ARIC - Affymetrix Genome-Wide Human SNP Array 6.0 platform
- $R S$ - Illumina 550K BeadChip array
- FHS - Affymetrix 500 K array and an additional gene-focused 50 K array
- inChianti - Illumina 550K BeadChip array
- BLSA - Infinium II HumanHap550 v. 1, Infinium II HumanHap550 v. 3, or a composite of Infinium HumanHap300 and Infinium II 240S


## COGENT

Sampath Arepalli, Melissa A. Austin, Diane M. Becker, Angela Britton, Zhao Chen, David Couper, J. David Curb, Eric Dean, Charles B. Eaton, Michele K. Evans, Aaron R. Folsom, Myriam Fornage, Santhi K. Ganesh, Struan F. A. Grant, Tamara B. Harris, Dena Hernandez, Naoyuki Kamatini, Brendan J. Keating, Michiaki Kubo, Andrea LaCroix, Leslie A. Lange, Guillaume Lettre, Simin Liu, Yongmei Liu, Kurt Lohman, Rasika Mathias, Yan Meng, Emile R. Mohler III, Solomon Musani, Yusuke Nakamura, Michael A. Nalls, Christopher J. O’Donnell, Yukinori Okada, Cameron D. Palmer, George J. Papanicolaou, Kushang V. Patel, Alexander P. Reiner, Andrew B. Singleton, Beverly M. Snively, Atsushi Takahashi, Hua Tang, Herman A. Taylor Jr., Kent Taylor, Cynthia Thomson, James G. Wilson, Lisa R. Yanek, Lingyao Yang, Elad Ziv, Alan B. Zonderman

COGENT genotyping platforms by study:
ARIC - Affymetrix Genome-Wide Human SNP Array 6.0 platform
CARDIA - Affymetrix Genome-Wide Human SNP Array 6.0 platform
GeneSTAR - Illumina BeadChip array

## HANDLS - Illumina BeadChip array

WHI - Affymetrix Affymetrix Genome-Wide Human SNP Array 6.0 platform
HealthABC - Illumina Human1M-Duo BeadChip platform

## RIKEN

Koichiro Higasa, Tomomitsu Hirota, Naoya Hosono, Yoichiro Kamatani, Naoyuki Kamatani, Michiaki Kubo, Natsuhiko Kumasaka, Koichi Matsuda, Yusuke Nakamura, Hiroko Ohmiya, Yukinori Okada, Atsushi Takahashi, Mayumi Tamari, Toshiko Tanaka, Toshihiro Tanaka, Tatsuhiko Tsunoda, Yumi Yamaguchi-Kabata, Kazuhiko Yamamoto

## Supplemental eQTL analysis

We applied expression quantitative trait loci (eQTL) analysis to significant index SNPs using SNAP (18974171), and identified proxy SNPs in high linkage disequilibrium ( $\mathrm{r}^{2}>0.8$ ). Index and proxy SNPs were searched for within a collected database of expression SNP (eSNP) results including the following tissues: fresh lymphocytes (17873875), fresh leukocytes (19966804), leukocyte samples in individuals with Celiac disease (19128478), whole blood samples (18344981, 21829388, 22692066), lymphoblastoid cell lines (LCL) derived from asthmatic children (17873877, 23345460), HapMap LCL from 3 populations (17873874), a separate study on HapMap CEU LCL (18193047), additional LCL population samples (19644074, 22286170,22941192, Mangravite et al., unpublished), CD19+ B cells (22446964), primary PHAstimulated T cells (19644074), CD4+ T cells (20833654), peripheral blood monocytes (19222302,20502693, 22446964), CD11+ dendritic cells before and after Mycobacterium tuberculosis infection (22233810), omental and subcutaneous adipose (18344981, 21602305, 22941192), stomach (21602305), endometrial carcinomas (21226949), ER+ and ER- breast cancer tumor cells (23374354), brain cortex (19222302, 19361613, 22685416), pre-frontal cortex (22031444, 20351726), frontal cortex (20485568), temporal cortex (20485568, 22685416), pons (20485568), cerebellum (20485568, 22685416), 3 additional large studies of brain regions including prefrontal cortex, visual cortex and cerebellum, respectively (Emilsson, Cell in press), liver (18462017, 21602305, 21637794), osteoblasts (19654370), lung (23209423), skin (21129726, 22941192) and primary fibroblasts (19644074). Micro-RNA QTLs were also queried for LCL (21691150), and gluteal and abdominal adipose (22102887). The cited original papers describe the criteria of association with gene transcript levels for eSNP results. The best eSNPs associated with a transcript were further examined using DIAGRAM, and LD between the DIAGRAM-selected eSNPs and the strongest associated eSNP was compared to determine the concordance of DIAGRAM and expression signals.

## Supplemental Figure Legends

Figure S1-11. Locus plots for known hits. Vertical axis indicates Bayes Factor, and horizontal axes indicate both chromosomal position and gene location.

## Supplemental Table Legends

Table S1. Random-effects meta-analysis of MANTRA identified loci.
Table S2.* Genome-wide summary statistics of the primary MANTRA analyses of Monocyte, Neutrophil, and White Blood Cell counts.
Table S3. Credible regions identified from MANTRA top hits are listed.
Table S4. Summarized eQTL analysis are reported.
*Located as a linked file through HMG’s sharing site.












Table S1. Random-effects meta-analysis of MANTRA identified loci.

| Phenotype | SNP | Chr | Position | A1 | A2 | Summary Frequency | Effect | Standard Error | P-value | Direction | Heterogeneity P-value |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| Monocytes | rs10209150 | 2 | 182329619 | a | g | 0.8526 | -0.0065 | 0.006 | 2.83E-01 | ?+- | $1.71 \mathrm{E}-01$ |
| Monocytes | rs10930969 | 2 | 182327957 | a | g | 0.1562 | -0.0126 | 0.0038 | 8.25E-04 | --+ | $2.27 \mathrm{E}-02$ |
| Monocytes | rs13415157 | 2 | 182330844 | a | c | 0.9791 | 0.0111 | 0.0119 | $3.51 \mathrm{E}-01$ | ??+ | $1.00 \mathrm{E}+00$ |
| Monocytes | rs1375493 | 2 | 182323766 | a | g | 0.5217 | 0.0192 | 0.0024 | $1.86 \mathrm{E}-15$ | +++ | 4.23E-06 |
| Monocytes | rs1449260 | 2 | 182333447 | a | g | 0.3123 | -0.0153 | 0.0026 | 4.01E-09 | --- | $1.65 \mathrm{E}-02$ |
| Monocytes | rs1449263 | 2 | 182319301 | $t$ | c | 0.4227 | 0.0087 | 0.0024 | 3.09E-04 | ++- | 1.28E-17 |
| Monocytes | rs155138 | 2 | 182319878 | a | C | 0.9485 | 0.0082 | 0.0071 | $2.45 \mathrm{E}-01$ | ??+ | $1.00 \mathrm{E}+00$ |
| Monocytes | rs155141 | 2 | 182317202 | a | t | 0.1367 | -0.0107 | 0.0034 | $1.68 \mathrm{E}-03$ | --- | $1.99 \mathrm{E}-02$ |
| Monocytes | rs155146 | 2 | 182332350 | t | c | 0.0589 | -0.0086 | 0.0066 | $1.93 \mathrm{E}-01$ | ?+- | $2.77 \mathrm{E}-01$ |
| Monocytes | rs155147 | 2 | 182332146 | a | g | 0.0981 | 8.00E-04 | 0.0049 | $8.64 \mathrm{E}-01$ | ?-+ | 2.26E-01 |
| Monocytes | rs155148 | 2 | 182331265 | a | g | 0.7178 | 0.0144 | 0.0038 | $1.44 \mathrm{E}-04$ | +++ | 5.26E-02 |
| Monocytes | rs155149 | 2 | 182329647 | t | g | 0.6017 | -0.002 | 0.0028 | $4.70 \mathrm{E}-01$ | --+ | 3.25E-11 |
| Monocytes | rs2124440 | 2 | 182328214 | a | g | 0.5145 | 0.0188 | 0.0024 | 2.37E-15 | +++ | 6.86E-07 |
| Monocytes | rs3731835 | 2 | 182321179 | t | C | 0.9794 | 0.0113 | 0.0118 | $3.38 \mathrm{E}-01$ | ??+ | $1.00 \mathrm{E}+00$ |
| Monocytes | rs3770132 | 2 | 182332961 | a | g | 0.7067 | 0.0139 | 0.0052 | 8.13E-03 | +++ | $1.53 \mathrm{E}-01$ |
| Monocytes | rs3770136 | 2 | 182329460 | t | c | 0.5251 | -0.012 | 0.0042 | 4.08E-03 | --- | 7.20E-02 |
| Monocytes | rs3770137 | 2 | 182326136 | c | g | 0.719 | 0.0197 | 0.0038 | $1.77 \mathrm{E}-07$ | +++ | $9.79 \mathrm{E}-01$ |
| Monocytes | rs3770138 | 2 | 182325907 | t | C | 0.4876 | -0.0129 | 0.0038 | $6.40 \mathrm{E}-04$ | --+ | 2.96E-02 |
| Monocytes | rs10188326 | 2 | 182311519 | t | g | 0.9237 | -0.0079 | 0.0057 | $1.69 \mathrm{E}-01$ | ??- | $1.00 \mathrm{E}+00$ |
| Monocytes | rs10930969 | 2 | 182327957 | a | g | 0.1562 | -0.0126 | 0.0038 | 8.25E-04 | --+ | 2.27E-02 |
| Monocytes | rs12988934 | 2 | 182323665 | $t$ | c | 0.2469 | 0.0382 | 0.0063 | 1.32E-09 | ++? | 5.90E-01 |
| Monocytes | rs13384671 | 2 | 182311594 | a | g | 0.6097 | 0.0123 | 0.0027 | 6.36E-06 | +++ | 1.83E-05 |
| Monocytes | rs1375493 | 2 | 182323766 | a | g | 0.5217 | 0.0192 | 0.0024 | $1.86 \mathrm{E}-15$ | +++ | 4.23E-06 |
| Monocytes | rs1449263 | 2 | 182319301 | t | c | 0.4227 | 0.0087 | 0.0024 | 3.09E-04 | ++- | $1.28 \mathrm{E}-17$ |
| Monocytes | rs155123 | 2 | 182312272 | a | c | 0.6856 | 0.0166 | 0.0036 | 3.61E-06 | +?+ | 6.03E-03 |
| Monocytes | rs155138 | 2 | 182319878 | a | c | 0.9485 | 0.0082 | 0.0071 | $2.45 \mathrm{E}-01$ | ??+ | $1.00 \mathrm{E}+00$ |
| Monocytes | rs155141 | 2 | 182317202 | a | t | 0.1367 | -0.0107 | 0.0034 | $1.68 \mathrm{E}-03$ | --- | $1.99 \mathrm{E}-02$ |
| Monocytes | rs2124440 | 2 | 182328214 | a | g | 0.5145 | 0.0188 | 0.0024 | 2.37E-15 | +++ | 6.86E-07 |
| Monocytes | rs3731835 | 2 | 182321179 | t | C | 0.9794 | 0.0113 | 0.0118 | $3.38 \mathrm{E}-01$ | ??+ | $1.00 \mathrm{E}+00$ |
| Monocytes | rs3770137 | 2 | 182326136 | c | g | 0.719 | 0.0197 | 0.0038 | $1.77 \mathrm{E}-07$ | +++ | $9.79 \mathrm{E}-01$ |
| Monocytes | rs3770138 | 2 | 182325907 | t | C | 0.4876 | -0.0129 | 0.0038 | $6.40 \mathrm{E}-04$ | --+ | 2.96E-02 |
| Monocytes | rs2010527 | 3 | 128303083 | c | g | 0.5977 | 0.0103 | 0.0029 | $3.95 \mathrm{E}-04$ | +++ | 4.53E-04 |
| Monocytes | rs2465354 | 3 | 128294897 | t | c | 0.6569 | -0.003 | 0.0034 | $3.65 \mathrm{E}-01$ | +?- | $5.07 \mathrm{E}-01$ |
| Monocytes | rs2712421 | 3 | 128287743 | t | g | 0.3218 | 0.0038 | 0.003 | 2.11E-01 | ++- | 1.67E-04 |
| Monocytes | rs2712423 | 3 | 128290938 | t | g | 0.6312 | 0.0079 | 0.003 | 9.02E-03 | +++ | 2.72E-01 |
| Monocytes | rs2713575 | 3 | 128294355 | a | g | 0.4891 | -8.00E-04 | 0.0027 | 7.81E-01 | -?- | 7.33E-01 |

Table S1. Random-effects meta-analysis of MANTRA identified loci.

| Monocytes | rs2713589 | 3 | 128290208 | t | c | 0.5501 | 7.00E-04 | 0.0027 | 7.93E-01 | -++ | $8.55 \mathrm{E}-01$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| Monocytes | rs2734046 | 3 | 128292213 | a | g | 0.1691 | 0.0016 | 0.0042 | 7.11E-01 | -++ | 7.76E-01 |
| Monocytes | rs6772849 | 3 | 128306418 | t | c | 0.3745 | -0.0093 | 0.0026 | 3.86E-04 | --- | $1.71 \mathrm{E}-04$ |
| Monocytes | rs6798431 | 3 | 128303373 | t | C | 0.5665 | 0.0102 | 0.0025 | $4.00 \mathrm{E}-05$ | +++ | $4.32 \mathrm{E}-05$ |
| Monocytes | rs7611156 | 3 | 128296868 | t | c | 0.7322 | -0.0049 | 0.0039 | $2.12 \mathrm{E}-01$ | -+- | $2.98 \mathrm{E}-01$ |
| Monocytes | rs9880192 | 3 | 128297569 | c | $g$ | 0.2892 | -0.0134 | 0.0033 | 5.27E-05 | --+ | 2.28E-06 |
| Monocytes | rs166175 | 5 | 76062954 | a | g | 0.671 | 0.0032 | 0.0034 | 0.3449 | +++ | 0.7236 |
| Monocytes | rs193943 | 5 | 76049832 | t | c | 0.4632 | 0.0017 | 0.0027 | 0.531 | +-+ | 0.7421 |
| Monocytes | rs2047076 | 5 | 76058509 | $t$ | c | 0.4863 | -0.0046 | 0.005 | 0.3544 | --- | 2.41E-07 |
| Monocytes | rs2054192 | 5 | 76048623 | a | g | 0.6932 | 3.00E-04 | 0.0038 | 0.9271 | --+ | 0.1092 |
| Monocytes | rs250718 | 5 | 76050452 | c | g | 0.6716 | 0.0033 | 0.0034 | 0.3259 | ++- | 0.08492 |
| Monocytes | rs250719 | 5 | 76050491 | a | g | 0.5465 | 0.0032 | 0.0025 | 0.197 | +++ | 0.9672 |
| Monocytes | rs250720 | 5 | 76051465 | a | t | 0.5113 | 0.0019 | 0.0025 | 0.4546 | +++ | 0.9669 |
| Monocytes | rs250723 | 5 | 76056054 | t | C | 0.6509 | 0.0031 | 0.003 | 0.2977 | +-+ | 0.6892 |
| Monocytes | rs250724 | 5 | 76061256 | a | c | 0.4776 | -0.002 | 0.0025 | 0.4279 | +-- | 0.8452 |
| Monocytes | rs250726 | 5 | 76059981 | a | g | 0.3473 | -0.0043 | 0.0028 | 0.1163 | +-- | 0.5026 |
| Monocytes | rs250727 | 5 | 76059301 | t | c | 0.4739 | -4.00E-04 | 0.0024 | 0.8828 | -+- | 0.7668 |
| Monocytes | rs250728 | 5 | 76059094 | t | g | 0.5791 | -0.0023 | 0.003 | 0.4411 | --+ | 0.544 |
| Monocytes | rs250729 | 5 | 76059055 | a | g | 0.4825 | -0.0013 | 0.0024 | 0.5883 | --- | 0.9574 |
| Monocytes | rs250759 | 5 | 76067728 | a | g | 0.422 | -0.0035 | 0.0024 | 0.1447 | +-- | 0.7495 |
| Monocytes | rs250760 | 5 | 76067624 | t | g | 0.6115 | -0.0025 | 0.0033 | 0.445 | --- | 0.615 |
| Monocytes | rs250761 | 5 | 76067376 | t | c | 0.5567 | -0.0051 | 0.0026 | 0.05333 | --- | 0.9884 |
| Monocytes | rs515362 | 5 | 76061924 | t | C | 0.5102 | $9.00 \mathrm{E}-04$ | 0.0024 | 0.6994 | +-+ | 0.6515 |
| Monocytes | rs6889169 | 5 | 76063338 | t | g | 0.9226 | -0.0059 | 0.0059 | 0.3212 | ??- | 1 |
| Monocytes | rs1639114 | 6 | 31219224 | a | c | 0.0361 | 0.0066 | 0.0072 | 3.63E-01 | ?++ | $2.58 \mathrm{E}-01$ |
| Monocytes | rs1793891 | 6 | 31221698 | t | C | 0.8454 | 0.0062 | 0.0044 | $1.58 \mathrm{E}-01$ | -?+ | $6.40 \mathrm{E}-01$ |
| Monocytes | rs1986997 | 6 | 31228410 | t | c | 0.6391 | -4.00E-04 | 0.0032 | 8.93E-01 | +-- | 3.81E-06 |
| Monocytes | rs2245822 | 6 | 31230800 | a | g | 0.1749 | -0.0116 | 0.0035 | 8.18E-04 | --- | 1.67E-01 |
| Monocytes | rs2394950 | 6 | 31230701 | a | g | 0.0809 | 0.0035 | 0.005 | 4.81E-01 | ?+- | 5.41E-02 |
| Monocytes | rs2394951 | 6 | 31230759 | a | g | 0.1812 | 0.0099 | 0.0031 | $1.46 \mathrm{E}-03$ | +++ | $4.06 \mathrm{E}-03$ |
| Monocytes | rs2394952 | 6 | 31230882 | a | g | 0.6833 | -0.0092 | 0.0031 | 3.56E-03 | --- | 5.25E-03 |
| Monocytes | rs2524117 | 6 | 31230869 | t | c | 0.3878 | 9.00E-04 | 0.0027 | 7.33E-01 | -+- | $1.93 \mathrm{E}-03$ |
| Monocytes | rs2524118 | 6 | 31230667 | a | t | 0.0849 | 0.0045 | 0.0054 | $4.09 \mathrm{E}-01$ | -?+ | $8.68 \mathrm{E}-01$ |
| Monocytes | rs2524121 | 6 | 31228972 | a | g | 0.11 | 0.0156 | 0.0038 | $3.49 \mathrm{E}-05$ | +++ | 7.74E-04 |
| Monocytes | rs2844626 | 6 | 31229552 | a | t | 0.57 | 0.007 | 0.0025 | 5.26E-03 | -++ | 6.72E-02 |
| Monocytes | rs2844627 | 6 | 31229462 | t | c | 0.6571 | -0.0035 | 0.0034 | $2.99 \mathrm{E}-01$ | --+ | 1.17E-03 |
| Monocytes | rs2894189 | 6 | 31217815 | a | c | 0.4675 | 0.0066 | 0.0024 | 5.14E-03 | +++ | 3.10E-02 |
| Monocytes | rs2894196 | 6 | 31230111 | a | t | 0.0756 | -0.0062 | 0.0053 | $2.44 \mathrm{E}-01$ | --- | 8.14E-01 |

Table S1. Random-effects meta-analysis of MANTRA identified loci.

| Monocytes | rs3095241 | 6 | 31215635 | a | c | 0.3707 | 0.008 | 0.0027 | 2.47E-03 | +++ | 3.12E-02 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| Monocytes | rs3095254 | 6 | 31221668 | c | $g$ | 0.5019 | 0.0091 | 0.0024 | 1.89E-04 | +++ | 2.42E-05 |
| Monocytes | rs3130408 | 6 | 31213791 | a | g | 0.7627 | -0.0073 | 0.0043 | 9.33E-02 | +-- | 2.62E-02 |
| Monocytes | rs3130409 | 6 | 31213811 | a | g | 0.366 | -0.0089 | 0.0028 | $1.66 \mathrm{E}-03$ | -?- | 2.45E-03 |
| Monocytes | rs3130433 | 6 | 31219638 | a | g | 0.1028 | 0.0075 | 0.0044 | 9.26E-02 | -++ | $2.64 \mathrm{E}-02$ |
| Monocytes | rs3130441 | 6 | 31228565 | a | g | 0.1039 | 0.0156 | 0.0039 | 5.48E-05 | +++ | $1.88 \mathrm{E}-04$ |
| Monocytes | rs3130444 | 6 | 31228634 | t | c | 0.2079 | -0.0044 | 0.0041 | 2.85E-01 | +?- | $5.22 \mathrm{E}-02$ |
| Monocytes | rs3899471 | 6 | 31215037 | t | C | 0.2635 | -0.001 | 0.0031 | 7.55E-01 | --+ | $6.02 \mathrm{E}-01$ |
| Monocytes | rs4084090 | 6 | 31218835 | a | g | 0.7796 | -0.0064 | 0.0043 | $1.34 \mathrm{E}-01$ | +-- | 4.12E-02 |
| Monocytes | rs4084262 | 6 | 31218889 | t | c | 0.6501 | -3.00E-04 | 0.0033 | $9.29 \mathrm{E}-01$ | ?-- | $9.25 \mathrm{E}-01$ |
| Monocytes | rs4332019 | 6 | 31213413 | t | g | 0.2873 | -0.0013 | 0.0031 | 6.62E-01 | --+ | $5.58 \mathrm{E}-01$ |
| Monocytes | rs4416711 | 6 | 31221039 | t | C | 0.7447 | 0.003 | 0.0042 | 4.83E-01 | -+- | 9.50E-03 |
| Monocytes | rs7747738 | 6 | 31230924 | t | C | 0.8998 | -0.0026 | 0.0051 | 6.13E-01 | ??- | $1.00 \mathrm{E}+00$ |
| Monocytes | rs9264180 | 6 | 31219902 | a | c | 0.5862 | -0.0061 | 0.0026 | 2.08E-02 | --- | $1.62 \mathrm{E}-01$ |
| Monocytes | rs9264374 | 6 | 31228666 | t | C | 0.2297 | 0.0026 | 0.0031 | 3.87E-01 | ?-+ | 5.22E-01 |
| Monocytes | rs9264386 | 6 | 31229130 | C | g | 0.6804 | -0.0104 | 0.003 | 6.01E-04 | --- | $6.36 \mathrm{E}-03$ |
| Monocytes | rs9264391 | 6 | 31229203 | a | g | 0.0369 | 0.0116 | 0.0079 | $1.40 \mathrm{E}-01$ | -?+ | $6.14 \mathrm{E}-01$ |
| Monocytes | rs9264409 | 6 | 31229738 | a | g | 0.1741 | 0.0097 | 0.0031 | 2.13E-03 | +++ | 9.08E-03 |
| Monocytes | rs9264416 | 6 | 31230042 | a | g | 0.668 | -0.01 | 0.0039 | $1.06 \mathrm{E}-02$ | -+- | $1.53 \mathrm{E}-02$ |
| Monocytes | rs9368665 | 6 | 31229248 | a | g | 0.6998 | 0.0039 | 0.0052 | $4.45 \mathrm{E}-01$ | +++ | $7.44 \mathrm{E}-01$ |
| Monocytes | rs9368666 | 6 | 31229644 | a | g | 0.7238 | 0.0045 | 0.0052 | 3.87E-01 | ++- | 6.19E-01 |
| Monocytes | rs9468908 | 6 | 31216227 | a | g | 0.0685 | -0.0049 | 0.0057 | $3.90 \mathrm{E}-01$ | ?+- | 8.11E-01 |
| Monocytes | rs9501364 | 6 | 31217026 | a | g | 0.0872 | 0.0095 | 0.0046 | 3.80E-02 | ++- | $1.08 \mathrm{E}-04$ |
| Monocytes | rs9501543 | 6 | 31216956 | t | C | 0.0474 | -0.0057 | 0.0069 | 4.07E-01 | +-- | 1.16E-02 |
| Monocytes | rs9501548 | 6 | 31228552 | a | g | 0.0812 | 0.0041 | 0.0049 | $4.03 \mathrm{E}-01$ | ?++ | 6.94E-02 |
| Monocytes | rs10092988 | 8 | 130624661 | a | g | 0.3854 | -0.0098 | 0.0028 | 4.81E-04 | +-- | 2.02E-04 |
| Monocytes | rs10100356 | 8 | 130626164 | a | g | 0.229 | -0.0074 | 0.0031 | 1.52E-02 | --- | $9.51 \mathrm{E}-01$ |
| Monocytes | rs10956485 | 8 | 130620585 | t | C | 0.3076 | -0.0141 | 0.0029 | $1.07 \mathrm{E}-06$ | --- | 6.60E-06 |
| Monocytes | rs12550563 | 8 | 130629300 | a | g | 0.2936 | -0.0086 | 0.0027 | $1.57 \mathrm{E}-03$ | --- | $6.24 \mathrm{E}-03$ |
| Monocytes | rs12681229 | 8 | 130623784 | t | C | 0.2914 | -0.0173 | 0.0036 | 1.87E-06 | --+ | 4.65E-04 |
| Monocytes | rs13253638 | 8 | 130623190 | t | C | 0.4147 | -0.0221 | 0.0086 | 1.03E-02 | ?-- | 8.29E-01 |
| Monocytes | rs13263823 | 8 | 130631510 | a | c | 0.4878 | -0.004 | 0.0025 | $1.10 \mathrm{E}-01$ | --+ | $1.46 \mathrm{E}-02$ |
| Monocytes | rs13265608 | 8 | 130624121 | t | C | 0.0989 | 0.0181 | 0.0058 | $1.81 \mathrm{E}-03$ | -++ | 1.03E-01 |
| Monocytes | rs1433585 | 8 | 130626097 | a | g | 0.1356 | 0.0135 | 0.0041 | $9.94 \mathrm{E}-04$ | ++- | 3.32E-03 |
| Monocytes | rs1469737 | 8 | 130626936 | a | g | 0.0209 | -0.0094 | 0.0138 | $4.96 \mathrm{E}-01$ | ??- | $1.00 \mathrm{E}+00$ |
| Monocytes | rs1583333 | 8 | 130614899 | a | C | 0.2211 | 0.0173 | 0.0037 | 2.66E-06 | +++ | 1.89E-03 |
| Monocytes | rs16893247 | 8 | 130625884 | t | c | 0.4523 | 0.0085 | 0.0042 | $4.28 \mathrm{E}-02$ | +-+ | 2.21E-02 |
| Monocytes | rs16904126 | 8 | 130627979 | a | g | 0.606 | -0.0131 | 0.0041 | $1.38 \mathrm{E}-03$ | --+ | 2.16E-03 |

Table S1. Random-effects meta-analysis of MANTRA identified loci.

| Monocytes | rs16904127 | 8 | 130628606 | a | g | 0.264 | -0.0079 | 0.0031 | 1.18E-02 | --- | 7.43E-01 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| Monocytes | rs17180246 | 8 | 130623468 | t | c | 0.6634 | 0.0079 | 0.0052 | $1.28 \mathrm{E}-01$ | ?++ | $1.28 \mathrm{E}-01$ |
| Monocytes | rs17180387 | 8 | 130624988 | t | C | 0.0729 | -0.0063 | 0.0059 | 2.91E-01 | ?-+ | 4.77E-02 |
| Monocytes | rs1982094 | 8 | 130624816 | t | c | 0.5272 | -0.0173 | 0.0049 | 3.87E-04 | --+ | 4.34E-04 |
| Monocytes | rs1991866 | 8 | 130624105 | c | $g$ | 0.5138 | -0.0156 | 0.0024 | 1.21E-10 | --- | 1.41E-07 |
| Monocytes | rs7005206 | 8 | 130620813 | a | g | 0.1894 | 0.0073 | 0.0036 | 3.98E-02 | -++ | 4.86E-01 |
| Monocytes | rs7814618 | 8 | 130627555 | t | c | 0.5916 | 0.0066 | 0.0028 | 1.83E-02 | +++ | 8.84E-03 |
| Monocytes | rs7819128 | 8 | 130630157 | t | c | 0.3757 | -0.0067 | 0.0028 | $1.52 \mathrm{E}-02$ | --- | 1.11E-02 |
| Monocytes | rs7823123 | 8 | 130628808 | t | C | 0.0262 | -0.0047 | 0.0117 | $6.88 \mathrm{E}-01$ | ??- | $1.00 \mathrm{E}+00$ |
| Monocytes | rs7825384 | 8 | 130626241 | a | C | 0.8726 | -0.0053 | 0.0045 | 2.43E-01 | ??- | $1.00 \mathrm{E}+00$ |
| Monocytes | rs7828331 | 8 | 130631930 | a | g | 0.425 | 0.0051 | 0.0027 | 5.61E-02 | 0 | 7.49E-03 |
| Monocytes | rs7830620 | 8 | 130627508 | a | C | 0.7442 | -0.0043 | 0.0035 | 2.23E-01 | ??- | $1.00 \mathrm{E}+00$ |
| Monocytes | rs7834911 | 8 | 130628036 | t | C | 0.3566 | -0.0078 | 0.0026 | 3.05E-03 | --- | 3.55E-03 |
| Monocytes | rs10116508 | 9 | 113913415 | t | c | 0.1843 | 0.0075 | 0.0057 | $1.91 \mathrm{E}-01$ | -?+ | 1.73E-01 |
| Monocytes | rs10126017 | 9 | 113918933 | a | g | 0.4515 | -6.00E-04 | 0.0026 | 8.22E-01 | --+ | 5.87E-01 |
| Monocytes | rs10217127 | 9 | 113919469 | t | g | 0.6003 | -0.0061 | 0.003 | 4.03E-02 | +-+ | 4.10E-03 |
| Monocytes | rs10217157 | 9 | 113919523 | t | c | 0.4802 | -0.0038 | 0.0026 | 1.49E-01 | --- | 9.52E-01 |
| Monocytes | rs10817156 | 9 | 113908195 | c | g | 0.4683 | -0.0011 | 0.0024 | $6.44 \mathrm{E}-01$ | +-+ | 2.42E-01 |
| Monocytes | rs10980798 | 9 | 113912649 | t | C | 0.1948 | -0.0042 | 0.0031 | 1.86E-01 | --- | $9.58 \mathrm{E}-01$ |
| Monocytes | rs10980800 | 9 | 113915905 | $t$ | c | 0.2554 | -0.0093 | 0.003 | 2.23E-03 | --+ | 7.40E-12 |
| Monocytes | rs10980802 | 9 | 113918856 | a | g | 0.7128 | 0.0122 | 0.0029 | 2.37E-05 | ++- | 1.53E-06 |
| Monocytes | rs10980805 | 9 | 113920552 | a | C | 0.4128 | 0.002 | 0.0026 | 4.35E-01 | --+ | 2.56E-01 |
| Monocytes | rs12342967 | 9 | 113924902 | t | c | 0.5967 | -0.0119 | 0.0036 | $1.02 \mathrm{E}-03$ | --- | 2.83E-02 |
| Monocytes | rs12345546 | 9 | 113919677 | a | c | 0.3243 | 5.00E-04 | 0.0028 | 8.43E-01 | ++- | 6.14E-01 |
| Monocytes | rs12346772 | 9 | 113920599 | a | g | 0.6638 | -0.0226 | 0.0035 | 1.11E-10 | --- | 8.14E-06 |
| Monocytes | rs12350763 | 9 | 113923723 | a | g | 0.6447 | -0.0166 | 0.0032 | 1.42E-07 | --- | $5.54 \mathrm{E}-08$ |
| Monocytes | rs1330280 | 9 | 113908312 | t | c | 0.8854 | 0.0033 | 0.0076 | $6.64 \mathrm{E}-01$ | ?++ | 3.70E-01 |
| Monocytes | rs1360284 | 9 | 113921689 | t | g | 0.2286 | -0.0028 | 0.0029 | $3.24 \mathrm{E}-01$ | --- | 9.06E-01 |
| Monocytes | rs16915755 | 9 | 113916135 | t | c | 0.1795 | -0.0031 | 0.004 | 4.42E-01 | ??- | $1.00 \mathrm{E}+00$ |
| Monocytes | rs16915757 | 9 | 113918391 | C | g | 0.8825 | 0.0047 | 0.0049 | 3.41E-01 | ??+ | $1.00 \mathrm{E}+00$ |
| Monocytes | rs16915769 | 9 | 113923169 | t | c | 0.1178 | -0.004 | 0.0048 | 4.09E-01 | ??- | $1.00 \mathrm{E}+00$ |
| Monocytes | rs17811749 | 9 | 113916949 | a | t | 0.5784 | 0.0165 | 0.0047 | 4.76E-04 | +++ | 2.81E-01 |
| Monocytes | rs2418136 | 9 | 113919300 | a | t | 0.5811 | 0.0162 | 0.0047 | 5.66E-04 | +++ | 2.83E-01 |
| Monocytes | rs4541999 | 9 | 113907759 | a | g | 0.472 | -0.0011 | 0.0024 | 6.55E-01 | +-+ | 2.58E-01 |
| Monocytes | rs6477805 | 9 | 113920108 | a | g | 0.3302 | 0.001 | 0.0028 | 7.22E-01 | ++- | 7.57E-01 |
| Monocytes | rs6477806 | 9 | 113920161 | t | C | 0.662 | 0.0023 | 0.0028 | 4.18E-01 | +++ | 8.45E-01 |
| Monocytes | rs7023923 | 9 | 113925534 | t | c | 0.5666 | 0.0082 | 0.0025 | 1.15E-03 | ++- | 2.08E-08 |
| Monocytes | rs7025176 | 9 | 113920511 | t | C | 0.0638 | -0.0076 | 0.0067 | 2.53E-01 | ??- | $1.00 \mathrm{E}+00$ |

Table S1. Random-effects meta-analysis of MANTRA identified loci.

| Monocytes | rs7034139 | 9 | 113924517 | a | c | 0.2196 | 0.0164 | 0.0032 | 1.99E-07 | +++ | 3.94E-08 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| Monocytes | rs7849209 | 9 | 113915676 | a | g | 0.8211 | 0.0057 | 0.0042 | $1.78 \mathrm{E}-01$ | ??+ | $1.00 \mathrm{E}+00$ |
| Monocytes | rs7867745 | 9 | 113908101 | a | g | 0.5084 | 0.0166 | 0.0054 | $1.96 \mathrm{E}-03$ | ++- | 5.55E-02 |
| Monocytes | rs7870066 | 9 | 113911613 | t | c | 0.6929 | 0.0037 | 0.003 | 2.10E-01 | +++ | 8.61E-01 |
| Monocytes | rs9299192 | 9 | 113918647 | a | t | 0.3278 | $7.00 \mathrm{E}-04$ | 0.0028 | 7.98E-01 | ++- | 5.70E-01 |
| Monocytes | rs1371798 | 4 | 74976781 | t | C | 0.4721 | -0.0095 | 0.0031 | $2.14 \mathrm{E}-03$ | --+ | 2.78E-09 |
| Neutrophils | rs1371799 | 4 | 74977837 | t | c | 0.4367 | 0.0088 | 0.0031 | 4.66E-03 | ++- | $1.11 \mathrm{E}-10$ |
| Neutrophils | rs1837559 | 4 | 74959093 | a | g | 0.6052 | -0.0159 | 0.0047 | 6.64E-04 | -?- | 3.55E-01 |
| Neutrophils | rs1893319 | 4 | 74973129 | t | c | 0.5039 | 0.0152 | 0.0049 | $1.76 \mathrm{E}-03$ | +?- | 5.59E-02 |
| Neutrophils | rs3806792 | 4 | 74965274 | t | c | 0.4472 | 0.0068 | 0.0033 | 3.95E-02 | ++- | 6.10E-08 |
| Neutrophils | rs505197 | 4 | 74968560 | t | C | 0.9414 | 0.006 | 0.0189 | 7.53E-01 | +?+ | 7.12E-01 |
| Neutrophils | rs549280 | 4 | 74971196 | a | g | 0.5094 | -0.0204 | 0.0042 | 9.52E-07 | -?- | 5.14E-01 |
| Neutrophils | rs7667376 | 4 | 74967890 | $t$ | c | 0.4765 | -0.0084 | 0.0031 | 6.04E-03 | --+ | 1.02E-10 |
| Neutrophils | rs9131 | 4 | 74963049 | t | c | 0.4904 | -0.006 | 0.0033 | $6.92 \mathrm{E}-02$ | --+ | 1.62E-08 |
| Neutrophils | rs10263804 | 7 | 92416590 | a | g | 0.877 | $5.00 \mathrm{E}-04$ | 0.0108 | 9.63E-01 | ??+ | $1.00 \mathrm{E}+00$ |
| Neutrophils | rs2282995 | 7 | 92411623 | a | g | 0.6026 | -0.0129 | 0.0045 | 4.43E-03 | --- | $8.34 \mathrm{E}-01$ |
| Neutrophils | rs3731303 | 7 | 92403859 | t | c | 0.3011 | 0.011 | 0.0045 | $1.52 \mathrm{E}-02$ | ++- | 2.75E-01 |
| Neutrophils | rs3731304 | 7 | 92403660 | a | c | 0.0161 | 0.0541 | 0.0309 | 7.96E-02 | ??+ | $1.00 \mathrm{E}+00$ |
| Neutrophils | rs3802072 | 7 | 92403053 | a | g | 0.1835 | 0.0152 | 0.0041 | 1.87E-04 | +++ | 7.54E-01 |
| Neutrophils | rs3802073 | 7 | 92402480 | t | c | 0.3564 | 0.0052 | 0.0041 | 2.05E-01 | ++- | $1.01 \mathrm{E}-02$ |
| Neutrophils | rs41461052 | 7 | 92407600 | t | c | 0.0564 | 0.0245 | 0.0158 | $1.21 \mathrm{E}-01$ | ??+ | $1.00 \mathrm{E}+00$ |
| Neutrophils | rs445 | 7 | 92408370 | $t$ | c | 0.3892 | -0.0093 | 0.0041 | 2.44E-02 | --+ | 8.29E-11 |
| Neutrophils | rs8 | 7 | 92408329 | t | c | 0.3711 | 0.002 | 0.0061 | 7.44E-01 | ?+- | 6.30E-01 |
| Neutrophils | rs10223710 | 6 | 32218681 | a | g | 0.5313 | -0.0066 | 0.0035 | 5.85E-02 | --- | 4.40E-01 |
| Neutrophils | rs10456405 | 6 | 32212867 | t | g | 0.1217 | -0.0076 | 0.0062 | 2.19E-01 | ?-- | 9.32E-01 |
| Neutrophils | rs12525893 | 6 | 32220577 | t | c | 0.3564 | 0.0017 | 0.0037 | 6.53E-01 | ++- | 8.87E-02 |
| Neutrophils | rs13194642 | 6 | 32211695 | t | C | 0.3186 | 0.0039 | 0.0045 | $3.89 \mathrm{E}-01$ | -++ | 3.39E-01 |
| Neutrophils | rs1559876 | 6 | 32215769 | C | g | 0.3299 | 0.0062 | 0.0032 | 5.55E-02 | ++- | 7.10E-02 |
| Neutrophils | rs17576984 | 6 | 32212985 | t | c | 0.27 | -0.0114 | 0.0046 | $1.28 \mathrm{E}-02$ | --- | 1.47E-01 |
| Neutrophils | rs2395110 | 6 | 32215876 | a | g | 0.5005 | -9.00E-04 | 0.0041 | 8.23E-01 | +?- | 1.83E-03 |
| Neutrophils | rs2395111 | 6 | 32215964 | t | c | 0.4188 | -0.0124 | 0.003 | 2.85E-05 | --- | 4.57E-01 |
| Neutrophils | rs2515892 | 6 | 32215252 | t | g | 0.5302 | 0.0127 | 0.0029 | $1.62 \mathrm{E}-05$ | +++ | 4.26E-01 |
| Neutrophils | rs2894239 | 6 | 32215796 | t | g | 0.4538 | -0.0129 | 0.0031 | 2.25E-05 | --- | 5.34E-01 |
| Neutrophils | rs2894240 | 6 | 32217846 | a | g | 0.4249 | 0.0063 | 0.003 | 3.17E-02 | ++- | 4.23E-04 |
| Neutrophils | rs3096700 | 6 | 32221782 | a | c | 0.3014 | -0.0067 | 0.0046 | 1.52E-01 | -?+ | 9.48E-03 |
| Neutrophils | rs3115571 | 6 | 32220918 | a | g | 0.3473 | -0.0121 | 0.0032 | $1.68 \mathrm{E}-04$ | --+ | $2.74 \mathrm{E}-03$ |
| Neutrophils | rs3115572 | 6 | 32220484 | c | g | 0.3515 | -0.0125 | 0.0032 | 7.21E-05 | --+ | 4.39E-03 |
| Neutrophils | rs3115573 | 6 | 32218843 | a | g | 0.5346 | -0.0039 | 0.0032 | 2.16E-01 | --+ | 1.61E-03 |

Table S1. Random-effects meta-analysis of MANTRA identified loci.

| Neutrophils | rs3115575 | 6 | 32216891 | t | g | 0.4655 | -0.013 | 0.003 | 1.11E-05 | --- | 4.64E-01 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| Neutrophils | rs3115576 | 6 | 32216850 | a | t | 0.426 | 0.0063 | 0.003 | 3.33E-02 | ++- | 4.52E-04 |
| Neutrophils | rs3130304 | 6 | 32207181 | a | g | 0.186 | -0.0112 | 0.004 | 4.82E-03 | --- | $7.30 \mathrm{E}-01$ |
| Neutrophils | rs3130309 | 6 | 32215472 | a | g | 0.3463 | -0.0125 | 0.0032 | 8.62E-05 | --+ | 3.82E-03 |
| Neutrophils | rs3130310 | 6 | 32216911 | C | g | 0.524 | -0.0063 | 0.003 | 3.26E-02 | --+ | 5.28E-04 |
| Neutrophils | rs3130311 | 6 | 32217367 | a | g | 0.5235 | -0.0062 | 0.003 | 3.55E-02 | --+ | 3.41E-04 |
| Neutrophils | rs3130315 | 6 | 32220685 | a | g | 0.4244 | 0.0041 | 0.0032 | $2.00 \mathrm{E}-01$ | ++- | 1.97E-03 |
| Neutrophils | rs3130316 | 6 | 32221228 | t | C | 0.4034 | -0.0153 | 0.0032 | $1.90 \mathrm{E}-06$ | --- | 2.96E-01 |
| Neutrophils | rs3130320 | 6 | 32223258 | t | c | 0.4139 | -0.0154 | 0.0048 | $1.32 \mathrm{E}-03$ | -?- | 3.51E-01 |
| Neutrophils | rs371156 | 6 | 32209963 | t | c | 0.431 | -0.0074 | 0.0032 | 2.08E-02 | --- | 6.95E-01 |
| Neutrophils | rs382259 | 6 | 32209027 | t | C | 0.438 | -0.0082 | 0.0034 | $1.46 \mathrm{E}-02$ | --- | $6.08 \mathrm{E}-01$ |
| Neutrophils | rs405875 | 6 | 32215188 | t | C | 0.4207 | -0.0125 | 0.003 | 2.36E-05 | --- | 5.05E-01 |
| Neutrophils | rs410283 | 6 | 32215198 | t | g | 0.5295 | 0.0126 | 0.003 | $1.90 \mathrm{E}-05$ | +++ | 4.50E-01 |
| Neutrophils | rs411326 | 6 | 32211317 | t | c | 0.4329 | -0.0044 | 0.0034 | 1.99E-01 | -+- | 1.67E-03 |
| Neutrophils | rs412492 | 6 | 32213831 | a | t | 0.4785 | -0.006 | 0.003 | 4.38E-02 | --+ | 3.90E-04 |
| Neutrophils | rs412657 | 6 | 32211085 | t | g | 0.4331 | 0.0058 | 0.0031 | 6.47E-02 | ++- | 6.92E-01 |
| Neutrophils | rs416352 | 6 | 32207393 | t | g | 0.4314 | 0.0017 | 0.0031 | 5.78E-01 | +-+ | 4.15E-03 |
| Neutrophils | rs419132 | 6 | 32210799 | a | g | 0.4703 | -0.0025 | 0.0032 | 4.29E-01 | --+ | 1.17E-01 |
| Neutrophils | rs424232 | 6 | 32208324 | t | C | 0.3828 | -7.00E-04 | 0.0033 | 8.43E-01 | +-+ | 1.30E-01 |
| Neutrophils | rs427037 | 6 | 32212264 | a | g | 0.5252 | -0.0021 | 0.0033 | 5.22E-01 | --+ | 3.73E-02 |
| Neutrophils | rs440169 | 6 | 32213788 | a | g | 0.4294 | 0.0066 | 0.003 | $2.88 \mathrm{E}-02$ | ++- | 1.19E-04 |
| Neutrophils | rs454748 | 6 | 32213210 | a | g | 0.5132 | 0.0074 | 0.003 | 1.33E-02 | ++- | $1.44 \mathrm{E}-03$ |
| Neutrophils | rs454875 | 6 | 32213008 | a | g | 0.1384 | 0.0059 | 0.005 | 2.39E-01 | ++- | $3.04 \mathrm{E}-01$ |
| Neutrophils | rs4959090 | 6 | 32219962 | a | g | 0.2316 | 0.0047 | 0.0037 | $1.98 \mathrm{E}-01$ | -++ | $3.58 \mathrm{E}-01$ |
| Neutrophils | rs507778 | 6 | 32209861 | t | C | 0.423 | 0.0012 | 0.0031 | $6.99 \mathrm{E}-01$ | +-+ | $3.38 \mathrm{E}-02$ |
| Neutrophils | rs563412 | 6 | 32215063 | a | g | 0.4711 | 0.006 | 0.0029 | 4.22E-02 | ++- | 4.22E-04 |
| Neutrophils | rs6457508 | 6 | 32216963 | t | c | 0.4857 | 0.0129 | 0.003 | 1.32E-05 | +++ | 5.33E-01 |
| Neutrophils | rs6457509 | 6 | 32217018 | C | g | 0.4247 | 0.0061 | 0.003 | 3.80E-02 | ++- | $3.88 \mathrm{E}-04$ |
| Neutrophils | rs6457510 | 6 | 32217046 | t | c | 0.4854 | 0.0128 | 0.003 | 1.52E-05 | +++ | 5.35E-01 |
| Neutrophils | rs6457515 | 6 | 32223785 | C | g | 0.2382 | -0.0095 | 0.0035 | 6.52E-03 | --+ | 5.07E-03 |
| Neutrophils | rs6457516 | 6 | 32223830 | a | t | 0.5404 | 0.0101 | 0.0035 | 3.93E-03 | ++- | $1.28 \mathrm{E}-02$ |
| Neutrophils | rs6908927 | 6 | 32224489 | a | g | 0.2385 | -0.01 | 0.0035 | 4.11E-03 | --+ | $1.30 \mathrm{E}-02$ |
| Neutrophils | rs6914780 | 6 | 32216529 | a | t | 0.0756 | 0.0063 | 0.0068 | 3.52E-01 | -++ | 9.14E-01 |
| Neutrophils | rs6915612 | 6 | 32218625 | a | g | 0.0628 | 0.0116 | 0.0145 | $4.22 \mathrm{E}-01$ | ??+ | $1.00 \mathrm{E}+00$ |
| Neutrophils | rs6916062 | 6 | 32219041 | t | c | 0.4791 | 0.0033 | 0.0086 | 7.01E-01 | ?+- | 3.03E-01 |
| Neutrophils | rs6921945 | 6 | 32220037 | a | C | 0.5291 | -0.0062 | 0.0035 | 7.55E-02 | --- | 5.29E-01 |
| Neutrophils | rs6936204 | 6 | 32217092 | $t$ | c | 0.4321 | -0.0196 | 0.0035 | 2.72E-08 | --- | 1.85E-01 |
| Neutrophils | rs9267947 | 6 | 32211218 | a | g | 0.3727 | 0.0013 | 0.0032 | 6.78E-01 | ++- | 2.08E-02 |

Table S1. Random-effects meta-analysis of MANTRA identified loci.

| Neutrophils | rs9267948 | 6 | 32212233 | a | g | 0.6607 | -0.0015 | 0.0034 | $6.61 \mathrm{E}-01$ | --+ | 2.82E-01 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| Neutrophils | rs9267954 | 6 | 32213052 | a | t | 0.826 | -0.0059 | 0.0055 | $2.84 \mathrm{E}-01$ | -?- | $7.49 \mathrm{E}-01$ |
| Neutrophils | rs9267955 | 6 | 32213150 | a | g | 0.0973 | 0.0033 | 0.012 | 7.84E-01 | ??+ | $1.00 \mathrm{E}+00$ |
| Neutrophils | rs9267956 | 6 | 32213638 | a | g | 0.6011 | -0.0039 | 0.0035 | 2.61E-01 | --+ | $9.66 \mathrm{E}-02$ |
| Neutrophils | rs9267958 | 6 | 32214616 | a | c | 0.5235 | -0.0062 | 0.003 | 3.67E-02 | --+ | 2.85E-04 |
| Neutrophils | rs9267970 | 6 | 32217160 | a | g | 0.5211 | -0.0059 | 0.003 | 4.45E-02 | --+ | 3.12E-04 |
| Neutrophils | rs9267971 | 6 | 32217185 | t | c | 0.4639 | -0.013 | 0.003 | 1.13E-05 | --- | $4.42 \mathrm{E}-01$ |
| Neutrophils | rs9267992 | 6 | 32220397 | a | g | 0.9256 | -0.0033 | 0.0127 | 7.95E-01 | ??- | $1.00 \mathrm{E}+00$ |
| Neutrophils | rs9268000 | 6 | 32223531 | a | C | 0.5759 | -0.0073 | 0.0033 | $2.56 \mathrm{E}-02$ | --+ | 7.84E-02 |
| Neutrophils | rs9296013 | 6 | 32216147 | a | c | 0.686 | 0.0098 | 0.0059 | 9.56E-02 | +-- | 4.43E-03 |
| Neutrophils | rs9296015 | 6 | 32218989 | a | g | 0.2758 | 0.0085 | 0.0036 | 1.97E-02 | +++ | $2.37 \mathrm{E}-01$ |
| Neutrophils | rs9296016 | 6 | 32219010 | a | t | 0.2668 | 0.0063 | 0.0035 | 7.07E-02 | +++ | $5.09 \mathrm{E}-01$ |
| Neutrophils | rs9357138 | 6 | 32219838 | t | C | 0.3014 | 0.0045 | 0.0046 | 3.29E-01 | -++ | $3.08 \mathrm{E}-01$ |
| Neutrophils | rs8078723 | 17 | 38166879 | $t$ | c | 0.4046 | -0.0307 | 3.00E-03 | 5.02E-24 | --+ | 4.23E-08 |
| Neutrophils | rs4794822 | 17 | 38156712 | $t$ | c | 0.496 | 3.12E-02 | 3.00E-03 | 8.55E-25 | ++- | 2.05E-07 |
| WBC count | rs16841659 | 1 | 159053650 | c | g | 0.9141 | 0.0122 | 0.004 | 2.33E-03 | -?+ | 2.62E-07 |
| WBC count | rs16841682 | 1 | 159064818 | a | C | 0.6964 | 0.0017 | 0.0029 | 5.61E-01 | +-+ | $4.63 \mathrm{E}-01$ |
| WBC count | rs1894043 | 1 | 159069211 | t | C | 0.2701 | 0.0166 | 0.0022 | 6.13E-14 | +-+ | $5.79 \mathrm{E}-166$ |
| WBC count | rs2078724 | 1 | 159072108 | t | C | 0.5111 | 0.0334 | 0.0032 | $2.84 \mathrm{E}-25$ | ?-- | $4.32 \mathrm{E}-18$ |
| WBC count | rs2518561 | 1 | 159055124 | t | C | 0.5007 | 0.0343 | 0.0031 | 1.37E-28 | ?-- | $9.41 \mathrm{E}-18$ |
| WBC count | rs2518562 | 1 | 159055656 | t | C | 0.5229 | -0.02 | 0.0022 | 2.42E-20 | ++- | $1.45 \mathrm{E}-40$ |
| WBC count | rs2518564 | 1 | 159062436 | $a$ | $g$ | 0.4639 | 0.0365 | 0.0021 | 6.92E-69 | +++ | 6.98E-258 |
| WBC count | rs2518565 | 1 | 159070113 | a | g | 0.3318 | -0.0236 | 0.0021 | 1.46E-29 | +-- | $1.46 \mathrm{E}-58$ |
| WBC count | rs2814762 | 1 | 159068326 | a | g | 0.6498 | 0.053 | 0.0039 | 2.60E-42 | ?++ | $1.64 \mathrm{E}-04$ |
| WBC count | rs2814764 | 1 | 159064568 | a | t | 0.354 | -0.0273 | 0.0021 | 2.12E-39 | --- | 3.17E-79 |
| WBC count | rs2852720 | 1 | 159060560 | t | C | 0.3129 | 0.038 | 0.0027 | 4.16E-45 | +-+ | 2.71E-41 |
| WBC count | rs2852721 | 1 | 159068064 | t | C | 0.6737 | 0.0472 | 0.0044 | 4.39E-27 | --+ | $4.83 \mathrm{E}-08$ |
| WBC count | rs855866 | 1 | 159052720 | a | g | 0.3516 | -0.0348 | 0.0021 | 6.43E-64 | --- | $1.98 \mathrm{E}-203$ |
| WBC count | rs855867 | 1 | 159052847 | a | g | 0.4913 | 0.0276 | 0.0021 | 2.27E-39 | +++ | 2.73E-83 |
| WBC count | rs1371798 | 4 | 74976781 | t | c | 0.4882 | -0.0058 | 0.0018 | $1.56 \mathrm{E}-03$ | --+ | $6.75 \mathrm{E}-11$ |
| WBC count | rs1371799 | 4 | 74977837 | $t$ | c | 0.4361 | 0.0052 | 0.0018 | 4.28E-03 | ++- | 1.09E-12 |
| WBC count | rs1866755 | 4 | 74978340 | t | C | 0.4908 | -0.0053 | 0.0018 | 3.59E-03 | --+ | $2.30 \mathrm{E}-12$ |
| WBC count | rs1893319 | 4 | 74973129 | t | C | 0.4683 | 0.0111 | 0.0019 | 9.67E-09 | +++ | $2.51 \mathrm{E}-01$ |
| WBC count | rs505197 | 4 | 74968560 | t | C | 0.9264 | -0.0037 | 0.0143 | 7.94E-01 | +?- | $6.04 \mathrm{E}-01$ |
| WBC count | rs549280 | 4 | 74971196 | a | g | 0.4733 | -0.011 | 0.0018 | 4.17E-10 | --- | $7.41 \mathrm{E}-01$ |
| WBC count | rs551055 | 4 | 74981560 | a | g | 0.9486 | -0.0086 | 0.014 | 5.39E-01 | ??- | $1.00 \mathrm{E}+00$ |
| WBC count | rs7667376 | 4 | 74967890 | t | c | 0.4918 | -0.005 | 0.0018 | 5.41E-03 | --+ | 2.92E-12 |
| WBC count | rs10484494 | 6 | 135430093 | a | g | 0.0945 | -0.0127 | 0.0053 | $1.66 \mathrm{E}-02$ | ?-+ | 1.18E-02 |

Table S1. Random-effects meta-analysis of MANTRA identified loci.

| WBC count | rs1074849 | 6 | 135423412 | a | g | 0.2266 | 0.0039 | 0.0022 | 7.07E-02 | +++ | 4.29E-01 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| WBC count | rs11154792 | 6 | 135431640 | t | c | 0.5275 | 0.0131 | 0.0026 | 5.39E-07 | +?- | 5.35E-04 |
| WBC count | rs11759553 | 6 | 135422296 | a | t | 0.5257 | 0.0108 | 0.0018 | 3.08E-09 | ++- | 1.43E-04 |
| WBC count | rs1411919 | 6 | 135432061 | a | g | 0.5694 | -0.0047 | 0.0021 | $2.39 \mathrm{E}-02$ | --- | 6.19E-01 |
| WBC count | rs2223385 | 6 | 135435171 | a | g | 0.2643 | 0.003 | 0.002 | $1.24 \mathrm{E}-01$ | +++ | 9.16E-01 |
| WBC count | rs4895440 | 6 | 135426558 | a | t | 0.5299 | 0.0107 | 0.0018 | 4.87E-09 | ++- | $5.78 \mathrm{E}-05$ |
| WBC count | rs4895441 | 6 | 135426573 | $a$ | $g$ | 0.5376 | 0.0131 | 0.002 | 2.50E-11 | +++ | 1.63E-02 |
| WBC count | rs6920211 | 6 | 135431318 | t | c | 0.4331 | 0.0126 | 0.0019 | $1.98 \mathrm{E}-11$ | +++ | $1.48 \mathrm{E}-02$ |
| WBC count | rs6930223 | 6 | 135424203 | , | g | 0.5024 | -0.0095 | 0.0018 | 1.73E-07 | --- | $6.28 \mathrm{E}-02$ |
| WBC count | rs7743042 | 6 | 135419834 | a | g | 0.3901 | 0.0058 | 0.0018 | $1.29 \mathrm{E}-03$ | ++- | $1.45 \mathrm{E}-01$ |
| WBC count | rs7743480 | 6 | 135420064 | t | C | 0.8608 | -0.0032 | 0.0082 | $6.98 \mathrm{E}-01$ | ??- | $1.00 \mathrm{E}+00$ |
| WBC count | rs7766963 | 6 | 135432883 | t | C | 0.4892 | 0.0089 | 0.0018 | $1.05 \mathrm{E}-06$ | +++ | 6.60E-01 |
| WBC count | rs7775698 | 6 | 135418635 | t | C | 0.4712 | -0.0102 | 0.0025 | 3.81E-05 | -?+ | 7.84E-04 |
| WBC count | rs7776054 | 6 | 135418916 | a | g | 0.552 | 0.012 | 0.0019 | 2.99E-10 | +++ | $8.84 \mathrm{E}-02$ |
| WBC count | rs9373124 | 6 | 135423209 | t | c | 0.4582 | 0.0113 | 0.0018 | $1.06 \mathrm{E}-09$ | ++- | 3.47E-04 |
| WBC count | rs9376092 | 6 | 135427144 | a | C | 0.305 | -0.0115 | 0.002 | 4.28E-09 | --+ | 5.63E-04 |
| WBC count | rs9389268 | 6 | 135419631 | a | g | 0.553 | 0.0115 | 0.0019 | 1.55E-09 | +++ | 2.31E-02 |
| WBC count | rs9389269 | 6 | 135427159 | t | C | 0.4243 | 0.013 | 0.002 | 5.11E-11 | ++- | 5.78E-03 |
| WBC count | rs9399137 | 6 | 135419018 | t | C | 0.4372 | 0.0116 | 0.002 | 8.29E-09 | ++- | 4.87E-04 |
| WBC count | rs9402684 | 6 | 135419305 | t | C | 0.4589 | 0.006 | 0.0018 | 8.67E-04 | ++- | 2.26E-01 |
| WBC count | rs9402686 | 6 | 135427817 | a | g | 0.3027 | -0.0138 | 0.002 | $4.45 \mathrm{E}-12$ | --- | 8.24E-02 |
| WBC count | rs9483787 | 6 | 135434609 | a | C | 0.1372 | -0.0015 | 0.0051 | 7.71E-01 | ??- | $1.00 \mathrm{E}+00$ |
| WBC count | rs9483788 | 6 | 135435501 | t | C | 0.4333 | 0.0128 | 0.0021 | 1.39E-09 | +++ | $4.18 \mathrm{E}-01$ |
| WBC count | rs9494140 | 6 | 135430029 | t | c | 0.883 | -7.00E-04 | 0.0057 | 9.02E-01 | ??- | $1.00 \mathrm{E}+00$ |
| WBC count | rs9494141 | 6 | 135430299 | t | C | 0.2024 | 0.0061 | 0.0045 | $1.78 \mathrm{E}-01$ | ??+ | $1.00 \mathrm{E}+00$ |
| WBC count | rs9494144 | 6 | 135432530 | t | C | 0.8832 | -0.0013 | 0.0056 | 8.15E-01 | ??- | $1.00 \mathrm{E}+00$ |
| WBC count | rs9494145 | 6 | 135432552 | t | C | 0.466 | 0.0124 | 0.0022 | 1.83E-08 | +++ | 2.33E-01 |
| WBC count | rs10263804 | 7 | 92416590 | a | g | 0.8781 | -0.0138 | 0.0057 | $1.48 \mathrm{E}-02$ | ??- | $1.00 \mathrm{E}+00$ |
| WBC count | rs2282995 | 7 | 92411623 | a | g | 0.6411 | -0.0064 | 0.0026 | $1.24 \mathrm{E}-02$ | --+ | $1.09 \mathrm{E}-01$ |
| WBC count | rs3731303 | 7 | 92403859 | t | C | 0.3141 | 0.0069 | 0.0026 | 7.41E-03 | +++ | $1.55 \mathrm{E}-01$ |
| WBC count | rs3731304 | 7 | 92403660 | a | C | 0.0157 | 0.0409 | 0.0235 | 8.13E-02 | ??+ | $1.00 \mathrm{E}+00$ |
| WBC count | rs3802072 | 7 | 92403053 | a | g | 0.1923 | 0.0078 | 0.0023 | 6.82E-04 | +++ | 2.15E-01 |
| WBC count | rs3802073 | 7 | 92402480 | t | c | 0.3701 | 0.0023 | 0.0023 | 3.16E-01 | ++- | 1.06E-03 |
| WBC count | rs41461052 | 7 | 92407600 | t | C | 0.0578 | 0.0086 | 0.0082 | 2.96E-01 | ??+ | $1.00 \mathrm{E}+00$ |
| WBC count | rs445 | 7 | 92408370 | $t$ | c | 0.4013 | -0.0098 | 0.0023 | 2.19E-05 | --+ | 8.92E-14 |
| WBC count | rs8 | 7 | 92408329 |  | C | 0.4051 | 0.0028 | 0.0037 | $4.55 \mathrm{E}-01$ | ?++ | $8.68 \mathrm{E}-01$ |
| WBC count | rs1006122 | 2 | 113785842 | C | g | 0.5553 | -3.00E-04 | 0.0024 | $9.04 \mathrm{E}-01$ | +-- | $5.58 \mathrm{E}-01$ |
| WBC count | rs1013477 | 2 | 113777408 | $t$ | C | 0.3527 | -0.0016 | 0.0019 | $3.78 \mathrm{E}-01$ | -+- | 7.87E-02 |

Table S1. Random-effects meta-analysis of MANTRA identified loci.

| WBC count | rs10165797 | 2 | 113828450 | a | c | 0.4294 | -0.004 | 0.0018 | $2.39 \mathrm{E}-02$ | --- | 6.89E-01 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| WBC count | rs10169599 | 2 | 113838652 | t | C | 0.3387 | 0.002 | 0.002 | 3.38E-01 | ++- | 1.27E-01 |
| WBC count | rs10176274 | 2 | 113840575 | C | g | 0.5542 | -0.0096 | 0.0021 | 7.81E-06 | --- | 5.14E-01 |
| WBC count | rs10179714 | 2 | 113805118 | a | g | 0.1941 | -0.0061 | 0.0028 | $2.97 \mathrm{E}-02$ | -?+ | 9.58E-02 |
| WBC count | rs10181720 | 2 | 113838608 | a | g | 0.4625 | -0.0044 | 0.002 | 2.98E-02 | --- | 7.17E-01 |
| WBC count | rs10184259 | 2 | 113838628 | t | c | 0.5216 | -0.0022 | 0.002 | $2.75 \mathrm{E}-01$ | --+ | 9.61E-02 |
| WBC count | rs10184277 | 2 | 113823485 | t | c | 0.2994 | -0.0076 | 0.0032 | $1.77 \mathrm{E}-02$ | -?+ | $1.38 \mathrm{E}-01$ |
| WBC count | rs10185781 | 2 | 113849150 | a | C | 0.2666 | -0.0046 | 0.0022 | 3.25E-02 | --+ | $1.56 \mathrm{E}-01$ |
| WBC count | rs10186133 | 2 | 113836944 |  | g | 0.3634 | 0.0037 | 0.0019 | 5.57E-02 | ++- | 3.67E-02 |
| WBC count | rs10188292 | 2 | 113840443 | a | t | 0.5807 | -0.0091 | 0.0021 | 1.52E-05 | --- | 1.83E-01 |
| WBC count | rs10192014 | 2 | 113922412 | a | g | 0.3613 | 0.0015 | 0.0036 | $6.67 \mathrm{E}-01$ | +?+ | $6.96 \mathrm{E}-01$ |
| WBC count | rs10199363 | 2 | 113840791 | a | g | 0.9191 | 0.0144 | 0.005 | 3.73E-03 | +?+ | $5.58 \mathrm{E}-01$ |
| WBC count | rs10205381 | 2 | 113753124 | a | g | 0.1217 | $5.00 \mathrm{E}-04$ | 0.0056 | 9.28E-01 | ??+ | $1.00 \mathrm{E}+00$ |
| WBC count | rs10206428 | 2 | 113826475 | a | g | 0.3829 | -0.0048 | 0.0018 | 7.06E-03 | --- | $2.56 \mathrm{E}-01$ |
| WBC count | rs10207930 | 2 | 113860861 | a | C | 0.4591 | 0.0041 | 0.0018 | 2.72E-02 | -++ | $6.02 \mathrm{E}-03$ |
| WBC count | rs10496447 | 2 | 113782059 | a | g | 0.5997 | 0.0019 | 0.003 | 5.31E-01 | --+ | 3.98E-02 |
| WBC count | rs10496448 | 2 | 113828690 | C | g | 0.197 | -0.0083 | 0.0029 | 4.83E-03 | -?- | 6.23E-01 |
| WBC count | rs10514809 | 2 | 113781077 | t | C | 0.022 | -0.0031 | 0.0187 | 8.69E-01 | ??- | $1.00 \mathrm{E}+00$ |
| WBC count | rs10864909 | 2 | 113750230 | a | g | 0.4359 | 0.0025 | 0.002 | 2.04E-01 | -++ | $3.01 \mathrm{E}-01$ |
| WBC count | rs10864910 | 2 | 113935222 | t | c | 0.5149 | 0 | 0.0021 | 9.87E-01 | -++ | 4.71E-01 |
| WBC count | rs11123158 | 2 | 113811057 | a | g | 0.2844 | -0.0062 | 0.0025 | $1.39 \mathrm{E}-02$ | -?- | 3.25E-01 |
| WBC count | rs11123159 | 2 | 113834030 | t | c | 0.451 | 0.0049 | 0.0024 | 3.56E-02 | ++- | $6.21 \mathrm{E}-02$ |
| WBC count | rs11123161 | 2 | 113845578 | t | C | 0.37 | -3.00E-04 | 0.0018 | 8.57E-01 | -+- | 4.59E-04 |
| WBC count | rs11123162 | 2 | 113850114 | a | g | 0.4476 | -0.0022 | 0.002 | 2.65E-01 | +-- | $3.71 \mathrm{E}-02$ |
| WBC count | rs11123167 | 2 | 113906869 | C | g | 0.3792 | 0.0055 | 0.0021 | 7.77E-03 | +++ | $6.62 \mathrm{E}-01$ |
| WBC count | rs1156701 | 2 | 113827845 | a | g | 0.09 | 0.0012 | 0.0031 | $6.94 \mathrm{E}-01$ | ++- | $1.58 \mathrm{E}-01$ |
| WBC count | rs11677140 | 2 | 113865808 | a | c | 0.5441 | 2.00E-04 | 0.0022 | 9.11E-01 | -+- | 3.03E-01 |
| WBC count | rs11677407 | 2 | 113845029 | t | c | 0.3456 | -0.0042 | 0.0022 | 5.36E-02 | --+ | 8.75E-02 |
| WBC count | rs11678375 | 2 | 113835691 | t | c | 0.5782 | -0.0039 | 0.0019 | $4.44 \mathrm{E}-02$ | --+ | 3.72E-02 |
| WBC count | rs11681884 | 2 | 113847628 | t | C | 0.3522 | -0.0044 | 0.0021 | $4.18 \mathrm{E}-02$ | --+ | $1.50 \mathrm{E}-01$ |
| WBC count | rs11684289 | 2 | 113845038 | t | g | 0.4912 | 0.0036 | 0.0021 | $9.64 \mathrm{E}-02$ | ++- | $4.58 \mathrm{E}-02$ |
| WBC count | rs11684719 | 2 | 113842401 | a | g | 0.2257 | -0.0089 | 0.0022 | 7.35E-05 | --- | $7.36 \mathrm{E}-01$ |
| WBC count | rs11687782 | 2 | 113841782 | a | t | 0.464 | 0.0036 | 0.0019 | 5.40E-02 | -++ | $6.19 \mathrm{E}-04$ |
| WBC count | rs11687786 | 2 | 113811731 | t | g | 0.6431 | 7.00E-04 | 0.0026 | 7.99E-01 | -?+ | $1.57 \mathrm{E}-01$ |
| WBC count | rs11693683 | 2 | 113850653 | a | t | 0.3944 | 0.0046 | 0.0031 | $1.46 \mathrm{E}-01$ | +++ | 6.89E-02 |
| WBC count | rs11883847 | 2 | 113792171 | t | C | 0.4891 | 0.0019 | 0.0037 | $6.01 \mathrm{E}-01$ | ?-+ | 7.66E-03 |
| WBC count | rs11885498 | 2 | 113900177 | a | g | 0.5914 | -0.0056 | 0.0021 | 7.00E-03 | --- | $8.64 \mathrm{E}-01$ |
| WBC count | rs11886754 | 2 | 113844342 | c | g | 0.5648 | 0.0056 | 0.0021 | 8.03E-03 | +++ | 3.53E-01 |

Table S1. Random-effects meta-analysis of MANTRA identified loci.

| WBC count | rs11891198 | 2 | 113844229 | a | g | 0.5745 | 0.0056 | 0.0022 | 1.04E-02 | +++ | 3.58E-01 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| WBC count | rs11893386 | 2 | 113844310 | t | c | 0.4927 | 0.0041 | 0.0022 | 5.97E-02 | ++- | 7.29E-02 |
| WBC count | rs11896207 | 2 | 113850882 | t | c | 0.7353 | -0.0111 | 0.0052 | $3.35 \mathrm{E}-02$ | ?+- | 4.49E-02 |
| WBC count | rs11897481 | 2 | 113851439 | t | C | 0.3521 | -0.0045 | 0.0021 | $3.79 \mathrm{E}-02$ | --+ | 1.63E-01 |
| WBC count | rs11898158 | 2 | 113823572 | a | g | 0.5168 | 0.0098 | 0.0035 | $4.71 \mathrm{E}-03$ | ?++ | $1.65 \mathrm{E}-01$ |
| WBC count | rs11898742 | 2 | 113845392 | a | g | 0.0721 | 0.002 | 0.0103 | $8.46 \mathrm{E}-01$ | ??+ | $1.00 \mathrm{E}+00$ |
| WBC count | rs11899198 | 2 | 113840539 | t | g | 0.3744 | -0.0022 | 0.0022 | $3.38 \mathrm{E}-01$ | --+ | $5.28 \mathrm{E}-04$ |
| WBC count | rs12328368 | 2 | 113847144 | c | g | 0.5653 | -0.0109 | 0.0021 | $1.21 \mathrm{E}-07$ | --- | $2.28 \mathrm{E}-01$ |
| WBC count | rs12328766 | 2 | 113846738 | a | g | 0.5217 | -0.0098 | 0.0021 | 3.43E-06 | --- | 4.62E-01 |
| WBC count | rs12329129 | 2 | 113847007 | a | g | 0.3445 | 0.0108 | 0.0021 | $2.00 \mathrm{E}-07$ | +++ | 2.27E-01 |
| WBC count | rs12466799 | 2 | 113835999 | t | C | 0.566 | -0.0025 | 0.0021 | 2.33E-01 | --+ | 2.52E-01 |
| WBC count | rs12469822 | 2 | 113830563 | a | g | 0.3218 | 0.0038 | 0.0019 | $4.90 \mathrm{E}-02$ | ++- | $5.14 \mathrm{E}-01$ |
| WBC count | rs12470990 | 2 | 113924738 | t | c | 0.5134 | 0.0026 | 0.0034 | $4.55 \mathrm{E}-01$ | +?+ | 7.17E-01 |
| WBC count | rs12475161 | 2 | 113823626 | t | C | 0.4316 | -0.0014 | 0.0018 | $4.44 \mathrm{E}-01$ | --+ | $1.01 \mathrm{E}-03$ |
| WBC count | rs12475781 | 2 | 113916912 | t | g | 0.3051 | 0.0042 | 0.002 | 3.57E-02 | +++ | 4.29E-01 |
| WBC count | rs12475887 | 2 | 113849986 | a | c | 0.4035 | -2.00E-04 | 0.0024 | $9.50 \mathrm{E}-01$ | -++ | $1.23 \mathrm{E}-01$ |
| WBC count | rs12614131 | 2 | 113930867 | t | C | 0.3588 | 0.0018 | 0.0027 | 5.04E-01 | ++- | $1.20 \mathrm{E}-01$ |
| WBC count | rs12711746 | 2 | 113752247 | a | C | 0.4277 | 0.0032 | 0.002 | $9.87 \mathrm{E}-02$ | -++ | 3.19E-01 |
| WBC count | rs12711747 | 2 | 113785020 | a | g | 0.4022 | $2.00 \mathrm{E}-04$ | 0.002 | $9.34 \mathrm{E}-01$ | -+- | 2.82E-01 |
| WBC count | rs12711749 | 2 | 113829709 | a | g | 0.4848 | -0.0015 | 0.0018 | 4.12E-01 | -+- | $9.60 \mathrm{E}-02$ |
| WBC count | rs12711751 | 2 | 113837765 | t | g | 0.5337 | -0.0023 | 0.002 | $2.59 \mathrm{E}-01$ | --+ | 2.61E-02 |
| WBC count | rs12711752 | 2 | 113837840 | t | c | 0.562 | -9.00E-04 | 0.002 | 6.36E-01 | --+ | 5.91E-01 |
| WBC count | rs12711762 | 2 | 113932800 | t | C | 0.3161 | -3.00E-04 | 0.002 | $9.02 \mathrm{E}-01$ | --+ | $5.81 \mathrm{E}-01$ |
| WBC count | rs12711763 | 2 | 113932811 | C | g | 0.7857 | -0.0045 | 0.0032 | $1.60 \mathrm{E}-01$ | -+- | 7.50E-01 |
| WBC count | rs12711765 | 2 | 113936036 | a | g | 0.5017 | -0.0028 | 0.0039 | $4.74 \mathrm{E}-01$ | ??- | $1.00 \mathrm{E}+00$ |
| WBC count | rs128964 | 2 | 113887803 | a | g | 0.4888 | 0.0059 | 0.0026 | $2.27 \mathrm{E}-02$ | +++ | $6.79 \mathrm{E}-01$ |
| WBC count | rs13011842 | 2 | 113833570 | t | c | 0.3867 | -0.0027 | 0.0026 | 2.87E-01 | --+ | $5.64 \mathrm{E}-01$ |
| WBC count | rs13019891 | 2 | 113829869 | t | g | 0.4386 | 0.0038 | 0.002 | $6.20 \mathrm{E}-02$ | +++ | $9.47 \mathrm{E}-01$ |
| WBC count | rs13030546 | 2 | 113840014 | a | g | 0.5508 | 0.0031 | 0.0027 | $2.48 \mathrm{E}-01$ | -++ | $4.04 \mathrm{E}-01$ |
| WBC count | rs13033104 | 2 | 113756777 | t | g | 0.4674 | 0.0049 | 0.002 | $1.24 \mathrm{E}-02$ | ++- | $4.90 \mathrm{E}-01$ |
| WBC count | rs13382561 | 2 | 113863536 | a | g | 0.5636 | -0.0104 | 0.0021 | $1.28 \mathrm{E}-06$ | --- | $1.52 \mathrm{E}-01$ |
| WBC count | rs13386602 | 2 | 113834820 | a | c | 0.5207 | -0.006 | 0.0019 | $1.77 \mathrm{E}-03$ | --- | 5.82E-01 |
| WBC count | rs13389431 | 2 | 113851830 | a | g | 0.0884 | 2.00E-04 | 0.0064 | $9.75 \mathrm{E}-01$ | ??+ | $1.00 \mathrm{E}+00$ |
| WBC count | rs13398728 | 2 | 113842506 | t | c | 0.4349 | -0.0052 | 0.0021 | $1.33 \mathrm{E}-02$ | --+ | $3.38 \mathrm{E}-04$ |
| WBC count | rs13404928 | 2 | 113866525 | a | g | 0.2362 | -2.00E-04 | 0.0022 | 9.23E-01 | --+ | 4.49E-01 |
| WBC count | rs13406085 | 2 | 113861561 | a | g | 0.2624 | -0.005 | 0.0022 | $2.05 \mathrm{E}-02$ | --- | 2.77E-01 |
| WBC count | rs13407508 | 2 | 113794759 | t | C | 0.6701 | 8.00E-04 | 0.0024 | $7.38 \mathrm{E}-01$ | +?- | 2.20E-02 |
| WBC count | rs13409360 | 2 | 113838102 | a | g | 0.3925 | 0.0056 | 0.0021 | $7.74 \mathrm{E}-03$ | +++ | 7.97E-02 |

Table S1. Random-effects meta-analysis of MANTRA identified loci.

| WBC count | rs13409371 | 2 | 113838145 | a | g | 0.4052 | 0.0057 | 0.0021 | 6.81E-03 | +++ | 1.20E-01 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| WBC count | rs13410552 | 2 | 113855406 | c | g | 0.5813 | 0.0043 | 0.0022 | 4.67E-02 | ++- | $1.36 \mathrm{E}-01$ |
| WBC count | rs13410964 | 2 | 113843283 | a | g | 0.3262 | 0.0096 | 0.0021 | 5.35E-06 | +++ | $2.22 \mathrm{E}-01$ |
| WBC count | rs13414333 | 2 | 113752731 | a | g | 0.9226 | 0.0044 | 0.0069 | 5.23E-01 | ??+ | $1.00 \mathrm{E}+00$ |
| WBC count | rs13419549 | 2 | 113933726 | t | c | 0.3229 | -4.00E-04 | 0.0034 | 9.13E-01 | +-- | $2.78 \mathrm{E}-01$ |
| WBC count | rs13424580 | 2 | 113852564 | a | g | 0.3039 | 0.0086 | 0.0022 | $9.81 \mathrm{E}-05$ | +++ | 1.13E-01 |
| WBC count | rs1374280 | 2 | 113796666 | t | c | 0.4015 | 6.00E-04 | 0.0021 | 7.68E-01 | -++ | 8.49E-01 |
| WBC count | rs1374281 | 2 | 113898789 | c | g | 0.5901 | -0.0051 | 0.0021 | $1.35 \mathrm{E}-02$ | --- | 8.37E-01 |
| WBC count | rs1374284 | 2 | 113753583 | t | c | 0.5857 | -9.00E-04 | 0.002 | $6.58 \mathrm{E}-01$ | +-+ | 8.74E-02 |
| WBC count | rs1446509 | 2 | 113851159 | a | t | 0.5637 | -0.0105 | 0.0021 | 5.29E-07 | --- | 2.65E-01 |
| WBC count | rs1446510 | 2 | 113851517 | t | c | 0.4689 | 0.0015 | 0.0022 | 4.83E-01 | ++- | 6.01E-06 |
| WBC count | rs1446511 | 2 | 113762543 | t | C | 0.4751 | -6.00E-04 | 0.0026 | $8.05 \mathrm{E}-01$ | ?+- | $3.87 \mathrm{E}-01$ |
| WBC count | rs1446512 | 2 | 113762532 | t | g | 0.4751 | -6.00E-04 | 0.0026 | $8.28 \mathrm{E}-01$ | ?+- | $3.78 \mathrm{E}-01$ |
| WBC count | rs1446514 | 2 | 113757973 | t | g | 0.2951 | 5.00E-04 | 0.0023 | 8.18E-01 | ?-+ | $3.78 \mathrm{E}-01$ |
| WBC count | rs1446515 | 2 | 113757654 | a | t | 0.4745 | -0.002 | 0.0023 | 3.92E-01 | ?-- | 5.66E-01 |
| WBC count | rs1446516 | 2 | 113750470 | a | g | 0.5939 | -0.0025 | 0.0023 | 2.89E-01 | +-- | 1.59E-01 |
| WBC count | rs1446519 | 2 | 113822448 | t | c | 0.3881 | -0.0053 | 0.0029 | 7.11E-02 | -?+ | 2.03E-02 |
| WBC count | rs1446521 | 2 | 113827396 | a | g | 0.4857 | 0.0022 | 0.0026 | $4.08 \mathrm{E}-01$ | +++ | 4.93E-01 |
| WBC count | rs1446522 | 2 | 113827503 | C | g | 0.0934 | 7.00E-04 | 0.0031 | $8.14 \mathrm{E}-01$ | ++- | $1.26 \mathrm{E}-01$ |
| WBC count | rs1530549 | 2 | 113817455 | t | c | 0.3364 | -0.0046 | 0.003 | $1.26 \mathrm{E}-01$ | --- | 5.52E-01 |
| WBC count | rs1530550 | 2 | 113817513 | a | t | 0.5784 | 0.0046 | 0.0022 | 3.25E-02 | -++ | $6.88 \mathrm{E}-02$ |
| WBC count | rs1530551 | 2 | 113817566 | t | C | 0.482 | 0.0018 | 0.0022 | $4.06 \mathrm{E}-01$ | +-+ | $2.46 \mathrm{E}-02$ |
| WBC count | rs1542176 | 2 | 113849837 | t | c | 0.3579 | -0.0013 | 0.0019 | $4.90 \mathrm{E}-01$ | -+- | $3.66 \mathrm{E}-03$ |
| WBC count | rs1562302 | 2 | 113810458 | a | g | 0.3777 | 0.0011 | 0.002 | 5.89E-01 | +-- | 1.85E-01 |
| WBC count | rs1562303 | 2 | 113782392 | t | c | 0.2935 | 0.0073 | 0.004 | 6.89E-02 | ??+ | $1.00 \mathrm{E}+00$ |
| WBC count | rs1562305 | 2 | 113751613 | a | g | 0.3701 | 0.0028 | 0.002 | $1.62 \mathrm{E}-01$ | ++- | 4.51E-01 |
| WBC count | rs1562306 | 2 | 113751380 | a | g | 0.2937 | 0.002 | 0.0023 | 3.86E-01 | -++ | 7.36E-01 |
| WBC count | rs1596894 | 2 | 113754754 | t | c | 0.4851 | 6.00E-04 | 0.0025 | 7.96E-01 | ?+- | 8.15E-01 |
| WBC count | rs1630153 | 2 | 113869803 | t | c | 0.34 | 0.0032 | 0.0019 | 8.82E-02 | -++ | 5.83E-03 |
| WBC count | rs1665188 | 2 | 113852789 | c | g | 0.0769 | -0.0143 | 0.0054 | 7.89E-03 | -?- | 4.03E-01 |
| WBC count | rs1688072 | 2 | 113869347 | a | g | 0.4789 | 0.0059 | 0.0029 | $3.90 \mathrm{E}-02$ | +++ | $1.35 \mathrm{E}-01$ |
| WBC count | rs1688075 | 2 | 113858196 | a | c | 0.3855 | -0.0016 | 0.0025 | 5.22E-01 | -++ | 8.58E-02 |
| WBC count | rs17042691 | 2 | 113745812 | t | c | 0.34 | 3.00E-04 | 0.0022 | 8.92E-01 | -+- | 4.36E-02 |
| WBC count | rs17042709 | 2 | 113788100 | a | g | 0.2321 | -0.0065 | 0.0027 | 1.63E-02 | -?- | 8.09E-01 |
| WBC count | rs17042712 | 2 | 113788194 | c | g | 0.7978 | 0.001 | 0.0047 | 8.33E-01 | ?+- | 5.56E-02 |
| WBC count | rs17042750 | 2 | 113823151 | t | g | 0.0983 | -0.0038 | 0.0037 | 3.01E-01 | ?+- | $1.42 \mathrm{E}-03$ |
| WBC count | rs17042751 | 2 | 113823365 | t | c | 0.395 | 0.0019 | 0.0026 | 4.67E-01 | ++- | 4.02E-01 |
| WBC count | rs17042755 | 2 | 113824585 | C | g | 0.1835 | 0.0023 | 0.0026 | 3.78E-01 | +++ | 4.61E-01 |

Table S1. Random-effects meta-analysis of MANTRA identified loci.

| WBC count | rs17042786 | 2 | 113827859 | t | c | 0.4314 | -0.007 | 0.0026 | 7.70E-03 | -?- | 3.96E-01 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| WBC count | rs17042795 | 2 | 113829840 | a | g | 0.6782 | -0.016 | 0.0039 | 3.61E-05 | ?-- | 4.31E-01 |
| WBC count | rs17042810 | 2 | 113838569 | t | c | 0.3626 | -0.0026 | 0.0025 | $3.03 \mathrm{E}-01$ | --+ | 1.93E-04 |
| WBC count | rs17042815 | 2 | 113839857 | c | g | 0.0865 | 0.0151 | 0.0052 | $3.47 \mathrm{E}-03$ | ?++ | $4.88 \mathrm{E}-01$ |
| WBC count | rs17042819 | 2 | 113840039 | a | g | 0.7307 | -0.0127 | 0.0051 | $1.26 \mathrm{E}-02$ | ?-- | $6.53 \mathrm{E}-01$ |
| WBC count | rs17042827 | 2 | 113840657 | a | g | 0.9737 | -0.0179 | 0.0116 | $1.24 \mathrm{E}-01$ | ??- | $1.00 \mathrm{E}+00$ |
| WBC count | rs17042828 | 2 | 113840934 | t | c | 0.087 | 0.0112 | 0.0051 | 2.76E-02 | ?-+ | 3.92E-02 |
| WBC count | rs17042833 | 2 | 113842279 | t | c | 0.7301 | -0.0093 | 0.0051 | 6.87E-02 | ?+- | 6.32E-02 |
| WBC count | rs17042838 | 2 | 113843337 | a | g | 0.1572 | $3.00 \mathrm{E}-04$ | 0.0049 | $9.52 \mathrm{E}-01$ | ??+ | $1.00 \mathrm{E}+00$ |
| WBC count | rs17042842 | 2 | 113843481 | a | g | 0.7399 | -0.016 | 0.0049 | $1.21 \mathrm{E}-03$ | ?-- | 4.29E-01 |
| WBC count | rs17042853 | 2 | 113848200 | a | t | 0.2659 | -0.0054 | 0.0021 | $1.14 \mathrm{E}-02$ | --- | 3.61E-01 |
| WBC count | rs17042869 | 2 | 113849691 | a | g | 0.0419 | 0.0255 | 0.0093 | 5.90E-03 | ??+ | $1.00 \mathrm{E}+00$ |
| WBC count | rs17042888 | 2 | 113862173 | a | g | 0.1744 | -0.0094 | 0.0024 | $8.74 \mathrm{E}-05$ | --- | 4.56E-01 |
| WBC count | rs17042894 | 2 | 113864047 | a | g | 0.0704 | 0.0152 | 0.0054 | 4.62E-03 | ?++ | 5.09E-01 |
| WBC count | rs17042905 | 2 | 113867956 | t | c | 0.36 | -0.0027 | 0.0024 | $2.61 \mathrm{E}-01$ | --+ | 5.60E-05 |
| WBC count | rs17042917 | 2 | 113870663 | a | g | 0.1702 | -0.0096 | 0.0024 | $7.04 \mathrm{E}-05$ | --- | $2.02 \mathrm{E}-01$ |
| WBC count | rs17043030 | 2 | 113926885 | t | c | 0.709 | -0.0028 | 0.0042 | $4.99 \mathrm{E}-01$ | ?+- | $1.26 \mathrm{E}-01$ |
| WBC count | rs17043031 | 2 | 113930864 | a | g | 0.8963 | 0.0043 | 0.0069 | $5.38 \mathrm{E}-01$ | +?+ | 6.99E-01 |
| WBC count | rs17043037 | 2 | 113932352 | t | g | 0.0471 | 0.0162 | 0.0144 | $2.61 \mathrm{E}-01$ | ??+ | $1.00 \mathrm{E}+00$ |
| WBC count | rs17043041 | 2 | 113933432 | t | g | 0.9406 | -0.0044 | 0.0113 | $6.98 \mathrm{E}-01$ | ??- | $1.00 \mathrm{E}+00$ |
| WBC count | rs17043051 | 2 | 113933804 | t | C | 0.5665 | -0.0015 | 0.0028 | 5.86E-01 | -?+ | $3.67 \mathrm{E}-02$ |
| WBC count | rs17043057 | 2 | 113934413 | t | C | 0.9525 | -0.0164 | 0.0142 | $2.48 \mathrm{E}-01$ | ??- | $1.00 \mathrm{E}+00$ |
| WBC count | rs17043069 | 2 | 113935807 | t | g | 0.8678 | 0.0042 | 0.0059 | $4.74 \mathrm{E}-01$ | ??+ | $1.00 \mathrm{E}+00$ |
| WBC count | rs17207494 | 2 | 113864010 | a | C | 0.582 | -0.0088 | 0.0022 | $6.34 \mathrm{E}-05$ | --- | 7.57E-02 |
| WBC count | rs17486819 | 2 | 113871806 | t | g | 0.0808 | -0.0027 | 0.004 | $4.90 \mathrm{E}-01$ | ?+- | $1.86 \mathrm{E}-03$ |
| WBC count | rs17624213 | 2 | 113933247 | t | g | 0.2778 | -0.0027 | 0.0026 | $3.07 \mathrm{E}-01$ | --- | 9.52E-01 |
| WBC count | rs17660913 | 2 | 113782329 | a | g | 0.559 | 7.00E-04 | 0.0028 | $8.09 \mathrm{E}-01$ | --+ | $2.30 \mathrm{E}-02$ |
| WBC count | rs17669228 | 2 | 113847712 | t | c | 0.3496 | 0.0059 | 0.003 | $4.71 \mathrm{E}-02$ | ++- | 4.05E-03 |
| WBC count | rs1794066 | 2 | 113886350 | a | g | 0.4233 | 0.0042 | 0.0018 | $1.74 \mathrm{E}-02$ | -++ | 3.37E-02 |
| WBC count | rs1794067 | 2 | 113886384 | a | g | 0.2199 | -0.0037 | 0.0023 | $1.09 \mathrm{E}-01$ | 0 | $5.57 \mathrm{E}-01$ |
| WBC count | rs1794068 | 2 | 113886503 | a | g | 0.2078 | -0.0063 | 0.0026 | $1.48 \mathrm{E}-02$ | --- | 5.55E-01 |
| WBC count | rs1800930 | 2 | 113820530 | a | g | 0.2762 | -0.009 | 0.0025 | $2.68 \mathrm{E}-04$ | -?- | 7.17E-01 |
| WBC count | rs1867761 | 2 | 113922983 | t | C | 0.328 | 0.0038 | 0.0029 | $1.91 \mathrm{E}-01$ | +++ | $9.74 \mathrm{E}-01$ |
| WBC count | rs1867828 | 2 | 113768620 | a | g | 0.3852 | 0.0038 | 0.0019 | $4.60 \mathrm{E}-02$ | +++ | 7.01E-01 |
| WBC count | rs1867829 | 2 | 113839151 | a | g | 0.46 | 0.0039 | 0.0021 | $6.36 \mathrm{E}-02$ | +-+ | 7.81E-02 |
| WBC count | rs1867831 | 2 | 113773561 | t | C | 0.4248 | $1.00 \mathrm{E}-04$ | 0.0025 | $9.80 \mathrm{E}-01$ | -++ | $3.97 \mathrm{E}-01$ |
| WBC count | rs1867834 | 2 | 113828099 | t | C | 0.4276 | 0.0039 | 0.0021 | 5.65E-02 | +++ | 9.69E-01 |
| WBC count | rs1900287 | 2 | 113797565 | a | g | 0.5339 | $1.00 \mathrm{E}-04$ | 0.0025 | $9.62 \mathrm{E}-01$ | -+- | 8.78E-01 |

Table S1. Random-effects meta-analysis of MANTRA identified loci.

| WBC count | rs1992761 | 2 | 113810286 | a | g | 0.113 | -0.0091 | 0.0035 | $9.09 \mathrm{E}-03$ | ?-- | 3.68E-01 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| WBC count | rs2029582 | 2 | 113863771 | t | c | 0.6258 | 6.00E-04 | 0.0018 | $7.34 \mathrm{E}-01$ | +-+ | 6.56E-04 |
| WBC count | rs2071459 | 2 | 113887483 | t | C | 0.567 | 7.00E-04 | 0.0021 | 7.23E-01 | +-+ | $4.25 \mathrm{E}-01$ |
| WBC count | rs2100071 | 2 | 113825386 | a | C | 0.522 | 0 | 0.0025 | 9.96E-01 | +-- | 8.73E-01 |
| WBC count | rs2121326 | 2 | 113846975 | c | g | 0.1516 | -0.0064 | 0.0028 | $2.14 \mathrm{E}-02$ | --- | 9.58E-02 |
| WBC count | rs2121329 | 2 | 113852629 | a | C | 0.2608 | -0.0045 | 0.0022 | $3.56 \mathrm{E}-02$ | --+ | $1.26 \mathrm{E}-01$ |
| WBC count | rs2197578 | 2 | 113790553 | a | g | 0.5528 | -5.00E-04 | 0.0023 | $8.31 \mathrm{E}-01$ | -+- | 7.85E-01 |
| WBC count | rs2248588 | 2 | 113898710 | t | c | 0.7957 | 0.0019 | 0.0067 | 7.76E-01 | ??+ | $1.00 \mathrm{E}+00$ |
| WBC count | rs2248596 | 2 | 113898827 | a | g | 0.1956 | $8.00 \mathrm{E}-04$ | 0.0047 | $8.66 \mathrm{E}-01$ | ??+ | $1.00 \mathrm{E}+00$ |
| WBC count | rs2248600 | 2 | 113898879 | a | g | 0.8494 | 0.0012 | 0.0076 | $8.75 \mathrm{E}-01$ | ??+ | $1.00 \mathrm{E}+00$ |
| WBC count | rs2248604 | 2 | 113898932 | a | c | 0.1594 | -0.0013 | 0.0071 | $8.55 \mathrm{E}-01$ | ??- | $1.00 \mathrm{E}+00$ |
| WBC count | rs2305150 | 2 | 113789316 | t | C | 0.5366 | 0.0015 | 0.0018 | $4.01 \mathrm{E}-01$ | ++- | $2.62 \mathrm{E}-01$ |
| WBC count | rs2305152 | 2 | 113763463 | a | C | 0.5444 | 0.0047 | 0.0021 | $2.41 \mathrm{E}-02$ | ++- | $7.75 \mathrm{E}-01$ |
| WBC count | rs2472188 | 2 | 113820814 | c | g | 0.3273 | -0.0056 | 0.0018 | 2.19E-03 | --- | $1.48 \mathrm{E}-01$ |
| WBC count | rs2515401 | 2 | 113820476 | t | c | 0.4305 | -0.0014 | 0.0018 | $4.32 \mathrm{E}-01$ | --+ | 2.15E-03 |
| WBC count | rs2515402 | 2 | 113820580 | a | C | 0.6096 | 0.0054 | 0.0019 | $4.64 \mathrm{E}-03$ | +++ | $9.77 \mathrm{E}-02$ |
| WBC count | rs2515404 | 2 | 113821117 | t | c | 0.3949 | -7.00E-04 | 0.0026 | $7.97 \mathrm{E}-01$ | +-- | 5.16E-01 |
| WBC count | rs2515406 | 2 | 113822168 | t | c | 0.4163 | 6.00E-04 | 0.0027 | 8.22E-01 | +-- | 2.56E-01 |
| WBC count | rs2637988 | 2 | 113876779 | a | g | 0.3912 | 0.0019 | 0.0018 | 2.97E-01 | -++ | $3.96 \mathrm{E}-01$ |
| WBC count | rs2853628 | 2 | 113880292 | C | g | 0.2497 | -0.0031 | 0.0026 | $2.35 \mathrm{E}-01$ | ?-+ | $3.50 \mathrm{E}-01$ |
| WBC count | rs2862772 | 2 | 113791482 | C | g | 0.5755 | 0.002 | 0.0018 | $2.55 \mathrm{E}-01$ | +-+ | 4.01E-01 |
| WBC count | rs2862774 | 2 | 113797677 | a | C | 0.5307 | 0.003 | 0.0026 | $2.37 \mathrm{E}-01$ | -++ | $3.35 \mathrm{E}-02$ |
| WBC count | rs2862853 | 2 | 113826759 | t | c | 0.3596 | -0.001 | 0.0024 | 6.82E-01 | ++- | $3.48 \mathrm{E}-01$ |
| WBC count | rs2902452 | 2 | 113895516 | a | C | 0.3077 | 0.0047 | 0.0021 | $2.61 \mathrm{E}-02$ | +++ | 7.71E-01 |
| WBC count | rs3087263 | 2 | 113885768 | a | g | 0.0757 | -0.0021 | 0.005 | 6.73E-01 | ?-- | $3.24 \mathrm{E}-01$ |
| WBC count | rs3087266 | 2 | 113889100 | t | c | 0.5215 | 2.00E-04 | 0.002 | $9.11 \mathrm{E}-01$ | +-+ | 1.06E-01 |
| WBC count | rs3087270 | 2 | 113893308 | a | g | 0.5737 | 0.0031 | 0.0021 | $1.34 \mathrm{E}-01$ | +++ | 8.43E-01 |
| WBC count | rs3087271 | 2 | 113895200 | a | c | 0.5097 | 0.0022 | 0.0026 | 3.97E-01 | -++ | $1.02 \mathrm{E}-01$ |
| WBC count | rs3099479 | 2 | 113898461 | a | g | 0.7958 | 0.0018 | 0.0067 | 7.88E-01 | ??+ | $1.00 \mathrm{E}+00$ |
| WBC count | rs315919 | 2 | 113876213 | t | g | 0.603 | 0.0017 | 0.0023 | $4.67 \mathrm{E}-01$ | +?+ | 8.51E-01 |
| WBC count | rs315920 | 2 | 113873018 | t | C | 0.4157 | -0.0019 | 0.0028 | 4.92E-01 | --- | 8.39E-01 |
| WBC count | rs315921 | 2 | 113872051 | a | g | 0.1649 | -0.0028 | 0.0033 | $3.99 \mathrm{E}-01$ | --- | 9.12E-01 |
| WBC count | rs315922 | 2 | 113862647 |  | c | 0.2908 | -0.0012 | 0.0031 | 7.11E-01 | --+ | 2.68E-02 |
| WBC count | rs315925 | 2 | 113865367 | t | C | 0.5852 | 0.0028 | 0.0025 | 2.51E-01 | +-+ | 7.56E-02 |
| WBC count | rs315927 | 2 | 113866369 | t | C | 0.3416 | 0.0028 | 0.0018 | $1.30 \mathrm{E}-01$ | -++ | $9.84 \mathrm{E}-03$ |
| WBC count | rs315928 | 2 | 113868263 | t | c | 0.3954 | 8.00E-04 | 0.0029 | 7.77E-01 | --+ | $3.64 \mathrm{E}-02$ |
| WBC count | rs315929 | 2 | 113869407 | t | C | 0.5832 | 0 | 0.0024 | $9.97 \mathrm{E}-01$ | +-- | $7.48 \mathrm{E}-02$ |
| WBC count | rs315931 | 2 | 113869843 | a | C | 0.3627 | -0.0022 | 0.0019 | 2.27E-01 | -+- | $5.10 \mathrm{E}-03$ |

Table S1. Random-effects meta-analysis of MANTRA identified loci.

| WBC count | rs315932 | 2 | 113869977 | t | c | 0.5742 | 0.0024 | 0.0025 | $3.46 \mathrm{E}-01$ | +-+ | 1.03E-01 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| WBC count | rs315935 | 2 | 113881365 | a | g | 0.8679 | -0.0024 | 0.0055 | $6.60 \mathrm{E}-01$ | ??- | $1.00 \mathrm{E}+00$ |
| WBC count | rs315936 | 2 | 113880947 | t | c | 0.4368 | -0.0033 | 0.0024 | $1.74 \mathrm{E}-01$ | --+ | 3.17E-01 |
| WBC count | rs315937 | 2 | 113897154 | c | g | 0.7557 | 0.0027 | 0.0063 | 6.67E-01 | ??+ | $1.00 \mathrm{E}+00$ |
| WBC count | rs315938 | 2 | 113896045 | a | t | 0.8452 | -0.0045 | 0.0049 | $3.62 \mathrm{E}-01$ | ??- | $1.00 \mathrm{E}+00$ |
| WBC count | rs315939 | 2 | 113895403 | a | C | 0.8433 | $1.00 \mathrm{E}-04$ | 0.0074 | $9.89 \mathrm{E}-01$ | ??+ | $1.00 \mathrm{E}+00$ |
| WBC count | rs315940 | 2 | 113895305 | t | C | 0.8476 | -0.0043 | 0.0051 | 4.03E-01 | ??- | $1.00 \mathrm{E}+00$ |
| WBC count | rs315941 | 2 | 113895235 | a | g | 0.8435 | 2.00E-04 | 0.0074 | $9.79 \mathrm{E}-01$ | ??+ | $1.00 \mathrm{E}+00$ |
| WBC count | rs315942 | 2 | 113895145 | t | g | 0.5534 | -0.003 | 0.002 | $1.36 \mathrm{E}-01$ | --+ | 4.59E-01 |
| WBC count | rs315943 | 2 | 113894338 | a | g | 0.5797 | -0.0063 | 0.0021 | 3.15E-03 | --- | $4.61 \mathrm{E}-01$ |
| WBC count | rs315944 | 2 | 113894187 | a | g | 0.147 | -0.0019 | 0.0078 | 8.08E-01 | ??- | $1.00 \mathrm{E}+00$ |
| WBC count | rs315947 | 2 | 113893839 | a | g | 0.4581 | -0.003 | 0.0018 | $1.02 \mathrm{E}-01$ | --- | $6.24 \mathrm{E}-01$ |
| WBC count | rs315948 | 2 | 113893256 | t | C | 0.4328 | -0.0018 | 0.003 | $5.39 \mathrm{E}-01$ | +-- | $9.00 \mathrm{E}-01$ |
| WBC count | rs315949 | 2 | 113892774 | a | g | 0.3496 | 0.007 | 0.002 | $6.40 \mathrm{E}-04$ | +++ | $5.20 \mathrm{E}-01$ |
| WBC count | rs315951 | 2 | 113890586 | C | g | 0.399 | 0.0025 | 0.0018 | $1.62 \mathrm{E}-01$ | +++ | 6.82E-01 |
| WBC count | rs315952 | 2 | 113890304 | t | C | 0.356 | 7.00E-04 | 0.0019 | 7.17E-01 | -+- | $1.98 \mathrm{E}-01$ |
| WBC count | rs315955 | 2 | 113889430 | C | g | 0.9107 | -0.0127 | 0.0065 | 5.01E-02 | ??- | $1.00 \mathrm{E}+00$ |
| WBC count | rs315957 | 2 | 113901313 | t | c | 0.7325 | -0.0046 | 0.0027 | 8.96E-02 | -?- | 1.11E-01 |
| WBC count | rs315958 | 2 | 113900851 | t | g | 0.5744 | -0.0043 | 0.002 | $2.94 \mathrm{E}-02$ | --- | $2.56 \mathrm{E}-01$ |
| WBC count | rs3180235 | 2 | 113820672 | a | g | 0.3279 | -0.0056 | 0.0018 | 2.39E-03 | --- | $1.56 \mathrm{E}-01$ |
| WBC count | rs3180238 | 2 | 113820833 | t | C | 0.6116 | 0.0114 | 0.009 | $2.08 \mathrm{E}-01$ | +?+ | 8.05E-01 |
| WBC count | rs3181052 | 2 | 113886049 | a | g | 0.4058 | 4.00E-04 | 0.0021 | 8.65E-01 | +-- | $4.36 \mathrm{E}-01$ |
| WBC count | rs3181053 | 2 | 113886142 | t | g | 0.9512 | 0.0211 | 0.0085 | $1.35 \mathrm{E}-02$ | ??+ | $1.00 \mathrm{E}+00$ |
| WBC count | rs3213448 | 2 | 113879297 | a | g | 0.4212 | 5.00E-04 | 0.0021 | 8.24E-01 | +-- | $4.14 \mathrm{E}-01$ |
| WBC count | rs35029104 | 2 | 113772596 | t | C | 0.0688 | 0.0041 | 0.0075 | 5.85E-01 | ??+ | $1.00 \mathrm{E}+00$ |
| WBC count | rs3748914 | 2 | 113940681 | t | c | 0.4307 | -9.00E-04 | 0.0018 | $6.13 \mathrm{E}-01$ | --+ | 5.49E-01 |
| WBC count | rs3752739 | 2 | 113788590 | C | g | 0.7706 | 0.0061 | 0.003 | $4.31 \mathrm{E}-02$ | +?+ | 9.43E-01 |
| WBC count | rs3762494 | 2 | 113937563 | a | g | 0.7537 | -0.0071 | 0.0029 | $1.49 \mathrm{E}-02$ | -+- | 3.81E-01 |
| WBC count | rs377086 | 2 | 113878715 | a | g | 0.1228 | 0.0028 | 0.0055 | 6.13E-01 | +?+ | 2.35E-01 |
| WBC count | rs380092 | 2 | 113888900 | a | t | 0.3581 | 8.00E-04 | 0.0019 | $6.54 \mathrm{E}-01$ | -++ | 3.62E-01 |
| WBC count | rs3811050 | 2 | 113830173 | t | C | 0.3958 | 8.00E-04 | 0.0021 | $6.92 \mathrm{E}-01$ | -++ | $2.76 \mathrm{E}-02$ |
| WBC count | rs3811053 | 2 | 113830926 | t | c | 0.523 | 0.0059 | 0.0021 | $5.62 \mathrm{E}-03$ | +++ | 4.59E-01 |
| WBC count | rs3827763 | 2 | 113831289 | a | g | 0.4054 | 0.0016 | 0.0019 | 4.07E-01 | +-+ | $2.40 \mathrm{E}-01$ |
| WBC count | rs396201 | 2 | 113891775 | t | c | 0.4582 | 0.0014 | 0.002 | $4.86 \mathrm{E}-01$ | ++- | $2.48 \mathrm{E}-01$ |
| WBC count | rs397211 | 2 | 113892141 | t | C | 0.4635 | 7.00E-04 | 0.0021 | $7.34 \mathrm{E}-01$ | ++- | 3.43E-01 |
| WBC count | rs408392 | 2 | 113887458 | t | g | 0.3666 | -0.0027 | 0.0025 | $2.86 \mathrm{E}-01$ | --+ | 7.57E-02 |
| WBC count | rs41334144 | 2 | 113766332 | t | c | 0.8487 | 0.0079 | 0.0086 | 3.61E-01 | ??+ | $1.00 \mathrm{E}+00$ |
| WBC count | rs41411944 | 2 | 113922486 | t | C | 0.9068 | -0.0034 | 0.0068 | 6.17E-01 | ??- | $1.00 \mathrm{E}+00$ |

Table S1. Random-effects meta-analysis of MANTRA identified loci.

| WBC count | rs4145013 | 2 | 113830688 | a | g | 0.5081 | 0.0053 | 0.0018 | 3.77E-03 | +++ | $2.18 \mathrm{E}-01$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| WBC count | rs419598 | 2 | 113887207 | t | c | 0.3546 | 0.0016 | 0.0027 | 5.57E-01 | ++- | 1.20E-01 |
| WBC count | rs423904 | 2 | 113887262 | t | C | 0.3659 | -0.0026 | 0.0025 | 3.06E-01 | --+ | $8.43 \mathrm{E}-02$ |
| WBC count | rs4251955 | 2 | 113873871 | a | g | 0.0479 | -0.0126 | 0.0086 | $1.45 \mathrm{E}-01$ | ??- | $1.00 \mathrm{E}+00$ |
| WBC count | rs4251961 | 2 | 113874467 | t | c | 0.4359 | -0.0034 | 0.0022 | 1.12E-01 | --+ | 2.03E-03 |
| WBC count | rs4251967 | 2 | 113874968 | C | g | 0.0578 | -0.0119 | 0.0052 | $2.29 \mathrm{E}-02$ | -?- | $6.80 \mathrm{E}-01$ |
| WBC count | rs4251984 | 2 | 113877226 | a | g | 0.5098 | 0.0055 | 0.0026 | 3.49E-02 | +++ | $8.68 \mathrm{E}-01$ |
| WBC count | rs4251985 | 2 | 113877413 | t | g | 0.3696 | -0.0031 | 0.0027 | 2.50E-01 | --+ | $1.09 \mathrm{E}-01$ |
| WBC count | rs4251995 | 2 | 113881050 | t | g | 0.9859 | 0.0149 | 0.0171 | 3.83E-01 | ??+ | $1.00 \mathrm{E}+00$ |
| WBC count | rs4252001 | 2 | 113883683 | a | g | 0.4541 | 0.0011 | 0.0021 | 6.12E-01 | -++ | 3.99E-01 |
| WBC count | rs4252012 | 2 | 113887349 | a | c | 0.046 | 0.0082 | 0.0089 | 3.54E-01 | ??+ | $1.00 \mathrm{E}+00$ |
| WBC count | rs4252019 | 2 | 113889119 | t | C | 0.5219 | 5.00E-04 | 0.002 | 8.23E-01 | +-+ | $1.01 \mathrm{E}-01$ |
| WBC count | rs4252022 | 2 | 113890161 | a | g | 0.0562 | -0.0055 | 0.0082 | $5.04 \mathrm{E}-01$ | ??- | $1.00 \mathrm{E}+00$ |
| WBC count | rs4252033 | 2 | 113891842 | a | c | 0.8539 | -5.00E-04 | 0.0052 | $9.24 \mathrm{E}-01$ | ??- | $1.00 \mathrm{E}+00$ |
| WBC count | rs4252035 | 2 | 113892563 | t | C | 0.0319 | 0.0072 | 0.016 | 6.52E-01 | ??+ | $1.00 \mathrm{E}+00$ |
| WBC count | rs4252036 | 2 | 113892964 | a | C | 0.017 | 0.0202 | 0.0149 | $1.76 \mathrm{E}-01$ | ??+ | $1.00 \mathrm{E}+00$ |
| WBC count | rs4252041 | 2 | 113890610 | t | c | 0.2199 | 0.0097 | 0.0075 | 1.98E-01 | ?+- | $1.41 \mathrm{E}-01$ |
| WBC count | rs431726 | 2 | 113889006 | t | g | 0.3182 | 0.0031 | 0.0025 | 2.19E-01 | ++- | 2.92E-01 |
| WBC count | rs432014 | 2 | 113888579 | t | c | 0.3062 | 5.00E-04 | 0.0024 | 8.20E-01 | ++- | 6.15E-03 |
| WBC count | rs4368340 | 2 | 113845080 | a | t | 0.5289 | -0.0032 | 0.0019 | $8.64 \mathrm{E}-02$ | +-- | $2.40 \mathrm{E}-03$ |
| WBC count | rs440286 | 2 | 113889469 | a | C | 0.2086 | -0.0036 | 0.0027 | 1.80E-01 | --- | $9.91 \mathrm{E}-01$ |
| WBC count | rs442710 | 2 | 113887399 | a | g | 0.2094 | -0.0062 | 0.0026 | 1.82E-02 | --- | 5.68E-01 |
| WBC count | rs444413 | 2 | 113885826 | a | g | 0.2049 | -0.0057 | 0.0026 | 2.79E-02 | --- | 8.10E-01 |
| WBC count | rs446433 | 2 | 113887273 | a | g | 0.2069 | -0.0059 | 0.0026 | $2.26 \mathrm{E}-02$ | --- | $6.25 \mathrm{E}-01$ |
| WBC count | rs447713 | 2 | 113887672 | a | g | 0.5006 | 0.0057 | 0.0026 | 2.51E-02 | +++ | 8.12E-01 |
| WBC count | rs4496335 | 2 | 113844475 | t | c | 0.5066 | 0.0057 | 0.0022 | 8.85E-03 | ++- | 1.16E-02 |
| WBC count | rs451578 | 2 | 113888557 | a | g | 0.2049 | -0.0064 | 0.0026 | $1.48 \mathrm{E}-02$ | --- | $4.27 \mathrm{E}-01$ |
| WBC count | rs452204 | 2 | 113889061 | a | g | 0.5077 | -0.0045 | 0.0018 | $1.18 \mathrm{E}-02$ | +-- | 1.17E-01 |
| WBC count | rs454078 | 2 | 113888793 | a | t | 0.4971 | 0.0058 | 0.0026 | 2.29E-02 | +++ | $8.01 \mathrm{E}-01$ |
| WBC count | rs4848314 | 2 | 113839286 | a | g | 0.4534 | 0.0013 | 0.0024 | 6.05E-01 | +-- | 5.80E-02 |
| WBC count | rs4848316 | 2 | 113934451 | t | c | 0.4194 | -7.00E-04 | 0.0018 | $6.81 \mathrm{E}-01$ | +-+ | 7.63E-01 |
| WBC count | rs4849136 | 2 | 113748202 | t | c | 0.2951 | 8.00E-04 | 0.0023 | 7.24E-01 | ?-+ | 2.96E-01 |
| WBC count | rs4849142 | 2 | 113779875 | a | g | 0.5068 | 0.001 | 0.0019 | $6.06 \mathrm{E}-01$ | +-- | 6.10E-02 |
| WBC count | rs4849143 | 2 | 113787356 | C | g | 0.5146 | 0.0011 | 0.0025 | $6.78 \mathrm{E}-01$ | -+- | $8.09 \mathrm{E}-01$ |
| WBC count | rs4849144 | 2 | 113800049 | a | g | 0.2431 | -7.00E-04 | 0.0025 | 7.90E-01 | + | 7.70E-01 |
| WBC count | rs4849147 | 2 | 113829236 | a | t | 0.2511 | -0.0022 | 0.0026 | $4.04 \mathrm{E}-01$ | --- | 4.14E-01 |
| WBC count | rs4849148 | 2 | 113829522 | t | C | 0.3513 | $4.00 \mathrm{E}-04$ | 0.0022 | 8.42E-01 | -++ | $9.80 \mathrm{E}-02$ |
| WBC count | rs4849152 | 2 | 113848574 | a | g | 0.5245 | 0.0069 | 0.0019 | 2.38E-04 | +++ | 3.20E-01 |

Table S1. Random-effects meta-analysis of MANTRA identified loci.

| WBC count | rs4849153 | 2 | 113848654 | t | c | 0.5702 | -4.00E-04 | 0.002 | 8.20E-01 | +-+ | 2.45E-02 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| WBC count | rs4849159 | 2 | 113925061 | t | c | 0.309 | 7.00E-04 | 0.0022 | $7.34 \mathrm{E}-01$ | +-- | 2.63E-01 |
| WBC count | rs4849160 | 2 | 113934085 | a | g | 0.261 | -0.0013 | 0.0022 | $5.48 \mathrm{E}-01$ | --- | 8.02E-01 |
| WBC count | rs4849163 | 2 | 113935028 | t | c | 0.3166 | 0.0017 | 0.0021 | 4.36E-01 | +++ | $4.68 \mathrm{E}-01$ |
| WBC count | rs4849166 | 2 | 113939763 | a | t | 0.5418 | 0.0011 | 0.0018 | $5.29 \mathrm{E}-01$ | ++- | 7.19E-01 |
| WBC count | rs4849167 | 2 | 113940839 | C | g | 0.5217 | $4.00 \mathrm{E}-04$ | 0.0018 | 8.09E-01 | ++- | 2.81E-01 |
| WBC count | rs495282 | 2 | 113887294 | C | g | 0.2071 | -0.0059 | 0.0026 | 2.27E-02 | --- | 6.25E-01 |
| WBC count | rs495410 | 2 | 113887338 | a | c | 0.9392 | 0.0099 | 0.0049 | $4.22 \mathrm{E}-02$ | +?+ | $6.81 \mathrm{E}-01$ |
| WBC count | rs579543 | 2 | 113889631 | a | g | 0.215 | -0.0022 | 0.0027 | $4.05 \mathrm{E}-01$ | --- | $9.34 \mathrm{E}-01$ |
| WBC count | rs6542108 | 2 | 113755190 | t | c | 0.3654 | 4.00E-04 | 0.0021 | $8.50 \mathrm{E}-01$ | -+- | 1.32E-01 |
| WBC count | rs6542109 | 2 | 113756821 | t | c | 0.3766 | 0.001 | 0.002 | 6.26E-01 | -+- | $9.79 \mathrm{E}-02$ |
| WBC count | rs6542111 | 2 | 113822559 | t | C | 0.0406 | 0.0165 | 0.0082 | $4.51 \mathrm{E}-02$ | ??+ | $1.00 \mathrm{E}+00$ |
| WBC count | rs6542113 | 2 | 113859276 | a | g | 0.0707 | 0.0155 | 0.0055 | 4.85E-03 | ?++ | $4.72 \mathrm{E}-01$ |
| WBC count | rs6542114 | 2 | 113862825 | a | g | 0.0595 | -0.0109 | 0.0078 | $1.63 \mathrm{E}-01$ | ??- | $1.00 \mathrm{E}+00$ |
| WBC count | rs6542118 | 2 | 113912081 | a | g | 0.4871 | -0.0037 | 0.002 | 6.10E-02 | --+ | $1.84 \mathrm{E}-01$ |
| WBC count | rs6708535 | 2 | 113833839 | t | C | 0.1678 | 0.0046 | 0.0042 | 2.72E-01 | ?-+ | 1.55E-01 |
| WBC count | rs6710007 | 2 | 113780930 | a | g | 0.505 | $9.00 \mathrm{E}-04$ | 0.0019 | $6.31 \mathrm{E}-01$ | +-- | $5.94 \mathrm{E}-02$ |
| WBC count | rs6714534 | 2 | 113759330 | t | c | 0.3954 | -0.0018 | 0.0023 | $4.32 \mathrm{E}-01$ | +-+ | 1.54E-01 |
| WBC count | rs6721033 | 2 | 113908086 | a | C | 0.3987 | 0.0045 | 0.002 | $2.38 \mathrm{E}-02$ | +++ | $2.87 \mathrm{E}-01$ |
| WBC count | rs6722922 | 2 | 113841515 | t | C | 0.4936 | 0.0039 | 0.0022 | $6.88 \mathrm{E}-02$ | ++- | $4.71 \mathrm{E}-04$ |
| WBC count | rs6723495 | 2 | 113922579 | t | c | 0.7273 | 0.001 | 0.0043 | 8.23E-01 | +?+ | $4.54 \mathrm{E}-01$ |
| WBC count | rs6724667 | 2 | 113789641 | a | g | 0.5307 | -0.0011 | 0.0025 | 6.53E-01 | -+- | $4.42 \mathrm{E}-01$ |
| WBC count | rs6728590 | 2 | 113844600 | a | g | 0.5257 | -0.0042 | 0.0018 | 2.19E-02 | +-- | 7.79E-03 |
| WBC count | rs6730516 | 2 | 113848961 | t | c | 0.4754 | 0.0071 | 0.0022 | 9.84E-04 | ++- | 9.52E-03 |
| WBC count | rs6731551 | 2 | 113844553 | t | C | 0.6277 | 0.0013 | 0.0019 | 4.90E-01 | +-+ | $1.74 \mathrm{E}-04$ |
| WBC count | rs6734238 | 2 | 113841030 | a | $g$ | 0.5182 | -0.0117 | 0.0021 | 2.37E-08 | --- | 2.09E-01 |
| WBC count | rs6738239 | 2 | 113850792 | a | c | 0.3446 | 0.0106 | 0.0021 | 3.63E-07 | +++ | $2.39 \mathrm{E}-01$ |
| WBC count | rs6738377 | 2 | 113850902 | C | g | 0.5289 | -0.004 | 0.0018 | 2.98E-02 | +-- | $1.21 \mathrm{E}-02$ |
| WBC count | rs6739883 | 2 | 113914312 | a | g | 0.1336 | 0.0022 | 0.0032 | 4.89E-01 | ++- | $6.48 \mathrm{E}-01$ |
| WBC count | rs6741180 | 2 | 113844384 | a | g | 0.3445 | 0.0102 | 0.0021 | 8.42E-07 | +++ | $2.48 \mathrm{E}-01$ |
| WBC count | rs6743171 | 2 | 113840058 | c | g | 0.3168 | 0.0079 | 0.0022 | 2.72E-04 | +++ | $5.69 \mathrm{E}-02$ |
| WBC count | rs6743376 | 2 | 113832333 | a | c | 0.7371 | 0.0122 | 0.0026 | $1.75 \mathrm{E}-06$ | +?+ | 5.76E-01 |
| WBC count | rs6744288 | 2 | 113783230 | a | t | 0.5023 | $4.00 \mathrm{E}-04$ | 0.0018 | 8.11E-01 | +-- | $4.79 \mathrm{E}-02$ |
| WBC count | rs6744874 | 2 | 113783909 | a | t | 0.1543 | -0.0032 | 0.0037 | 3.89E-01 | -?- | $6.47 \mathrm{E}-01$ |
| WBC count | rs6746979 | 2 | 113854120 | a | t | 0.335 | 0.0106 | 0.0021 | $5.59 \mathrm{E}-07$ | +++ | $2.39 \mathrm{E}-01$ |
| WBC count | rs6750559 | 2 | 113841532 | a | g | 0.3589 | 0.0104 | 0.0021 | 9.57E-07 | +++ | $3.94 \mathrm{E}-01$ |
| WBC count | rs6754298 | 2 | 113896213 | a | t | 0.3063 | 0.0047 | 0.0021 | 2.80E-02 | +++ | 7.59E-01 |
| WBC count | rs6754538 | 2 | 113777393 | C | g | 0.3001 | 0.0066 | 0.0039 | 9.15E-02 | ??+ | $1.00 \mathrm{E}+00$ |

Table S1. Random-effects meta-analysis of MANTRA identified loci.

| WBC count | rs6755354 | 2 | 113826072 | t | g | 0.4443 | 0.002 | 0.0021 | $3.48 \mathrm{E}-01$ | +-- | 3.73E-01 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| WBC count | rs6756735 | 2 | 113806121 | t | c | 0.3324 | -0.0082 | 0.0031 | 7.73E-03 | -?- | 3.66E-01 |
| WBC count | rs6758965 | 2 | 113791193 | t | C | 0.638 | 3.00E-04 | 0.0024 | $9.02 \mathrm{E}-01$ | +?- | 1.21E-02 |
| WBC count | rs6759676 | 2 | 113836348 | t | C | 0.3562 | 0.0034 | 0.002 | $9.42 \mathrm{E}-02$ | ++- | 4.01E-02 |
| WBC count | rs6760120 | 2 | 113932431 | t | C | 0.4236 | -2.00E-04 | 0.0019 | 9.13E-01 | --+ | 8.65E-01 |
| WBC count | rs6761276 | 2 | 113832312 | t | C | 0.5895 | -0.0027 | 0.002 | $1.74 \mathrm{E}-01$ | --+ | 2.56E-02 |
| WBC count | rs724496 | 2 | 113922276 | a | g | 0.4256 | 2.00E-04 | 0.0027 | $9.28 \mathrm{E}-01$ | --+ | $5.29 \mathrm{E}-01$ |
| WBC count | rs7557928 | 2 | 113797968 | t | g | 0.445 | 5.00E-04 | 0.0023 | $8.44 \mathrm{E}-01$ | +-+ | 7.45E-01 |
| WBC count | rs7558672 | 2 | 113825303 | a | g | 0.1901 | -0.0085 | 0.0032 | 7.14E-03 | -?- | 4.79E-01 |
| WBC count | rs7561080 | 2 | 113864920 | a | g | 0.6021 | 0.0054 | 0.0023 | $1.94 \mathrm{E}-02$ | +++ | $2.88 \mathrm{E}-01$ |
| WBC count | rs7569284 | 2 | 113803563 | a | g | 0.5537 | 0.0011 | 0.0023 | $6.47 \mathrm{E}-01$ | -++ | 8.43E-01 |
| WBC count | rs7570267 | 2 | 113828578 | a | g | 0.627 | 0.0029 | 0.002 | $1.52 \mathrm{E}-01$ | -++ | 1.95E-01 |
| WBC count | rs7571656 | 2 | 113758863 | t | g | 0.5891 | -6.00E-04 | 0.002 | $7.40 \mathrm{E}-01$ | +-+ | 7.33E-02 |
| WBC count | rs7573950 | 2 | 113859645 | a | g | 0.101 | 0.0234 | 0.009 | 8.96E-03 | ??+ | $1.00 \mathrm{E}+00$ |
| WBC count | rs7574159 | 2 | 113859761 | a | g | 0.3351 | 0.0101 | 0.0021 | 2.69E-06 | +++ | 2.61E-01 |
| WBC count | rs7574427 | 2 | 113860038 | a | g | 0.3084 | 0.0085 | 0.0022 | $1.22 \mathrm{E}-04$ | +++ | $1.31 \mathrm{E}-01$ |
| WBC count | rs7574787 | 2 | 113837416 | t | g | 0.624 | 0.009 | 0.0055 | $9.90 \mathrm{E}-02$ | -?+ | $3.14 \mathrm{E}-02$ |
| WBC count | rs7575934 | 2 | 113817334 | t | c | 0.4188 | -0.0069 | 0.0027 | $1.21 \mathrm{E}-02$ | -?- | 1.94E-01 |
| WBC count | rs7583835 | 2 | 113754725 | a | C | 0.0463 | -0.0078 | 0.0136 | $5.66 \mathrm{E}-01$ | ??- | $1.00 \mathrm{E}+00$ |
| WBC count | rs7584409 | 2 | 113742734 | a | g | 0.4811 | -5.00E-04 | 0.0025 | $8.48 \mathrm{E}-01$ | ?-+ | 8.32E-01 |
| WBC count | rs7593760 | 2 | 113753731 | a | g | 0.9537 | 0.0077 | 0.0136 | $5.71 \mathrm{E}-01$ | ??+ | $1.00 \mathrm{E}+00$ |
| WBC count | rs7595962 | 2 | 113934226 | t | C | 0.516 | -9.00E-04 | 0.0021 | $6.56 \mathrm{E}-01$ | -+- | $2.26 \mathrm{E}-01$ |
| WBC count | rs7599662 | 2 | 113825085 | t | C | 0.3882 | -4.00E-04 | 0.0025 | $8.74 \mathrm{E}-01$ | -+- | 9.44E-01 |
| WBC count | rs7606584 | 2 | 113796484 | t | C | 0.186 | -6.00E-04 | 0.007 | $9.32 \mathrm{E}-01$ | ??- | $1.00 \mathrm{E}+00$ |
| WBC count | rs768627 | 2 | 113821653 | t | C | 0.4295 | -0.0014 | 0.0018 | 4.35E-01 | --+ | $1.37 \mathrm{E}-03$ |
| WBC count | rs895495 | 2 | 113899758 | a | g | 0.464 | -0.0038 | 0.002 | $6.06 \mathrm{E}-02$ | --+ | 2.16E-01 |
| WBC count | rs895496 | 2 | 113899025 | t | C | 0.3891 | 0.0024 | 0.002 | $2.45 \mathrm{E}-01$ | ++- | 6.32E-02 |
| WBC count | rs895497 | 2 | 113763575 | a | g | 0.2718 | 0.004 | 0.0026 | $1.23 \mathrm{E}-01$ | ?++ | 8.67E-02 |
| WBC count | rs9005 | 2 | 113891412 | a | g | 0.2583 | -0.0033 | 0.002 | $9.80 \mathrm{E}-02$ | --- | 8.09E-01 |
| WBC count | rs921065 | 2 | 113825966 | t | c | 0.4131 | -3.00E-04 | 0.0018 | 8.47E-01 | --+ | $1.02 \mathrm{E}-02$ |
| WBC count | rs928940 | 2 | 113877495 | t | g | 0.3068 | 2.00E-04 | 0.002 | 9.23E-01 | -++ | $3.00 \mathrm{E}-01$ |
| WBC count | rs9308681 | 2 | 113781335 | C | g | 0.5484 | 6.00E-04 | 0.0024 | 8.13E-01 | +-- | $6.80 \mathrm{E}-01$ |
| WBC count | rs9308682 | 2 | 113828425 | a | g | 0.3779 | 0.0027 | 0.0022 | $2.11 \mathrm{E}-01$ | +-+ | $1.86 \mathrm{E}-01$ |
| WBC count | rs931471 | 2 | 113912693 | t | C | 0.3665 | 0.0051 | 0.0021 | $1.40 \mathrm{E}-02$ | +++ | 3.25E-01 |
| WBC count | rs957200 | 2 | 113821336 | a | C | 0.182 | 0.0022 | 0.0026 | 3.85E-01 | +++ | 4.66E-01 |
| WBC count | rs957201 | 2 | 113821200 | t | C | 0.4283 | -0.0014 | 0.0018 | 4.51E-01 | --+ | $1.20 \mathrm{E}-03$ |
| WBC count | rs9678578 | 2 | 113810947 | a | t | 0.5205 | -2.00E-04 | 0.0025 | $9.27 \mathrm{E}-01$ | -0- | 9.82E-01 |
| WBC count | rs973635 | 2 | 113889134 | a | g | 0.4377 | -0.0026 | 0.002 | 1.93E-01 | +-- | 3.69E-01 |

Table S1. Random-effects meta-analysis of MANTRA identified loci.

| WBC count | rs990524 | 2 | 113816481 | t | c | 0.363 | 2.00E-04 | 0.0025 | 9.41E-01 | -0+ | 9.73E-01 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| WBC count | rs996879 | 2 | 113822241 | a | g | 0.2204 | 0.0031 | 0.0032 | 3.41E-01 | +?+ | $2.44 \mathrm{E}-01$ |
| WBC count | rs9973741 | 2 | 113853381 | a | g | 0.7481 | -0.0131 | 0.0034 | $1.45 \mathrm{E}-04$ | -?- | $2.20 \mathrm{E}-01$ |
| WBC count | rs9973878 | 2 | 113752761 | a | g | 0.1357 | 0.0052 | 0.0052 | 3.22E-01 | ??+ | $1.00 \mathrm{E}+00$ |
| WBC count | rs10169718 | 2 | 219103580 | a | g | 0.5134 | -0.0098 | 0.0018 | 7.67E-08 | --+ | 7.59E-02 |
| WBC count | rs10182823 | 2 | 219107367 | a | g | 0.8973 | -4.00E-04 | 0.0083 | $9.62 \mathrm{E}-01$ | ??- | $1.00 \mathrm{E}+00$ |
| WBC count | rs1018288 | 2 | 219099260 | c | g | 0.7837 | 5.00E-04 | 0.0065 | 9.39E-01 | ??+ | $1.00 \mathrm{E}+00$ |
| WBC count | rs10932765 | 2 | 219099484 | $t$ | c | 0.4251 | -0.0106 | 0.0019 | 9.42E-09 | --- | 4.55E-01 |
| WBC count | rs11901011 | 2 | 219106136 | a | g | 0.0217 | 0.0149 | 0.0199 | 4.53E-01 | ??+ | $1.00 \mathrm{E}+00$ |
| WBC count | rs16858888 | 2 | 219090451 | t | c | 0.0885 | -0.0128 | 0.0094 | 1.72E-01 | ??- | $1.00 \mathrm{E}+00$ |
| WBC count | rs17572109 | 2 | 219093934 | a | g | 0.2382 | -0.0068 | 0.0032 | 3.37E-02 | ?-- | 2.69E-01 |
| WBC count | rs4674275 | 2 | 219097738 | a | g | 0.4084 | -0.0087 | 0.0018 | $9.10 \mathrm{E}-07$ | --+ | 2.96E-02 |
| WBC count | rs6436046 | 2 | 219098248 | t | g | 0.0184 | -0.0236 | 0.0227 | 2.99E-01 | ??- | $1.00 \mathrm{E}+00$ |
| WBC count | rs6436047 | 2 | 219108913 | a | C | 0.4046 | -0.0091 | 0.0018 | 6.59E-07 | --+ | 2.62E-02 |
| WBC count | rs6436048 | 2 | 219109033 | a | g | 0.5661 | 0.0087 | 0.0018 | 8.77E-07 | ++- | 3.15E-02 |
| WBC count | rs6704974 | 2 | 219109391 | a | g | 0.1313 | -0.0024 | 0.0076 | 7.53E-01 | ??- | $1.00 \mathrm{E}+00$ |
| WBC count | rs6742439 | 2 | 219098672 | a | g | 0.2163 | -5.00E-04 | 0.0065 | $9.39 \mathrm{E}-01$ | ??- | $1.00 \mathrm{E}+00$ |
| WBC count | rs6757318 | 2 | 219090422 | a | g | 0.784 | 8.00E-04 | 0.0065 | 9.02E-01 | ??+ | $1.00 \mathrm{E}+00$ |
| WBC count | rs7605980 | 2 | 219100869 | C | g | 0.5673 | 0.0087 | 0.0018 | $8.74 \mathrm{E}-07$ | ++- | 3.15E-02 |
| WBC count | rs9288536 | 2 | 219089616 | C | g | 0.1232 | -0.0011 | 0.0056 | 8.43E-01 | ??- | $1.00 \mathrm{E}+00$ |
| WBC count | rs9288537 | 2 | 219096333 | c | g | 0.7003 | -0.0024 | 0.004 | 5.50E-01 | ??- | $1.00 \mathrm{E}+00$ |
| WBC count | rs9679606 | 2 | 219107700 | a | g | 0.6942 | -0.0017 | 0.0059 | 7.72E-01 | ??- | $1.00 \mathrm{E}+00$ |
| WBC count | rs9789363 | 2 | 219090365 | t | c | 0.6969 | -0.0015 | 0.0039 | 7.01E-01 | ??- | $1.00 \mathrm{E}+00$ |
| WBC count | rs10456057 | 6 | 31245534 | a | g | 0.4735 | -0.0028 | 0.0031 | $3.66 \mathrm{E}-01$ | --+ | $5.71 \mathrm{E}-01$ |
| WBC count | rs12111032 | 6 | 31242191 | a | g | 0.4782 | 0.0054 | 0.0027 | $4.95 \mathrm{E}-02$ | ?++ | 3.92E-01 |
| WBC count | rs12191877 | 6 | 31252925 | t | c | 0.4245 | 0.0039 | 0.0031 | 2.07E-01 | +++ | 5.68E-01 |
| WBC count | rs12234123 | 6 | 31243604 | a | g | 0.2245 | -0.0045 | 0.0028 | 1.10E-01 | -++ | 2.07E-02 |
| WBC count | rs12529015 | 6 | 31248262 | a | g | 0.1416 | -0.0058 | 0.0031 | $6.47 \mathrm{E}-02$ | -++ | 7.21E-02 |
| WBC count | rs13191343 | 6 | 31241109 | t | c | 0.3155 | 0.0054 | 0.0026 | $3.90 \mathrm{E}-02$ | +++ | $6.08 \mathrm{E}-01$ |
| WBC count | rs13200569 | 6 | 31243615 | a | g | 0.3146 | -0.0033 | 0.0019 | 7.63E-02 | -+- | 2.09E-01 |
| WBC count | rs13203895 | 6 | 31244082 | t | c | 0.4222 | 0.004 | 0.0031 | $1.94 \mathrm{E}-01$ | +++ | 9.02E-01 |
| WBC count | rs13207315 | 6 | 31241127 | t | c | 0.4271 | -0.0034 | 0.0026 | 1.97E-01 | --+ | $3.25 \mathrm{E}-01$ |
| WBC count | rs13208617 | 6 | 31248819 | t | C | 0.4217 | 0.003 | 0.0031 | 3.26E-01 | ++- | 6.05E-01 |
| WBC count | rs16899147 | 6 | 31255201 | a | g | 0.6685 | 0.0027 | 0.0026 | 3.07E-01 | +-- | 6.92E-02 |
| WBC count | rs16899160 | 6 | 31256667 | a | g | 0.0655 | -0.0125 | 0.0046 | 6.92E-03 | ?-- | 7.63E-01 |
| WBC count | rs2040748 | 6 | 31243785 | t | g | 0.3754 | 0.0042 | 0.002 | 3.91E-02 | ++- | 2.82E-04 |
| WBC count | rs2074488 | 6 | 31240431 | t | g | 0.3562 | -0.0055 | 0.0024 | 2.01E-02 | --+ | 1.52E-02 |
| WBC count | rs2074489 | 6 | 31240128 | t | C | 0.4071 | -0.0023 | 0.0019 | $2.40 \mathrm{E}-01$ | -+- | 2.03E-02 |

Table S1. Random-effects meta-analysis of MANTRA identified loci.

| WBC count | rs2074491 | 6 | 31239896 | t | c | 0.8493 | 0.003 | 0.0098 | 7.64E-01 | +?+ | 9.12E-01 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| WBC count | rs2249741 | 6 | 31240712 | a | c | 0.3549 | -0.0105 | 0.0041 | $1.08 \mathrm{E}-02$ | ??- | $1.00 \mathrm{E}+00$ |
| WBC count | rs2249742 | 6 | 31240721 | t | C | 0.5109 | 0.0051 | 0.0017 | 3.17E-03 | ++- | 2.37E-08 |
| WBC count | rs2394963 | 6 | 31251462 | t | C | 0.4608 | -0.0018 | 0.0018 | 3.31E-01 | -+- | 4.27E-02 |
| WBC count | rs2395471 | 6 | 31240692 | a | g | 0.4747 | 0.0033 | 0.0017 | 4.92E-02 | -++ | 8.88E-02 |
| WBC count | rs2524042 | 6 | 31257160 | c | g | 0.5922 | -0.012 | 0.0024 | $4.98 \mathrm{E}-07$ | --- | $4.31 \mathrm{E}-01$ |
| WBC count | rs2524043 | 6 | 31257012 | a | g | 0.5902 | -0.0175 | 0.0028 | $8.24 \mathrm{E}-10$ | --- | $1.51 \mathrm{E}-01$ |
| WBC count | rs2524044 | 6 | 31256753 | t | g | 0.4573 | -0.0083 | 0.0024 | 4.50E-04 | --+ | 6.39E-03 |
| WBC count | rs2524047 | 6 | 31256591 | a | g | 0.5906 | -0.0174 | 0.0028 | 8.57E-10 | --- | $1.47 \mathrm{E}-01$ |
| WBC count | rs2524048 | 6 | 31256561 | t | C | 0.3832 | -0.0154 | 0.0028 | 5.89E-08 | --+ | $2.40 \mathrm{E}-03$ |
| WBC count | rs2524049 | 6 | 31255953 | c | g | 0.5814 | -0.0137 | 0.0025 | $2.18 \mathrm{E}-08$ | --- | $3.20 \mathrm{E}-01$ |
| WBC count | rs2524050 | 6 | 31255541 | t | C | 0.4065 | -0.0132 | 0.003 | 8.99E-06 | --+ | 6.70E-03 |
| WBC count | rs2524051 | 6 | 31255500 | a | g | 0.5907 | -0.0175 | 0.0028 | 8.41E-10 | --- | $1.47 \mathrm{E}-01$ |
| WBC count | rs2524052 | 6 | 31253996 | a | g | 0.3813 | -0.0055 | 0.0019 | 4.33E-03 | +-- | 7.24E-03 |
| WBC count | rs2524053 | 6 | 31252469 | t | g | 0.4984 | -0.0025 | 0.0019 | 2.02E-01 | +-+ | $3.34 \mathrm{E}-04$ |
| WBC count | rs2524054 | 6 | 31252396 | a | C | 0.2011 | -0.0077 | 0.0022 | 4.03E-04 | +- | 5.57E-05 |
| WBC count | rs2524055 | 6 | 31252237 | a | C | 0.5899 | -0.0172 | 0.0029 | $1.70 \mathrm{E}-09$ | --- | 9.47E-02 |
| WBC count | rs2524057 | 6 | 31251895 | a | g | 0.5922 | -0.012 | 0.0024 | 5.58E-07 | --- | 4.33E-01 |
| WBC count | rs2524059 | 6 | 31250823 | C | g | 0.2066 | -0.0024 | 0.0028 | $3.99 \mathrm{E}-01$ | +?- | 7.55E-02 |
| WBC count | rs2524063 | 6 | 31248155 | a | g | 0.1072 | 0.0173 | 0.0028 | 4.05E-10 | +++ | 2.11E-01 |
| WBC count | rs2524067 | 6 | 31245821 | a | g | 0.4562 | 0.0121 | 0.0031 | $7.70 \mathrm{E}-05$ | ?++ | 6.02E-01 |
| WBC count | rs2524068 | 6 | 31245061 | t | g | 0.3804 | 0.0083 | 0.0026 | $1.38 \mathrm{E}-03$ | 0 | $1.41 \mathrm{E}-02$ |
| WBC count | rs2524069 | 6 | 31244789 | a | t | 0.4054 | 0.0135 | 0.0033 | 3.59E-05 | ?++ | 8.67E-01 |
| WBC count | rs2524070 | 6 | 31244520 | a | g | 0.1551 | 0.0101 | 0.0024 | 3.69E-05 | ++- | 8.59E-03 |
| WBC count | rs2524074 | 6 | 31244021 | a | g | 0.5899 | 0.0088 | 0.0021 | 2.82E-05 | -++ | 2.32E-03 |
| WBC count | rs2524077 | 6 | 31243603 | t | c | 0.3468 | 0.0133 | 0.0028 | 1.42E-06 | ++- | $6.74 \mathrm{E}-05$ |
| WBC count | rs2524078 | 6 | 31242649 | a | g | 0.45 | 0.0112 | 0.003 | 2.20E-04 | ?++ | 9.03E-01 |
| WBC count | rs2524079 | 6 | 31242174 | a | g | 0.4084 | 0.0121 | 0.0019 | $1.97 \mathrm{E}-10$ | +++ | 2.92E-01 |
| WBC count | rs2524082 | 6 | 31241761 | a | t | 0.4934 | 0.0074 | 0.0017 | 1.63E-05 | +++ | $1.24 \mathrm{E}-01$ |
| WBC count | rs2524083 | 6 | 31241737 | t | c | 0.3452 | 0.0133 | 0.0028 | $2.66 \mathrm{E}-06$ | ++- | $6.70 \mathrm{E}-05$ |
| WBC count | rs2524084 | 6 | 31241639 | a | g | 0.478 | 0.0107 | 0.0018 | 2.53E-09 | +++ | 3.43E-01 |
| WBC count | rs2524094 | 6 | 31240041 | t | c | 0.5217 | 0.0033 | 0.0019 | 7.91E-02 | -++ | 1.63E-02 |
| WBC count | rs2844599 | 6 | 31256005 | C | g | 0.102 | 0.017 | 0.0028 | 1.85E-09 | +++ | 8.32E-02 |
| WBC count | rs2844600 | 6 | 31253771 | t | g | 0.3158 | 0.015 | 0.0028 | $1.21 \mathrm{E}-07$ | ++- | 1.15E-03 |
| WBC count | rs2844603 | 6 | 31250854 | a | g | 0.2085 | -0.0026 | 0.0028 | 3.51E-01 | +?- | 6.60E-02 |
| WBC count | rs2844605 | 6 | 31248591 | a | c | 0.4656 | -0.0083 | 0.0017 | 1.13E-06 | --- | $1.80 \mathrm{E}-01$ |
| WBC count | rs2844611 | 6 | 31243979 | a | t | 0.5905 | 0.0088 | 0.0021 | 3.07E-05 | -++ | 2.08E-03 |
| WBC count | rs2844612 | 6 | 31243971 | a | g | 0.2291 | -0.0086 | 0.0021 | 4.46E-05 | +-- | 2.29E-03 |

Table S1. Random-effects meta-analysis of MANTRA identified loci.

| WBC count | rs2844613 | 6 | 31243846 | t | g | 0.4509 | -0.0067 | 0.0031 | 2.92E-02 | ?-+ | $2.01 \mathrm{E}-03$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| WBC count | rs2844615 | 6 | 31242959 | t | c | 0.3368 | 0.0135 | 0.0028 | $8.59 \mathrm{E}-07$ | ++- | $1.76 \mathrm{E}-04$ |
| WBC count | rs2844618 | 6 | 31242439 | t | C | 0.3377 | 0.0137 | 0.0028 | $6.38 \mathrm{E}-07$ | ++- | 2.22E-04 |
| WBC count | rs2844619 | 6 | 31242223 | C | g | 0.31 | -0.0097 | 0.002 | $1.89 \mathrm{E}-06$ | +-- | $1.55 \mathrm{E}-04$ |
| WBC count | rs2844621 | 6 | 31241537 | t | C | 0.4445 | -0.0051 | 0.002 | 9.73E-03 | --+ | $1.95 \mathrm{E}-02$ |
| WBC count | rs2844622 | 6 | 31240060 | a | g | 0.2562 | -0.0071 | 0.0021 | 5.14E-04 | +-- | 3.43E-03 |
| WBC count | rs28481932 | 6 | 31242900 | a | g | 0.625 | 7.00E-04 | 0.0023 | 7.53E-01 | +?- | $9.09 \mathrm{E}-03$ |
| WBC count | rs2853929 | 6 | 31255434 | t | c | 0.3101 | 0.0155 | 0.0029 | $5.99 \mathrm{E}-08$ | ++- | 2.39E-03 |
| WBC count | rs2853930 | 6 | 31255424 | a | C | 0.6214 | -0.0153 | 0.003 | $2.35 \mathrm{E}-07$ | --- | 2.23E-01 |
| WBC count | rs2853933 | 6 | 31254088 | t | C | 0.4906 | -0.0012 | 0.0021 | $5.74 \mathrm{E}-01$ | +-+ | 2.94E-03 |
| WBC count | rs2853934 | 6 | 31253928 | a | t | 0.5945 | -0.0173 | 0.0028 | $8.77 \mathrm{E}-10$ | --- | $1.46 \mathrm{E}-01$ |
| WBC count | rs2853935 | 6 | 31253878 | t | C | 0.576 | -0.0059 | 0.0027 | $2.95 \mathrm{E}-02$ | -?- | 5.02E-01 |
| WBC count | rs2853937 | 6 | 31252569 | t | C | 0.3787 | -0.0156 | 0.0028 | 3.36E-08 | --+ | 4.92E-03 |
| WBC count | rs2853939 | 6 | 31250642 | t | C | 0.4989 | -0.003 | 0.0019 | $1.22 \mathrm{E}-01$ | +-+ | 1.16E-03 |
| WBC count | rs2853943 | 6 | 31247871 | t | C | 0.4963 | 0.0021 | 0.0024 | 3.84E-01 | -?+ | $1.49 \mathrm{E}-01$ |
| WBC count | rs2853946 | 6 | 31247203 | a | $t$ | 0.5198 | -0.0139 | 0.0018 | 2.67E-14 | --- | 4.37E-01 |
| WBC count | rs2853948 | 6 | 31245573 | C | g | 0.1071 | 0.0176 | 0.0027 | $1.66 \mathrm{E}-10$ | +++ | 3.29E-01 |
| WBC count | rs3130693 | 6 | 31242859 | t | c | 0.5511 | -0.0021 | 0.0026 | $4.30 \mathrm{E}-01$ | -?- | 8.13E-01 |
| WBC count | rs3130696 | 6 | 31243884 | a | g | 0.2691 | -0.0074 | 0.002 | 1.85E-04 | --- | $6.18 \mathrm{E}-01$ |
| WBC count | rs3132485 | 6 | 31243389 | a | C | 0.4261 | -9.00E-04 | 0.0017 | 5.94E-01 | -0- | $6.06 \mathrm{E}-01$ |
| WBC count | rs3132486 | 6 | 31243170 | a | g | 0.5518 | -0.003 | 0.0018 | $9.26 \mathrm{E}-02$ | -+- | 1.16E-01 |
| WBC count | rs3132488 | 6 | 31242695 | t | g | 0.6175 | -0.004 | 0.0033 | 2.27E-01 | -?- | 7.01E-01 |
| WBC count | rs3132489 | 6 | 31242674 | t | C | 0.5637 | -5.00E-04 | 0.0019 | $7.90 \mathrm{E}-01$ | -++ | $3.97 \mathrm{E}-01$ |
| WBC count | rs3132490 | 6 | 31242560 | a | g | 0.6104 | 0.0037 | 0.0025 | $1.37 \mathrm{E}-01$ | -++ | 2.37E-02 |
| WBC count | rs3132491 | 6 | 31242272 | t | c | 0.4201 | 0.0048 | 0.0025 | 5.85E-02 | -++ | 3.05E-02 |
| WBC count | rs3134745 | 6 | 31242762 | t | C | 0.4455 | 0.0088 | 0.0033 | $8.54 \mathrm{E}-03$ | +?- | 2.61E-03 |
| WBC count | rs3873374 | 6 | 31251311 | t | C | 0.4898 | 0.0045 | 0.0022 | $4.30 \mathrm{E}-02$ | +?+ | $3.28 \mathrm{E}-01$ |
| WBC count | rs3873375 | 6 | 31251360 | t | C | 0.4643 | -0.0019 | 0.0018 | 2.85E-01 | -+- | $1.04 \mathrm{E}-01$ |
| WBC count | rs3930575 | 6 | 31244654 | a | g | 0.1303 | -0.0047 | 0.0026 | 7.03E-02 | +-- | 4.57E-02 |
| WBC count | rs3931670 | 6 | 31243767 | C | g | 0.2886 | 0.0061 | 0.0019 | $1.69 \mathrm{E}-03$ | +++ | $6.06 \mathrm{E}-01$ |
| WBC count | rs4361609 | 6 | 31240635 | C | g | 0.1906 | 0.0044 | 0.0022 | 4.89E-02 | -++ | $9.48 \mathrm{E}-05$ |
| WBC count | rs4386816 | 6 | 31247135 | t | c | 0.4409 | -0.0042 | 0.0024 | 7.41E-02 | --+ | $2.68 \mathrm{E}-05$ |
| WBC count | rs4628144 | 6 | 31257104 | t | c | 0.4329 | 0.0017 | 0.0018 | $3.27 \mathrm{E}-01$ | +-+ | $8.48 \mathrm{E}-02$ |
| WBC count | rs5009853 | 6 | 31240636 | t | C | 0.2671 | -0.0031 | 0.0048 | 5.22E-01 | -+- | 8.67E-01 |
| WBC count | rs5010528 | 6 | 31241032 | a | g | 0.5863 | -0.0113 | 0.0028 | 6.92E-05 | --- | $7.80 \mathrm{E}-01$ |
| WBC count | rs6457358 | 6 | 31239999 | a | t | 0.5449 | 0.0081 | 0.0039 | 3.82E-02 | +++ | $8.45 \mathrm{E}-01$ |
| WBC count | rs6457372 | 6 | 31247121 | a | g | 0.4975 | -0.0044 | 0.0017 | 1.11E-02 | --- | 9.63E-01 |
| WBC count | rs6906846 | 6 | 31245736 | a | g | 0.3114 | 0.0022 | 0.0018 | 2.30E-01 | +-+ | 4.04E-02 |

Table S1. Random-effects meta-analysis of MANTRA identified loci.

| WBC count | rs6913377 | 6 | 31243495 | a | g | 0.0711 | -7.00E-04 | 0.0045 | $8.78 \mathrm{E}-01$ | ?-+ | 7.11E-01 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| WBC count | rs6919409 | 6 | 31244532 | a | g | 0.0677 | -0.0109 | 0.0045 | $1.61 \mathrm{E}-02$ | ?-- | 9.82E-01 |
| WBC count | rs6919908 | 6 | 31244960 | t | c | 0.4206 | 0.0033 | 0.0024 | $1.75 \mathrm{E}-01$ | +?- | $3.06 \mathrm{E}-02$ |
| WBC count | rs6923313 | 6 | 31241370 | t | C | 0.5801 | -0.0067 | 0.0024 | 5.60E-03 | -?- | $6.36 \mathrm{E}-01$ |
| WBC count | rs7381988 | 6 | 31246703 | a | g | 0.5936 | 0.0044 | 0.0025 | 7.92E-02 | -++ | $9.90 \mathrm{E}-03$ |
| WBC count | rs7382297 | 6 | 31247067 | t | g | 0.3092 | -0.001 | 0.0027 | 7.19E-01 | +-+ | 1.23E-01 |
| WBC count | rs7382307 | 6 | 31247149 | a | g | 0.1298 | -0.0043 | 0.0026 | $9.09 \mathrm{E}-02$ | +-- | $4.59 \mathrm{E}-02$ |
| WBC count | rs7754443 | 6 | 31254263 | a | g | 0.4021 | 0.0018 | 0.0018 | 3.22E-01 | -++ | 7.98E-02 |
| WBC count | rs7755686 | 6 | 31247796 | C | g | 0.114 | -1.00E-04 | 0.0036 | $9.86 \mathrm{E}-01$ | --+ | $3.68 \mathrm{E}-01$ |
| WBC count | rs7759127 | 6 | 31240988 | t | g | 0.4004 | -0.011 | 0.0029 | 1.37E-04 | --+ | 5.01E-04 |
| WBC count | rs7767581 | 6 | 31239855 | C | g | 0.248 | -0.0074 | 0.0021 | 3.73E-04 | +- | $1.98 \mathrm{E}-04$ |
| WBC count | rs7773175 | 6 | 31240959 | C | g | 0.585 | -0.0119 | 0.002 | 1.96E-09 | --- | $1.75 \mathrm{E}-01$ |
| WBC count | rs9264602 | 6 | 31237405 | t | C | 0.5731 | -0.0066 | 0.003 | $2.90 \mathrm{E}-02$ | -?- | $6.07 \mathrm{E}-01$ |
| WBC count | rs9264603 | 6 | 31237437 | t | c | 0.4216 | 0.0051 | 0.0028 | $7.38 \mathrm{E}-02$ | +?+ | $5.29 \mathrm{E}-01$ |
| WBC count | rs9264604 | 6 | 31237449 | t | C | 0.4344 | 0.0049 | 0.003 | $1.06 \mathrm{E}-01$ | +?+ | $6.36 \mathrm{E}-01$ |
| WBC count | rs9264606 | 6 | 31237592 | a | C | 0.2397 | -0.0075 | 0.0021 | 3.40E-04 | +-- | $1.51 \mathrm{E}-04$ |
| WBC count | rs9264608 | 6 | 31237660 | a | g | 0.2553 | -0.0082 | 0.0021 | 6.72E-05 | +-- | 7.19E-04 |
| WBC count | rs9264636 | 6 | 31238297 | t | c | 0.4458 | 0.0046 | 0.002 | $2.48 \mathrm{E}-02$ | -+- | $1.50 \mathrm{E}-06$ |
| WBC count | rs9264647 | 6 | 31238613 | a | g | 0.2419 | -0.0082 | 0.0021 | 8.33E-05 | +-- | 2.05E-04 |
| WBC count | rs9264664 | 6 | 31239227 | t | c | 0.4264 | 0.0045 | 0.0021 | 3.54E-02 | -+- | 5.31E-07 |
| WBC count | rs9264672 | 6 | 31239766 | t | c | 0.3186 | 0.0143 | 0.0028 | 3.45E-07 | ++- | $3.74 \mathrm{E}-04$ |
| WBC count | rs9264679 | 6 | 31240663 | a | t | 0.5891 | 0.0044 | 0.0026 | 8.65E-02 | -++ | $3.80 \mathrm{E}-02$ |
| WBC count | rs9264681 | 6 | 31240871 | c | g | 0.305 | 0.0025 | 0.0019 | 1.83E-01 | -+- | $1.36 \mathrm{E}-01$ |
| WBC count | rs9264731 | 6 | 31244068 | a | g | 0.5917 | 0.0046 | 0.0026 | 7.19E-02 | -++ | $4.50 \mathrm{E}-02$ |
| WBC count | rs9264733 | 6 | 31244214 | C | g | 0.2797 | 0.0083 | 0.0025 | 7.66E-04 | +?+ | 5.97E-01 |
| WBC count | rs9264742 | 6 | 31244616 | t | c | 0.4047 | -0.0087 | 0.0037 | $2.06 \mathrm{E}-02$ | ?-+ | 3.16E-02 |
| WBC count | rs9264745 | 6 | 31245478 | a | g | 0.6158 | 0.0037 | 0.0024 | $1.24 \mathrm{E}-01$ | -++ | 2.92E-02 |
| WBC count | rs9264759 | 6 | 31248678 | t | c | 0.4067 | -0.0029 | 0.002 | $1.51 \mathrm{E}-01$ | +-- | 5.88E-02 |
| WBC count | rs9348860 | 6 | 31252667 | t | c | 0.4748 | 0.0041 | 0.0026 | $1.21 \mathrm{E}-01$ | +-+ | $4.08 \mathrm{E}-02$ |
| WBC count | rs9348861 | 6 | 31252722 | a | g | 0.6626 | 0.0044 | 0.0026 | $9.17 \mathrm{E}-02$ | +-+ | $4.41 \mathrm{E}-02$ |
| WBC count | rs9348862 | 6 | 31252747 | t | C | 0.0626 | -0.0045 | 0.0048 | 3.43E-01 | ?-- | 6.17E-01 |
| WBC count | rs9366776 | 6 | 31256630 | a | g | 0.1425 | -0.0031 | 0.0026 | $2.27 \mathrm{E}-01$ | -+- | $1.12 \mathrm{E}-02$ |
| WBC count | rs9368669 | 6 | 31248493 |  | c | 0.6627 | 0.0059 | 0.0032 | $6.68 \mathrm{E}-02$ | +-- | 6.82E-02 |
| WBC count | rs9368670 | 6 | 31252882 | t | C | 0.3253 | -0.0037 | 0.0026 | $1.53 \mathrm{E}-01$ | -++ | $3.38 \mathrm{E}-02$ |
| WBC count | rs9368671 | 6 | 31254704 | t | C | 0.3252 | -0.0038 | 0.0026 | $1.46 \mathrm{E}-01$ | -++ | 3.49E-02 |
| WBC count | rs9380232 | 6 | 31242741 | t | c | 0.0403 | -0.0026 | 0.0055 | $6.36 \mathrm{E}-01$ | ?-+ | $1.80 \mathrm{E}-01$ |
| WBC count | rs9380234 | 6 | 31248568 | a | g | 0.1492 | -0.0062 | 0.0032 | 5.61E-02 | -+- | 6.14E-02 |
| WBC count | rs9380236 | 6 | 31254664 | a | g | 0.6842 | 0.0031 | 0.0028 | $2.66 \mathrm{E}-01$ | +-- | 1.45E-02 |

Table S1. Random-effects meta-analysis of MANTRA identified loci.

| WBC count | rs9391714 | 6 | 31245080 | a | g | 0.1481 | -0.0059 | 0.0032 | 6.68E-02 | -+- | $6.70 \mathrm{E}-02$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| WBC count | rs9394049 | 6 | 31244028 | a | c | 0.0937 | 0.0019 | 0.0038 | $6.25 \mathrm{E}-01$ | ?++ | 7.73E-01 |
| WBC count | rs9461680 | 6 | 31243347 | t | C | 0.4008 | -0.0035 | 0.002 | $8.64 \mathrm{E}-02$ | --+ | 6.93E-02 |
| WBC count | rs9461681 | 6 | 31251040 | a | g | 0.4077 | 0.0015 | 0.0017 | 3.99E-01 | -++ | $1.66 \mathrm{E}-01$ |
| WBC count | rs9461684 | 6 | 31253444 | t | c | 0.4404 | 0.001 | 0.003 | 7.44E-01 | ++- | $2.24 \mathrm{E}-05$ |
| WBC count | rs9468915 | 6 | 31250801 | a | C | 0.5258 | -0.006 | 0.0019 | 1.85E-03 | --- | $9.64 \mathrm{E}-02$ |
| WBC count | rs9468919 | 6 | 31253866 | a | g | 0.1486 | 0.013 | 0.0031 | 3.20E-05 | -++ | 5.96E-01 |
| WBC count | rs9468920 | 6 | 31254936 | a | g | 0.4072 | 0.0017 | 0.0018 | 3.36E-01 | -++ | 3.89E-02 |
| WBC count | rs9468922 | 6 | 31255286 | a | g | 0.4359 | -4.00E-04 | 0.0017 | 8.32E-01 | -++ | 4.53E-01 |
| WBC count | rs10094039 | 8 | 130589676 | a | g | 0.4504 | 0.0056 | 0.0017 | $1.31 \mathrm{E}-03$ | ++- | 7.97E-03 |
| WBC count | rs10103048 | 8 | 130602281 | a | c | 0.4571 | 0.008 | 0.0023 | $4.30 \mathrm{E}-04$ | +?+ | $1.65 \mathrm{E}-02$ |
| WBC count | rs10107630 | 8 | 130603635 | t | c | 0.4953 | -0.006 | 0.0017 | 5.04E-04 | --+ | 2.30E-02 |
| WBC count | rs10458304 | 8 | 130596459 | t | C | 0.3851 | -4.00E-04 | 0.0043 | $9.32 \mathrm{E}-01$ | -?+ | 7.73E-01 |
| WBC count | rs10956484 | 8 | 130588520 | a | g | 0.1551 | -0.0179 | 0.0032 | $2.44 \mathrm{E}-08$ | -?- | 2.49E-02 |
| WBC count | rs11785889 | 8 | 130603247 | t | c | 0.4256 | 0.0073 | 0.0021 | 3.92E-04 | ++- | $1.79 \mathrm{E}-02$ |
| WBC count | rs11995855 | 8 | 130601919 | t | C | 0.971 | 0.0093 | 0.0199 | $6.40 \mathrm{E}-01$ | ??+ | $1.00 \mathrm{E}+00$ |
| WBC count | rs13277237 | 8 | 130604563 | a | g | 0.497 | -0.0063 | 0.0017 | $2.47 \mathrm{E}-04$ | --- | $3.54 \mathrm{E}-02$ |
| WBC count | rs1368700 | 8 | 130600091 | a | g | 0.3388 | 0.0043 | 0.0018 | $1.80 \mathrm{E}-02$ | ++- | $1.84 \mathrm{E}-03$ |
| WBC count | rs1433577 | 8 | 130595881 | a | g | 0.3429 | 0.0064 | 0.0019 | 5.67E-04 | 0 | $1.85 \mathrm{E}-02$ |
| WBC count | rs1433578 | 8 | 130601089 | t | c | 0.5297 | 0.0148 | 0.0028 | 9.77E-08 | ++- | 3.50E-03 |
| WBC count | rs1433580 | 8 | 130603707 | t | c | 0.4096 | -0.0067 | 0.0018 | 2.33E-04 | --- | $3.49 \mathrm{E}-02$ |
| WBC count | rs16904119 | 8 | 130593742 | t | g | 0.3902 | -0.0061 | 0.0019 | $1.24 \mathrm{E}-03$ | --- | $1.19 \mathrm{E}-02$ |
| WBC count | rs16904121 | 8 | 130600781 | t | c | 0.4246 | 0.0068 | 0.002 | $9.39 \mathrm{E}-04$ | ++- | 4.76E-02 |
| WBC count | rs17265074 | 8 | 130603007 | t | g | 0.0811 | -6.00E-04 | 0.004 | 8.88E-01 | ?+- | 1.72E-01 |
| WBC count | rs1821340 | 8 | 130605291 | C | g | 0.1665 | 0.0049 | 0.0041 | $2.34 \mathrm{E}-01$ | -?+ | 4.21E-01 |
| WBC count | rs1865223 | 8 | 130606476 | t | g | 0.5206 | -0.0044 | 0.0018 | $1.44 \mathrm{E}-02$ | --+ | $1.01 \mathrm{E}-03$ |
| WBC count | rs2043405 | 8 | 130596548 | a | g | 0.1455 | 0.0031 | 0.0043 | $4.70 \mathrm{E}-01$ | +?+ | 9.75E-01 |
| WBC count | rs2163950 | 8 | 130597585 | a | c | 0.1404 | -0.0167 | 0.0028 | 1.90E-09 | --- | 8.93E-02 |
| WBC count | rs2163951 | 8 | 130603357 | t | c | 0.5054 | -0.0034 | 0.0031 | 2.81E-01 | --- | 8.37E-01 |
| WBC count | rs6470743 | 8 | 130606310 | a | g | 0.327 | 0.0057 | 0.0019 | 2.61E-03 | ++- | 2.09E-02 |
| WBC count | rs6982402 | 8 | 130589924 | t | c | 0.6347 | -0.001 | 0.0042 | 8.11E-01 | +?- | 7.68E-01 |
| WBC count | rs6982553 | 8 | 130589974 | a | t | 0.3543 | 0.0047 | 0.0018 | $9.56 \mathrm{E}-03$ | ++- | 9.89E-04 |
| WBC count | rs6986451 | 8 | 130589711 | a | g | 0.1445 | 0.0037 | 0.0042 | 3.77E-01 | +?+ | 7.47E-01 |
| WBC count | rs7000289 | 8 | 130597230 | a | g | 0.5107 | -0.0054 | 0.0018 | $2.90 \mathrm{E}-03$ | --+ | 4.87E-03 |
| WBC count | rs7000320 | 8 | 130589906 | t | C | 0.4408 | 0.0067 | 0.0018 | $2.68 \mathrm{E}-04$ | +++ | $2.64 \mathrm{E}-02$ |
| WBC count | rs7000372 | 8 | 130590017 | a | g | 0.5079 | -0.0053 | 0.0019 | $4.35 \mathrm{E}-03$ | --+ | $2.45 \mathrm{E}-03$ |
| WBC count | rs7000917 | 8 | 130590087 | a | g | 0.9601 | 0.0015 | 0.006 | 7.99E-01 | -?+ | $6.35 \mathrm{E}-01$ |
| WBC count | rs7004211 | 8 | 130594056 | t | C | 0.631 | -7.00E-04 | 0.0043 | 8.72E-01 | +?- | 7.37E-01 |

Table S1. Random-effects meta-analysis of MANTRA identified loci.

| WBC count | rs7005227 | 8 | 130597858 | a | c | 0.3249 | 0.0055 | 0.0019 | $3.76 \mathrm{E}-03$ | ++- | $1.50 \mathrm{E}-02$ |
| :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- |
| WBC count | rs7008811 | 8 | 130600611 | t | g | 0.4125 | -0.0064 | 0.0018 | $4.31 \mathrm{E}-04$ | --- | $3.31 \mathrm{E}-02$ |
| WBC count | rs7011870 | 8 | 130592498 | a | g | 0.4961 | -0.0049 | 0.0018 | $8.06 \mathrm{E}-03$ | --+ | $1.31 \mathrm{E}-03$ |
| WBC count | rs 7840212 | 8 | 130599247 | t | c | 0.4977 | $-1.00 \mathrm{E}-04$ | 0.0018 | $9.45 \mathrm{E}-01$ | --+ | $2.67 \mathrm{E}-01$ |
| WBC count | rs892353 | 8 | 130601285 | a | g | 0.0793 | -0.002 | 0.0042 | $6.30 \mathrm{E}-01$ | ?+- | $2.55 \mathrm{E}-01$ |
| WBC count | rs 996313 | 8 | 130598217 | a | g | 0.3249 | 0.0057 | 0.0019 | $2.37 \mathrm{E}-03$ | ++- | $2.59 \mathrm{E}-02$ |
| WBC count | rs4794822 | $\mathbf{1 7}$ | $\mathbf{3 8 1 5 6 7 1 2}$ | $\boldsymbol{t}$ | $\boldsymbol{c}$ | $\mathbf{0 . 5 0 9 8}$ | $\mathbf{0 . 0 1 5 8}$ | $\mathbf{1 . 8 0 E - 0 3}$ | $\mathbf{2 . 7 5 E - 1 9}$ | ++- | $\mathbf{2 . 4 9 E - 1 6}$ |

Random-effects meta-analysis performed using METAL. MANTRA identified variants are in bold italics.

Table S3. Credible Interval Statistics

| Trait | SNP | Chr | Position (b37) | Effect Allele | Other Allele | Study Number | $\log 10$ (BF) | PPH | Sample Size | Effect Direction | Genes | set95* |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| Neutrophils | rs7667376 | 4 | 74967890 | C | T | 3 | 11.12528 | 1 | 33743 | -++ | NA | 0.52118827 |
| Neutrophils | rs1371798 | 4 | 74976781 | C | T | 3 | 9.94375 | - 1 | 33753 | -++ | NA | 0.997634695 |
| Neutrophils | rs1371799 | 4 | 74977837 | C | T | 3 | 10.9749 | 1 | 33753 | +-- | NA | 0.889838606 |
| Neutrophils | rs1866755 | 4 | 74978340 | C | T | 3 | 10.27447 | - 1 | 33705 | -++ | NA | 0.963321226 |
| Neutrophils | rs2072633 | 6 | 31919578 | A | G | 3 | 3.44008 | 0.509 | 33698 | --- | Several, HLA | 0.977220747 |
| Neutrophils | rs592229 | 6 | 31930441 | G | T | 3 | 5.91581 | 1 | 33737 | ++- | Several, HLA | 0.941088647 |
| Neutrophils | rs204989 | 6 | 32161852 | A | G | 3 | 3.12774 | 0.041 | 33716 | --- | Several, HLA | 0.986900655 |
| Neutrophils | rs405875 | 6 | 32215188 | C | T | 3 | 3.44774 | 0.026 | 33743 | +++ | Several, HLA | 0.976511099 |
| Neutrophils | rs410283 | 6 | 32215198 | G | T | 3 | 3.65833 | 0.019 | 33753 | -- | Several, HLA | 0.974006156 |
| Neutrophils | rs2515892 | 6 | 32215252 | G | T | 3 | 3.58835 | 0.025 | 33753 | --- | Several, HLA | 0.975004576 |
| Neutrophils | rs3130309 | 6 | 32215472 | A | G | 3 | 3.22801 | 0.529 | 33753 | +-- | Several, HLA | 0.984631838 |
| Neutrophils | rs2395111 | 6 | 32215964 | C | T | 3 | 3.26049 | 0.02 | 32931 | +++ | Several, HLA | 0.983314935 |
| Neutrophils | rs3115575 | 6 | 32216891 | G | T | 3 | 3.73571 | 0.025 | 33753 | +++ | Several, HLA | 0.971611839 |
| Neutrophils | rs6457508 | 6 | 32216963 | C | T | 3 | 3.76889 | 0.013 | 33753 | --- | Several, HLA | 0.970210081 |
| Neutrophils | rs6457510 | 6 | 32217046 | C | T | 3 | 3.22972 | 0.024 | 33753 | --- | Several, HLA | 0.984196353 |
| Neutrophils | rs6936204 | 6 | 32217092 | C | T | 3 | 6.45169 | 0.048 | 33706 | +++ | Several, HLA | 0.728875097 |
| Neutrophils | rs9267971 | 6 | 32217185 | C | T | 3 | 3.79636 | 0.022 | 33597 | +++ | Several, HLA | 0.968697032 |
| Neutrophils | rs9267972 | 6 | 32217232 | A | G | 2 | 4.72678 | 0.016 | 26352 | ?-- | Several, HLA | 0.954820942 |
| Neutrophils | rs3115572 | 6 | 32220484 | C | G | 3 | 3.32271 | 0.487 | 33680 | +-- | Several, HLA | 0.981399393 |
| Neutrophils | rs3115571 | 6 | 32220918 | A | G | 3 | 3.08206 | 0.616 | 33753 | +-- | Several, HLA | 0.988222552 |
| Neutrophils | rs3130316 | 6 | 32221228 | C | T | 3 | 4.61145 | 0.027 | 33737 | +++ | Several, HLA | 0.965350572 |
| Neutrophils | rs3132931 | 6 | 32235895 | G | T | 3 | 2.97088 | 0.019 | 33744 | --- | Several, HLA | 0.98932872 |
| Neutrophils | rs3096673 | 6 | 32238013 | C | T | 3 | 3.02775 | 0.021 | 33744 | --- | Several, HLA | 0.989087816 |
| Neutrophils | rs1559873 | 6 | 32243129 | C | T | 3 | 2.96203 | 0.019 | 33717 | --- | Several, HLA | 0.989564764 |
| Neutrophils | rs9268142 | 6 | 32256418 | C | T | 3 | 3.1193 | 0.02 | 33753 | --- | Several, HLA | 0.987582941 |
| Neutrophils | rs3749966 | 6 | 32261507 | C | T | 3 | 2.94944 | 0.023 | 33744 | --- | Several, HLA | 0.989794063 |
| Neutrophils | rs9268163 | 6 | 32270875 | C | T | 3 | 3.67587 | 0.02 | 33753 | +++ | Several, HLA | 0.972833169 |
| Neutrophils | rs3864301 | 6 | 32271955 | C | T | 3 | 3.26717 | 0.014 | 33753 | +++ | Several, HLA | 0.982845633 |
| Neutrophils | rs9268176 | 6 | 32274079 | C | T | 3 | 3.39956 | 0.023 | 33753 | +++ | Several, HLA | 0.978515431 |
| Neutrophils | rs7341300 | 6 | 32275027 | C | T | 3 | 3.37453 | 0.013 | 33753 | +++ | Several, HLA | 0.979747993 |
| Neutrophils | rs9268181 | 6 | 32276009 | C | T | 3 | 3.38306 | 0.013 | 33697 | --- | Several, HLA | 0.979137764 |
| Neutrophils | rs9268182 | 6 | 32276027 | G | T | 3 | 3.0606 | 0.022 | 33753 | --- | Several, HLA | 0.988518737 |
| Neutrophils | rs9268204 | 6 | 32279500 | C | T | 3 | 3.33732 | 0.018 | 33753 | +++ | Several, HLA | 0.980308116 |
| Neutrophils | rs6934429 | 6 | 32279622 | G | T | 3 | 3.40078 | 0.017 | 33743 | +++ | Several, HLA | 0.977868998 |
| Neutrophils | rs9268207 | 6 | 32280462 | C | T | 3 | 3.05808 | 0.02 | 33702 | +++ | Several, HLA | 0.988813208 |
| Neutrophils | rs1018434 | 6 | 32281360 | C | T | 3 | 3.26851 | 0.025 | 33738 | --- | Several, HLA | 0.982369056 |
| Neutrophils | rs3129959 | 6 | 32375796 | A | T | 2 | 3.82824 | 0.017 | 26352 | ?++ | Several, HLA | 0.967085187 |
| Neutrophils | rs9268503 | 6 | 32377061 | C | T | 2 | 3.17613 | 0.025 | 26352 | ?-- | Several, HLA | 0.985430664 |
| Neutrophils | rs6932810 | 6 | 32380190 | A | G | 2 | 2.94422 | 0.029 | 26352 | ?++ | Several, HLA | 0.990020623 |
| Neutrophils | rs6918317 | 6 | 32381210 | C | T | 2 | 3.32915 | 0.016 | 26352 | ?++ | Several, HLA | 0.9808578 |
| Neutrophils | rs2239806 | 6 | 32411307 | C | T | 3 | 3.1711 | 0.055 | 33744 | -++ | Several, HLA | 0.985812664 |
| Neutrophils | rs1051336 | 6 | 32412592 | A | G | 3 | 3.20433 | 0.036 | 33713 | --- | Several, HLA | 0.985044214 |
| Neutrophils | rs1041885 | 6 | 32412809 | A | T | 3 | 3.23663 | 0.034 | 33744 | +-- | Several, HLA | 0.98375915 |

Table S3. Credible Interval Statistics

| Neutrophils | rs9286790 | 6 | 32439828 | A | G | 3 | 3.15656 | 0.027 | 33753 | +++ | Several, HLA | 0.986554952 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| Neutrophils | rs602875 | 6 | 32573629 | A | G | 2 | 3.10547 | 1 | 26352 | ?-+ | Several, HLA | 0.987911363 |
| Neutrophils | rs2395225 | 6 | 32590624 | C | T | 2 | 3.12463 | 0.018 | 26352 | ?-- | Several, HLA | 0.987243891 |
| Neutrophils | rs4713581 | 6 | 32660023 | C | T | 2 | 3.28066 | 0.059 | 26352 | ?++ | Several, HLA | 0.981891006 |
| Neutrophils | rs2858310 | 6 | 32668323 | A | G | 3 | 3.16059 | 0.02 | 33753 | +++ | Several, HLA | 0.98618553 |
| Neutrophils | rs9469246 | 6 | 32692227 | G | T | 3 | 3.48349 | 0.109 | 33701 | -++ | Several, HLA | 0.975788823 |
| Neutrophils | rs445 | 7 | 92408370 | C | T | 3 | 10.52497 | 1 | 33744 | -++ | CDK6 | 0.999977503 |
| Neutrophils | rs4794822 | 17 | 38156712 | C | T | 3 | 28.91982 | 1 | 33753 | +-- | NA | 0.874563179 |
| Neutrophils | rs8078723 | 17 | 38166879 | C | T | 3 | 28.06213 | 1 | 33693 | -++ | NA | 0.995930339 |

Table S3. Credible Interval Statistics

| Trait | SNP | Chr | Position (b37) | Effect Allele | Other Allele | Study Number | $\begin{aligned} & \log 10 \\ & (B F) \end{aligned}$ | PPH | Sample Size | Effect Direction | Genes | set95* |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| Monocytes | rs1449263 | 2 | 182319301 | C | T | 3 | 19.23857 | 1 | 33729 | +-- | ITGA4 | 0.513404537 |
| Monocytes | rs2124440 | 2 | 182328214 | A | G | 3 | 19.1084 | 1 | 33681 | +++ | ITGA4 | 0.893847628 |
| Monocytes | rs1375493 | 2 | 182323766 | A | G | 3 | 18.55404 | 1 | 33745 | +++ | ITGA4 | 1 |
| Monocytes | rs9880192 | 3 | 128297569 | C | G | 3 | 8.30125 | 1 | 33745 | +- | NA | 0.958237747 |
| Monocytes | rs9839508 | 3 | 128310794 | A | T | 3 | 6.92642 | 1 | 33704 | -++ | NA | 0.998662118 |
| Monocytes | rs9849126 | 3 | 128312069 | C | T | 3 | 5.30501 | 0.982 | 33708 | +++ | NA | 0.999628688 |
| Monocytes | rs2047076 | 5 | 76058509 | C | T | 3 | 6.03105 | 1 | 33729 | +++ | NA | 0.998865282 |
| Monocytes | rs1264376 | 6 | 30765579 | A | C | 3 | 3.99291 | 1 | 28260 | +-- | Several, HLA | 0.987763272 |
| Monocytes | rs1264344 | 6 | 30800577 | C | T | 3 | 4.0451 | 1 | 33693 | -+- | Several, HLA | 0.985309756 |
| Monocytes | rs2535326 | 6 | 30832063 | C | T | 3 | 4.12249 | 1 | 33728 | -+- | Several, HLA | 0.983486311 |
| Monocytes | rs2844654 | 6 | 30838688 | G | T | 3 | 4.02148 | 1 | 33729 | +-+ | Several, HLA | 0.986991592 |
| Monocytes | rs1264333 | 6 | 30844314 | C | T | 3 | 4.08467 | 1 | 33729 | +-+ | Several, HLA | 0.98443954 |
| Monocytes | rs2240804 | 6 | 30920890 | A | G | 3 | 4.46125 | 1 | 33729 | --+ | Several, HLA | 0.97929274 |
| Monocytes | rs3131012 | 6 | 31115441 | C | T | 3 | 5.00397 | 1 | 33729 | + | Several, HLA | 0.945313282 |
| Monocytes | rs2073717 | 6 | 31122126 | C | G | 3 | 4.65235 | 0.288 | 33671 | --- | Several, HLA | 0.973796085 |
| Monocytes | rs3130456 | 6 | 31126252 | A | C | 3 | 4.29037 | 0.405 | 33729 | --- | Several, HLA | 0.982446349 |
| Monocytes | rs3095239 | 6 | 31126790 | A | G | 3 | 4.74009 | 0.475 | 33621 | +++ | Several, HLA | 0.966295715 |
| Monocytes | rs1419880 | 6 | 31135197 | C | T | 3 | 6.51791 | 1 | 33745 | +-- | Several, HLA | 0.767942886 |
| Monocytes | rs6904669 | 6 | 31192796 | A | G | 3 | 3.87778 | 1 | 33728 | ++- | Several, HLA | 0.9896044 |
| Monocytes | rs3130712 | 6 | 31209510 | C | T | 3 | 3.90702 | 1 | 33674 | -++ | Several, HLA | 0.988396482 |
| Monocytes | rs28570051 | 6 | 31210601 | C | T | 2 | 4.7051 | 1 | 15304 | -+? | Several, HLA | 0.970273372 |
| Monocytes | rs3095254 | 6 | 31221668 | C | G | 3 | 6.81258 | 1 | 33745 | +++ | Several, HLA | 0.509456724 |
| Monocytes | rs3130441 | 6 | 31228565 | A | G | 3 | 4.31576 | 0.889 | 33745 | +++ | Several, HLA | 0.980915624 |
| Monocytes | rs2844623 | 6 | 31232543 | C | T | 3 | 6.21948 | 1 | 33720 | +-- | Several, HLA | 0.897962028 |
| Monocytes | rs1049281 | 6 | 31236567 | C | T | 2 | 4.6144 | 1 | 26360 | ?-+ | Several, HLA | 0.977024039 |
| Monocytes | rs9264636 | 6 | 31238297 | C | T | 3 | 5.11292 | 1 | 33745 | -+- | Several, HLA | 0.928636566 |
| Monocytes | rs9264664 | 6 | 31239227 | C | T | 3 | 5.04802 | 1 | 33745 | -+- | Several, HLA | 0.937397437 |
| Monocytes | rs2524084 | 6 | 31241639 | A | G | 3 | 3.84627 | 0.969 | 33729 | +++ | Several, HLA | 0.990154951 |
| Monocytes | rs9264699 | 6 | 31241933 | A | C | 2 | 3.89501 | 0.106 | 26360 | ?-- | Several, HLA | 0.98901242 |
| Monocytes | rs6919908 | 6 | 31244960 | C | T | 2 | 4.82581 | 1 | 16373 | +-? | Several, HLA | 0.961984325 |
| Monocytes | rs2524064 | 6 | 31247817 | A | G | 2 | 4.88433 | 0.269 | 26360 | ?-+ | Several, HLA | 0.951323045 |
| Monocytes | rs2524053 | 6 | 31252469 | G | T | 3 | 4.0388 | 1 | 33745 | --+ | Several, HLA | 0.98616744 |
| Monocytes | rs2853928 | 6 | 31257511 | A | C | 3 | 4.8386 | 1 | 33745 | ++- | Several, HLA | 0.956732173 |
| Monocytes | rs6457374 | 6 | 31272261 | C | T | 3 | 5.41726 | 1 | 33727 | -+- | Several, HLA | 0.918463607 |
| Monocytes | rs10107630 | 8 | 130603635 | C | T | 3 | 14.3687 | 1 | 33745 | -++ | CCDC26 | 0.994867393 |
| Monocytes | rs13277237 | 8 | 130604563 | A | G | 3 | 15.9041 | 1 | 33728 | --- | CCDC26 | 0.793234363 |
| Monocytes | rs4480083 | 8 | 130609320 | C | T | 3 | 13.95825 | 1 | 33745 | -++ | CCDC26 | 0.999230753 |
| Monocytes | rs4407843 | 8 | 130609368 | A | C | 2 | 15.28985 | 0.012 | 26360 | ?++ | CCDC26 | 0.983640202 |
| Monocytes | rs2163952 | 8 | 130610389 | C | T | 3 | 15.92914 | 1 | 33697 | -++ | CCDC26 | 0.40804784 |
| Monocytes | rs1991866 | 8 | 130624105 | C | G | 3 | 15.30419 | 1 | 33598 | -- | CCDC26 | 0.890008896 |

Table S3. Credible Interval Statistics

| Monocytes | rs10980800 | 9 | 113915905 | C | T | 3 | 11.63069 | 1 | 33692 | -++ | RP11-202G18.1 | 0.943693203 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| Monocytes | rs12346772 | 9 | 113920599 | A | G | 3 | 11.92453 | 0.999 | 33745 | --- | RP11-202G18.1 | 0.862140888 |
| Monocytes | rs12350763 | 9 | 113923723 | A | G | 3 | 12.55646 | 1 | 33745 | --- | RP11-202G18.1 | 0.262246809 |
| Monocytes | rs7034139 | 9 | 113924517 | A | C | 3 | 12.44428 | 1 | 33745 | +++ | RP11-202G18.1 | 0.46479586 |
| Monocytes | rs1330279 | 9 | 113926908 | A | G | 3 | 11.84052 | 1 | 33745 | +++ | RP11-202G18.1 | 0.912580323 |
| Monocytes | rs12337595 | 9 | 113927801 | A | G | 3 | 12.27785 | 1 | 33745 | --- | RP11-202G18.1 | 0.800936831 |
| Monocytes | rs10124626 | 9 | 113940107 | A | C | 3 | 11.40469 | 1 | 33745 | --- | RP11-202G18.1 | 0.992935828 |
| Monocytes | rs10123393 | 9 | 113941670 | C | T | 3 | 12.43457 | 1 | 33745 | -++ | RP11-202G18.1 | 0.662866549 |
| Monocytes | rs17812386 | 9 | 113942888 | A | G | 3 | 11.62563 | 1 | 33745 | +++ | RP11-202G18.1 | 0.974445688 |

Table S3. Credible Interval Statistics

| Trait | SNP | Chr | Position (b37) | Effect <br> Allele | Other Allele | Study Number | $\begin{aligned} & \log 10 \\ & (B F) \end{aligned}$ | PPH | $\begin{aligned} & \text { Sample } \\ & \text { Size } \end{aligned}$ | Effect Direction | Genes | set95* |  |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| WBC | rs2518564 | 1 | 159062436 | A | G | 3 | 341.5851¢ | 1 | 51768 | +++ | DARC | Inf |  |
| WBC | rs6734238 | 2 | 113841030 | A | G | 3 | 6.61886 | 0.028 | 52691 | --- | IL1RN | 0.680435948 | A |
| WBC | rs12328368 | 2 | 113847144 | C | G | 3 | 5.82474 | 0.021 | 52702 | --- | IL1RN | 0.789747799 | A |
| WBC | rs12329129 | 2 | 113847007 | A | G | 3 | 5.56842 | 0.037 | 52740 | +++ | IL1RN | 0.850330308 | A |
| WBC | rs6738239 | 2 | 113850792 | A | C | 3 | 5.34698 | 0.022 | 52700 | +++ | IL1RN | 0.88671404 | A |
| WBC | rs1446509 | 2 | 113851159 | A | T | 3 | 5.26256 | 0.028 | 52740 | --- | IL1RN | 0.916670276 | A |
| WBC | rs6741180 | 2 | 113844384 | A | G | 3 | 5.02042 | 0.035 | 52740 | +++ | IL1RN | 0.933823559 | A |
| WBC | rs6746979 | 2 | 113854120 | A | T | 3 | 4.99966 | 0.026 | 52740 | +++ | IL1RN | 0.950176176 | A |
| WBC | rs6750559 | 2 | 113841532 | A | G | 3 | 4.95959 | 0.025 | 52693 | +++ | IL1RN | 0.965087536 | A |
| WBC | rs13382561 | 2 | 113863536 | A | G | 3 | 4.91913 | 0.037 | 52740 | --- | IL1RN | 0.978672461 | A |
| WBC | rs6743376 | 2 | 113832333 | A | C | 2 | 4.70749 | 0.014 | 33230 | ++? | IL1RN | 0.987017286 | A |
| WBC | rs7574159 | 2 | 113859761 | A | G | 3 | 4.43428 | 0.025 | 52740 | +++ | IL1RN | 0.991465721 | A |
| WBC | rs10171839 | 2 | 219051314 | A | G | 3 | 4.99431 | 0.71 | 52685 | -++ | ARPC2,TRNA | 0.9604962 | B |
| WBC | rs13026485 | 2 | 219057984 | C | T | 3 | 5.63031 | 0.011 | 52698 | +++ | ARPC2,TRNA | 0.910468592 | B |
| WBC | rs6752254 | 2 | 219058743 | C | T | 3 | 4.9914 | 0.026 | 52740 | --- | ARPC2,TRNA | 0.966542104 | B |
| WBC | rs10206984 | 2 | 219066980 | C | T | 3 | 4.99748 | 0.039 | 52740 | +-- | ARPC2,TRNA | 0.954409649 | B |
| WBC | rs13384682 | 2 | 219069137 | A | G | 3 | 5.16233 | 0.056 | 52740 | -++ | ARPC2,TRNA | 0.933189765 | B |
| WBC | rs13393821 | 2 | 219076742 | C | G | 3 | 4.61738 | 0.155 | 52740 | +-- | ARPC2,TRNA | 0.990878058 | B |
| WBC | rs12694432 | 2 | 219082330 | A | G | 3 | 5.34854 | 0.049 | 52740 | +- | ARPC2,TRNA | 0.924228048 | B |
| WBC | rs7578940 | 2 | 219083318 | G | T | 3 | 6.17796 | 0.026 | 52737 | +++ | ARPC2,TRNA | 0.805936872 | B |
| WBC | rs4674274 | 2 | 219089131 | A | T | 3 | 4.64791 | 0.167 | 52740 | -++ | ARPC2,TRNA | 0.988322762 | B |
| WBC | rs4674275 | 2 | 219097738 | A | G | 3 | 4.86409 | 0.093 | 52740 | +-- | ARPC2,TRNA | 0.982434553 | B |
| WBC | rs10932765 | 2 | 219099484 | C | T | 3 | 6.95452 | 0.025 | 52740 | +++ | ARPC2,TRNA | 0.55536852 | B |
| WBC | rs7605980 | 2 | 219100869 | C | G | 3 | 4.95231 | 0.099 | 52683 | -++ | ARPC2,TRNA | 0.977924816 | B |
| WBC | rs10169718 | 2 | 219103580 | A | G | 3 | 5.84272 | 0.048 | 52701 | +-- | ARPC2,TRNA | 0.848868885 | B |
| WBC | rs6436047 | 2 | 219108913 | A | C | 3 | 5.07854 | 0.146 | 52701 | +-- | ARPC2,TRNA | 0.948278508 | B |
| WBC | rs6436048 | 2 | 219109033 | A | G | 3 | 4.97763 | 0.097 | 52701 | -++ | ARPC2,TRNA | 0.97239932 | B |
| WBC | rs6738394 | 2 | 219110625 | A | G | 3 | 5.0964 | 0.097 | 52740 | +-- | ARPC2,TRNA | 0.940889243 | B |
| WBC | rs13392177 | 2 | 219112956 | C | T | 3 | 5.7574 | 0.029 | 52665 | +++ | ARPC2,TRNA | 0.884143417 | B |
| WBC | rs13430006 | 2 | 219113089 | G | T | 3 | 6.40768 | 0.016 | 52668 | --- | ARPC2,TRNA | 0.713035686 | B |
| WBC | rs3731859 | 2 | 219124222 | A | G | 3 | 4.70781 | 0.024 | 52740 | +++ | ARPC2,TRNA | 0.985581369 | B |
| WBC | rs3806792 | 4 | 74965274 | C | T | 3 | 11.40926 | 1 | 52739 | +-- | NA | 0.988197424 |  |
| WBC | rs7667376 | 4 | 74967890 | C | T | 3 | 12.95485 | 1 | 52740 | -++ | NA | 0.983622655 |  |
| WBC | rs1371798 | 4 | 74976781 | C | T | 3 | 11.35837 | 1 | 52740 | -++ | NA | 0.992266343 |  |
| WBC | rs1371799 | 4 | 74977837 | C | T | 3 | 13.49611 | 1 | 52740 | +-- | NA | 0.558752008 |  |
| WBC | rs1866755 | 4 | 74978340 | C | T | 3 | 13.17081 | 1 | 52690 | -++ | NA | 0.822943664 |  |
| WBC | rs2853946 | 6 | 31247203 | A | T | 3 | 12.15511 | 0.009 | 52740 | --- | HLA-B | 0.996632812 | A |
| WBC | rs45855 | 6 | 32189481 | A | T | 3 | 4.70607 | 0.009 | 51806 | +++ | Several, HLA | 0.990319437 | B |
| WBC | rs9267972 | 6 | 32217232 | A | G | 2 | 6.0505 | 0.012 | 36352 | ?-- | Several, HLA | 0.93845529 | B |
| WBC | rs3115572 | 6 | 32220484 | C | G | 3 | 5.38859 | 0.037 | 52673 | --- | Several, HLA | 0.971244947 | B |
| WBC | rs9268402 | 6 | 32341353 | A | G | 3 | 6.51568 | 0.006 | 52702 | +++ | Several, HLA | 0.866294331 | B |
| WBC | rs9268473 | 6 | 32355683 | A | G | 3 | 5.5005 | 0.01 | 52739 | --- | Several, HLA | 0.965853416 | B |

Table S3. Credible Interval Statistics

| WBC | rs7194 | 6 | 32412480 | A | G | 3 | 5.45412 | 0.017 | 41911 | +++ | Several, HLA | 0.968752176 | B |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| WBC | rs1051336 | 6 | 32412592 A | A | G | 3 | 4.95743 | 0.014 | 52692 | --- | Several, HLA | 0.982602579 | B |
| WBC | rs1041885 | 6 | 32412809 | A | T | 3 | 4.85516 | 0.011 | 51806 | --- | Several, HLA | 0.986669768 | B |
| WBC | rs2227139 | 6 | 32413459 | A | G | 3 | 4.94958 | 0.024 | 52740 | +++ | Several, HLA | 0.983509715 | B |
| WBC | rs3763327 | 6 | 32413830 | C | G | 3 | 5.09542 | 0.017 | 52740 | +++ | Several, HLA | 0.98051656 | B |
| WBC | rs9286790 | 6 | 32439828 | A | G | 3 | 5.20455 | 0.015 | 52740 | +++ | Several, HLA | 0.974799402 | B |
| WBC | rs12194148 | 6 | 32444198 | G | T | 3 | 5.13929 | 0.854 | 52740 | +-- | Several, HLA | 0.979247412 | B |
| WBC | rs9378212 | 6 | 32445691 | C | T | 3 | 5.17552 | 0.762 | 52740 | +-- | Several, HLA | 0.976325605 | B |
| WBC | rs9378213 | 6 | 32448398 | G | T | 3 | 4.93964 | 0.781 | 52740 | -++ | Several, HLA | 0.984396325 | B |
| WBC | rs9391786 | 6 | 32448561 | A | G | 3 | 7.05769 | 0.026 | 52740 | --- | Several, HLA | 0.832892343 | B |
| WBC | rs5020946 | 6 | 32450089 | G | T | 3 | 4.85706 | 0.733 | 52710 | +-- | Several, HLA | 0.985939886 | B |
| WBC | rs9270657 | 6 | 32566021 | G | T | 3 | 5.8223 | 0.995 | 52740 | +-- | Several, HLA | 0.95652103 | B |
| WBC | rs602875 | 6 | 32573629 | A | G | 2 | 7.59229 | 1 | 36352 | ?-+ | Several, HLA | 0.398457786 | B |
| WBC | rs615672 | 6 | 32574171 | C | G | 3 | 4.90065 | 0.014 | 52614 | --- | Several, HLA | 0.985206805 | B |
| WBC | rs9271562 | 6 | 32590411 | A | G | 2 | 6.50348 | 0.01 | 36352 | ?-- | Several, HLA | 0.898771063 | B |
| WBC | rs2395225 | 6 | 32590624 | C | T | 2 | 7.23567 | 0.011 | 36352 | ?-- | Several, HLA | 0.573749876 | B |
| WBC | rs9271586 | 6 | 32590899 | G | T | 2 | 5.17311 | 0.169 | 36352 | ?-- | Several, HLA | 0.977843363 | B |
| WBC | rs9271850 | 6 | 32595060 | A | G | 2 | 7.1466 | 0.011 | 36352 | ?++ | Several, HLA | 0.716538075 | B |
| WBC | rs2395237 | 6 | 32690945 | A | C | 3 | 5.27583 | 0.012 | 51716 | --- | Several, HLA | 0.973167695 | B |
| WBC | rs9469246 | 6 | 32692227 | G | T | 3 | 4.84978 | 0.797 | 52673 | -++ | Several, HLA | 0.987390663 | B |
| WBC | rs1383264 | 6 | 32739967 | A | T | 3 | 5.77773 | 0.008 | 52640 | --- | Several, HLA | 0.962627951 | B |
| WBC | rs3117016 | 6 | 33095516 | A | G | 3 | 6.14184 | 0.035 | 52701 | +++ | Several, HLA | 0.912894237 | B |
| WBC | rs3117014 | 6 | 33095615 | C | T | 3 | 5.05724 | 0.214 | 52740 | -++ | Several, HLA | 0.981678898 | B |
| WBC | rs3117013 | 6 | 33095636 | A | G | 3 | 6.14164 | 0.018 | 52740 | --- | Several, HLA | 0.927010909 | B |
| WBC | rs3130234 | 6 | 33096009 | A | C | 3 | 4.77032 | 0.022 | 52740 | --- | Several, HLA | 0.989258646 | B |
| WBC | rs3117008 | 6 | 33096274 | A | G | 3 | 4.7826 | 0.029 | 52740 | +++ | Several, HLA | 0.988658285 | B |
| WBC | rs3117007 | 6 | 33096514 | C | G | 3 | 4.7267 | 0.014 | 52740 | +++ | Several, HLA | 0.989801637 | B |
| WBC | rs3117005 | 6 | 33096755 | A | G | 3 | 4.80485 | 0.023 | 52740 | --- | Several, HLA | 0.988040706 | B |
| WBC | rs3129207 | 6 | 33125312 | C | G | 3 | 6.04494 | 0.011 | 52699 | +++ | Several, HLA | 0.94975409 | B |
| WBC | rs9376090 | 6 | 135411228 | C | T | 3 | 8.98253 | 0.022 | 43711 | --- | HBS1L | 0.95209774 | C |
| WBC | rs7776054 | 6 | 135418916 | A | G | 3 | 8.47363 | 0.054 | 52740 | +++ | HBS1L | 0.988485197 | C |
| WBC | rs11759553 | 6 | 135422296 | A | T | 3 | 8.6132 | 0.895 | 52740 | -++ | HBS1L | 0.981048367 | C |
| WBC | rs9373124 | 6 | 135423209 | C | T | 3 | 8.38179 | 0.767 | 52740 | +-- | HBS1L | 0.994504523 | C |
| WBC | rs4895440 | 6 | 135426558 | A | T | 3 | 8.87397 | 0.969 | 52740 | -++ | HBS1L | 0.970792843 | C |
| WBC | rs4895441 | 6 | 135426573 | A | G | 3 | 9.63521 | 0.165 | 52703 | +++ | HBS1L | 0.86513885 | C |
| WBC | rs9389269 | 6 | 135427159 | C | T | 3 | 9.40127 | 0.276 | 52740 | +-- | HBS1L | 0.928093506 | C |
| WBC | rs9402686 | 6 | 135427817 | A | G | 3 | 10.38729 | 0.055 | 52740 | --- | HBS1L | 0.609604066 | C |
| WBC | rs6920211 | 6 | 135431318 | C | T | 3 | 9.77147 | 0.167 | 52732 | --- | HBS1L | 0.757252163 | C |
| WBC | rs445 | 7 | 92408370 | C | T | 3 | 16.80458 | 1 | 52740 | -++ | CDK6 | 1 |  |
| WBC | rs2894479 | 8 | 130584161 | A | C | 2 | 6.64341 | 0.116 | 33223 | --? | CCDC26 | 0.950697794 |  |
| WBC | rs10956484 | 8 | 130588520 | A | G | 2 | 6.66953 | 0.15 | 32297 | --? | CCDC26 | 0.886495614 |  |
| WBC | rs2163950 | 8 | 130597585 | A | C | 3 | 7.74878 | 0.065 | 52740 | --- | CCDC26 | 0.818313608 |  |
| WBC | rs1433578 | 8 | 130601089 | C | T | 3 | 6.31136 | 0.593 | 52740 | +-- | CCDC26 | 0.980585995 |  |
| WBC | rs13263733 | 8 | 130661418 | A | G | 2 | 5.90041 | 0.026 | 36352 | ?-- | CCDC26 | 0.992188447 |  |

Table S3. Credible Interval Statistics

| WBC | rs2241245 | 17 | 38151014 | C | T | 3 | 30.26561 | 1 | 52740 | -++ | PSMD3 | 0.47512314 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| WBC | rs4065321 | 17 | 38143548 | C | T | 3 | 29.70093 | 1 | 52740 | -++ | PSMD3 | 0.604580332 |
| WBC | rs4794822 | 17 | 38156712 | C | T | 3 | 29.64331 | 1 | 52740 | +-- | PSMD3 | 0.71795243 |
| WBC | rs9915252 | 17 | 38145088 | C | G | 3 | 29.58448 | 0.029 | 52740 | +++ | PSMD3 | 0.816961774 |
| WBC | rs8073254 | 17 | 38149350 | C | G | 3 | 29.55371 | 0.034 | 52740 | +++ | PSMD3 | 0.909198989 |
| WBC | rs2305482 | 17 | 38140927 | A | C | 3 | 29.36374 | 0.018 | 52701 | +++ | PSMD3 | 0.968756451 |
| WBC | rs11658328 | 17 | 38149236 | C | T | 3 | 29.08263 | 1 | 52740 | +-- | PSMD3 | 0.999932867 |

Variants encompassing the $95 \%$ credible region of an associated region, as identified using the top hits from the MANTRA analysis, are presented for each subtype. PPA = posterior probability of association; the probability that a SNP is truly associated with a phenotype. *set $95=95 \%$ credible set.

Table S4. Summarized eQTL Results

| Blood Cell GWAS SNP | Position (b37) | Blood Cell Trait | Relation To Esnp | R^2 | eQTL SNP | chr | pos(hg18) | Tissue | eQTL p-value | Transcript | Best eSNP For Transcript | Best eSNP p-value | $\mathrm{R}^{\wedge} 2$ (CEU) to Blood Cell GWAS SNP* |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| rs2853946 | 31247203 | WBC | C_SentinelSNP | 1 | rs2853946 | 6 | 31355182 | LCL (MuTHER) | 7.53E-05 | BAT1 | rs206017 | 2.06E-06 | >500kb |
| rs2853946 | 31247203 | WBC | Proxy:ASN,Rel22 | 0.645 | rs3094691 | 6 | 31382672 | LCL (MuTHER) | $1.09 \mathrm{E}-06$ | MICB | rs2844498 | $4.06 \mathrm{E}-12$ | 0.34 |
| rs2853946 | 31247203 | WBC | Proxy:YRI,Rel22 | 0.554 | rs3130521 | 6 | 31304355 | LCL (MuTHER) | $2.28 \mathrm{E}-09$ | TCF19 | rs887464 | $1.20 \mathrm{E}-14$ | 0.333 |
| rs6936204 | 32217092 | Neutrophil | C_SentinelSNP | 1 | rs6936204 | 6 | 32325070 | Whole Blood** | $0.00 \mathrm{E}+00$ | HLA-DRB5 | rs9270986 | $0.00 \mathrm{E}+00$ | 0.278 |
| rs6936204 | 32217092 | Neutrophil | Proxy:YRI,Rel22 | 0.848 | rs1559873 | 6 | 32351107 | Whole Blood** | $0.00 \mathrm{E}+00$ | HLA-DRB5 | rs9270986 | 0.00E+00 | 0.057 |
| rs6936204 | 32217092 | Neutrophil | Proxy:ASN,Rel22 | 0.653 | rs9268557 | 6 | 32497283 | Whole Blood** | $0.00 \mathrm{E}+00$ | HLA-DRB5 | rs9270986 | 0.00E+00 | 0.256 |
| rs6936204 | 32217092 | Neutrophil | C_SentinelSNP | 1 | rs6936204 | 6 | 32325070 | LCL (MuTHER) | $1.06 \mathrm{E}-05$ | ATP6V1G2 | rs3132454 | $4.90 \mathrm{E}-14$ | $>500 \mathrm{~kb}$ |
| rs6936204 | 32217092 | Neutrophil | Proxy:ASN,Rel22 | 1 | rs3130320 | 6 | 32331236 | LCL (MuTHER) | 3.01E-06 | ATP6V1G2 | rs3132454 | $4.90 \mathrm{E}-14$ | $>500 \mathrm{~kb}$ |
| rs6936204 | 32217092 | Neutrophil | Proxy:YRI,H3 | 0.781 | rs3096674 | 6 | 32346197 | LCL (MuTHER) | 1.13E-06 | HLA-DPB1 | rs7772134 | 0.00E+00 | >500kb |
| rs3095254 | 31221668 | Monocyte | Proxy:ASN,Rel22 | 0.565 | rs879882 | 6 | 31247431 | LCL (MuTHER) | $2.25 \mathrm{E}-10$ | TCF19 | rs887464 | $1.20 \mathrm{E}-14$ | 0.198 |
| rs3095254 | 31221668 | Monocyte | Proxy:YRI,Rel22 | 0.737 | rs3130409 | 6 | 31321790 | LCL (MuTHER) | 2.00E-06 | TCF19 | rs887464 | $1.20 \mathrm{E}-14$ | 0.301 |
| rs3095254 | 31221668 | Monocyte | C_SentinelSNP | 1 | rs3095254 | 6 | 31329647 | LCL (MuTHER) | $1.54 \mathrm{E}-12$ | TCF19 | rs887464 | $1.20 \mathrm{E}-14$ | 0.371 |
| rs1449263 | 182319301 | Monocyte | Proxy:YRI,Rel22 | 0.539 | rs1449260 | 2 | 182041692 | Monocytes | $4.16 \mathrm{E}-13$ | ITGA4 | rs2124440 | $0.00 \mathrm{E}+00$ | 0.329 |
| rs1449263 | 182319301 | Monocyte | Proxy:ASN,H3 | 0.983 | rs6740847 | 2 | 182016597 | Monocytes (CD14+) | $0.00 \mathrm{E}+00$ | ITGA4 | rs1449263 | 0.00E+00 | 0.967 |
| rs1449263 | 182319301 | Monocyte | C_SentinelSNP | 1 | rs1449263 | 2 | 182027546 | Monocytes (CD14+) | $0.00 \mathrm{E}+00$ | ITGA4 | rs1449263 | 0.00E+00 | 1-sameSNP |

Table 6. Summarized results of the eQTL analysis are reported (all cis). *Not in HapMap build, ** Pax Gene Whole Blood (Mehta). The final R^2 column includes only European ancestry tissue types.


[^0]:    *To whom correspondence should be addressed at: Division of Cardiovascular Medicine, Department of Internal Medicine, University of Michigan, 1500 E. Medical Center Drive, Ann Arbor, MI 48109, USA. Tel: + 17347644500 ; Fax: + 17349368266 ; Email: sganesh@med.umich.edu; Laboratory of Neurogenetics, Building 35, 1A-1014, 35 Convent Drive Bethesda, MD 20892, USA. Tel: 301451 3831; E-mail: nallsm@mail.nih.gov
    ${ }^{\dagger}$ These authors contributed equally to this work.
    ${ }^{\ddagger}$ These authors contributed equally to this work.
    ${ }^{\top}$ A full list of collaborators from CHARGE, COGENT and RIKEN can be found in Supplementary Material.

