

List of Abbreviations

AC allele count

APC antigen-presenting cell

ASC antibody-secreting cell

ASE allele-specific expression

AUC area under the curve

BCR B cell receptor

BH Benjamini-Hochberg

BMI body mass index

BTM blood transcription module

CD Crohn's disease

CDR complementarity-determining region

ChIP-seq chromatin immunoprecipitation sequencing

CPM counts per million

CRP C-reactive protein

CyTOF cytometry by time-of-flight

DAMP damage-associated molecular pattern

DC dendritic cell

df degree of freedom

DGE differential gene expression

ELISA enzyme-linked immunosorbent assay

ELISPOT enzyme-linked immune absorbent spot

eQTL expression quantitative trait locus

FACS fluorescence-activated cell sorting

FC fold change

FDR false discovery rate

FWER family-wise error rate

GO Gene Ontology

GWAS genome-wide association study

HA haemagglutinin

HAI haemagglutination inhibition

HBI Harvey Bradshaw index

HIRD Human Immune Response Dynamics

HLA human leukocyte antigen

HSC hematopoietic stem cell

HWE Hardy-Weinberg equilibrium

IBD inflammatory bowel disease

IIV inactivated influenza vaccine

IMID immune-mediated inflammatory disease

INT inverse normal transformation

IV instrumental variable

LAI live attenuated influenza vaccine

LD linkage disequilibrium

LFSR local false sign rate

LMM linear mixed model

LOCO leave-one-chromosome-out

LOR loss of response

LRT likelihood ratio test

MAF minor allele frequency

MANOVA multivariate analysis of variance

MAR missing at random

MCAR missing completely at random

MHC major histocompatibility complex

ML maximum likelihood

MN microneutralisation

MNAR missing not at random

molQTL molecular quantitative trait locus

MR Mendelian randomisation

mRNA messenger RNA

MS multiple sclerosis

NA neuraminidase

ncRNA non-coding RNA

NK natural killer

PAMP pathogen-associated molecular pattern

PANTS Personalised Anti-TNF Therapy in Crohn's Disease

PBMC peripheral blood mononuclear cell

PC principal component

PCA principal component analysis

PNR primary non-response

PRR pattern recognition receptor

PVE proportion of variance explained

QTL quantitative trait locus

RA rheumatoid arthritis

RBC red blood cell

REML restricted maximum likelihood

reQTL response expression quantitative trait locus

RNA-seq RNA sequencing

rRNA ribosomal RNA

scRNA-seq single-cell RNA sequencing

SLE systemic lupus erythematosus

SNP single nucleotide polymorphism

T1D type 1 diabetes

T2D type 2 diabetes

TF transcription factor

TIV trivalent inactivated influenza vaccine

TLR toll-like receptor

TMM trimmed mean of M-values

TNF tumour necrosis factor

TPM transcripts per million

TRI titre response index

TSS transcription start site

TWAS transcriptome-wide association study

UC ulcerative colitis

UTR untranslated region

WES whole-exome sequencing

WGS whole-genome sequencing

WHO World Health Organization