| Chr | Selection signature | Gene symbol | Gene name | Documented immunity function |
|-----|------------------------------------------------|----------------|---------------------------------------------------------|---------------------------------------------------------------------------------------------------------------------------------------------------------|
| 1 | 12964810-17425222 (3Mb sliding windows) | NCAM | neural cell adhesion molecule 1 | affects the innate immune system in the lung [1] |
| 1 | 63951419-65580210 | IGSF11 | immunoglobulin superfamily member 11 | a novel target for cancer immunotherapy of gastrointestinal and hepatocellular carcinomas [2] |
| | | CD80 | cluster of differentiation 80 | costimulatory signals for T cell proliferation, cytokine production, and generation of CTL[3] |
| | | GSK3B | glycogen synthase kinase-3 beta | negatively regulate IFN-ß Production by TLR4- stimulated innate immune cells [4] |
| 1 | 85294656-88698235 (2Mb sliding windows) | SOX2 | SRY-box 2 | a sequence-specific DNA sensor in neutrophils to initiate innate immunity against microbial infection [5] |
| | | USP13 | ubiquitin specific peptidase 13 | negatively regulates antiviral responses by deubiquitinating STING [6] |
| 2 | 79257749-80298164 | STAT1 | signal transducer and activator of transcription 1 | targeted disruption of the mouse STAT1 gene results in compromised innate immunity to viral disease [7] |
| | | STAT4 | signal transducer and activator of transcription 4 | mediated immune mechanisms in protection against plague [8] |
| 2 | 84373550-86035295 | SLC39A10 | solute carrier family 39 member 10 | controls humoral immunity by modulating B-cell receptor signal strength [9] |
| 2 | 91221891-92326427 | CD28 | cluster of differentiation 28 | promote T cell survival [10] |
| 2 | 120914934-121503634 | TRIM64 | tripartite motif containing 64 | Trim62-deficient mice had increased susceptibility to fungal infection [11] |
| 2 | 128695085-129554709 | GRHL3 | grainyhead like transcription factor 3 | A GRHL3-regulated repair pathway suppresses immune-mediated epidermal hyperplasia [12] |
| | | IFNLR1 | interferon lambda receptor 1 interleukin 22 receptor | expression of Ifnlr1 on intestinal epithelial cells is critical to the antiviral effects of interferon lambda against norovirus and reovirus [13] |
| | | IL22RA1 | subunit alpha 1 | critical role of IL-22/IL22-RA1 signaling in pneumococcal pneumonia [14] |

Additional file 11: Table S4. Immunity genes in or near chromosome regions subjected to genetic selection since 1964.

| 3 | 8685390-9810337 | CD244 CD48 CD84 | cluster of differentiation 244 cluster of differentiation 48 cluster of differentiation 84 | long noncoding RNA derived from CD244 signaling epigenetically controls CD8+ T-cell immune responses in tuberculosis infection [15] CD48-deficient mice have a pronounced defect in CD4+ T cell activation [16] |
|---|---------------------|-----------------------|--------------------------------------------------------------------------------------------------|--------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| 3 | 11105182-11770282 | IGSF8 | immunoglobulin superfamily member 8 | CD4+ T cell activation [10] This gene encodes a member the EWI subfamily of the immunoglobulin protein superfamily. Members of this family contain a single transmembrane domain, an EWI (Glu-Trp-Ile)-motif and a variable number of immunoglobulin domains. This protein interacts with the tetraspanins CD81 and CD9 and may regulate their role in certain cellular functions including cell migration and viral infection. [17] |
| 3 | 16444546-16881296 | CRTC2 | CREB regulated transcription coactivator 2 | the CREB/CRTC2 pathway modulates autoimmune disease by promoting Th17 differentiation [18] |
| 3 | 79282515-80265751 | PDE4B | phosphodiesterase 4B | PDE4B gene activation by LPS constitutes a feedback regulation essential for an efficient immune response [19] |
| 4 | 39012513-39435115 | HGF | hepatocyte growth factor | key regulator of dendritic cell migration in skin immunity [20] |
| 5 | 104116518-105105448 | CD27 CD9 | cluster of differentiation 27 cluster of differentiation 9 | CD27 is required for generation and long-term maintenance of T cell immunity [21] CD9 tetraspanin is not required for the development of peripheral B cells or for humoral immunity [22] |
| 6 | 90075383-90188972 | ANKRD17 | ankyrin repeat domain 17 | Ankrd17 positively regulates RIG-I-like receptor (RLR)-mediated immune signaling [23] |
| 7 | 11486814-12752376 | IL-27Ra | interleukin 27 receptor subunit alpha | IL-27Ra-deficient mice were hypersusceptible to experimental autoimmune encephalomyelitis and generated more IL-17-producing T helper cells [24] |
| 7 | 54455147-57844905 | RNF14 NDFIP1 | ring finger protein 14 Nedd4 family interacting protein 1 | RNF14 is a regulator of mitochondrial and immune function in muscle [25] mediates peripheral tolerance to self and exogenous antigen by inducing cell cycle exit in responding |

| | | | | CD4+ T cells [26] |
|----|---------------------------------------------------------------------|---------------------------------------------------------------------------------------------------------------------------------------------------------------------|--------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| 8 | 37706259-39306605 | IL33 | interleukin 33 | a crucial amplifier of innate rather than acquired immunity [27] |
| 8 | 108652993-109059409 | TLR4 | toll like receptor 4 | links innate immunity and fatty acid–induced insulin resistance [28] |
| 9 | 87190233-88964135 | LATS1 | large tumor suppressor kinase 1 | The Hippo pathway kinases LATS1/2 suppress cancer immunity [29] |
| 10 | 21516136-22008344 (the gene clusters are at 22.1-25.5 Mb) | 135 genes 4 genes 3 genes 3 genes 1 gene 1 gene 18 genes 2 genes 1 gene | T-cell receptor alpha, T-cell receptor alpha chain V region and C region Ig lambda chain V-I region Ig lambda chain V-II region Ig lambda chain V-III region Ig lambda chain V-VI region Ig kappa chain V-I region Ig kappa chain V-VI region Ig kappa chain V-VI region Ig heavy chain V region | T cell receptors recognize foreign antigens and bound to MHC molecules at the surface of antigen presenting cells [30] Immunoglobulins recognize foreign antigens and initiate immune responses such as phagocytosis and the complement system [31] |
| 10 | 25598535-26289588 | NDRG2 | NDRG2 | one of novel intrinsic factors for regulation of IL-10 production in human myeloid cell [32] |
| 10 | 36465813-37290812 | TYRO3 | TYRO3 protein tyrosine kinase | a negative regulator of type 2 immunity [33] |
| 10 | 51001410-52033595 | ADAM10 | ADAM metallopeptidase domain 10 | regulates the production of multiple secreted factors that contribute to autoimmune reactions [34] |
| 11 | 6602495-7047686 | MAP4K4 IL1R2 IL1R1 IL1RL2 IL1RL1 IL18R IL18RAP | mitogen-activated protein kinase kinase kinase kinase 4 interleukin 1 receptor type 2 interleukin 1 receptor type 1 interleukin 1 receptor like 2 interleukin 1 receptor like 1 interleukin 18 receptor 1 interleukin 18 receptor accessory protein | promotes vascular inflammation and atherosclerosis [35] The IL-1 family: regulators of immunity [36] The IL-1 family: regulators of immunity [36] |

| 11 | 15814741-16759475 | RASGRP3 | RAS, guanyl releasing protein 3 | limits inflammatory response [37] |
|----|-------------------|--------------------------------------------|--------------------------------------------------------------------------------------------------------------------------------------|---------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| 11 | 28577854-29495322 | EPAS1 | endothelial PAS domain protein 1 | links DOCK8 deficiency to atopic skin inflammation via IL-31 induction [38] |
| 12 | 35886159-37211264 | LATS2 | large tumor suppressor kinase 2 | The Hippo pathway kinases LATS1/2 suppress cancer immunity [29] |
| 18 | 34794005-35027876 | TRADD | TNFRSF1A associated via death domain | an essential component of the RIG-like helicase antiviral pathway [39] |
| 18 | 50581375-51369666 | AXL CEACAMI | AXL receptor tyrosine kinase carcinoembryonic antigen related cell adhesion molecule 1 | required for T cell priming and antiviral immunity [40] a regulatory co-receptor for both lymphoid and myeloid cell types [41] |
| 18 | 58696066-61421776 | 33 genes9 genes | cationic amino acid transporters sialic acid-binding Ig-like lectin | play key roles in the survival and transmission of apicomplexan parasites [42] Sialic acid-binding immunoglobulin-like lectins (SIGLECs) are a family of cell surface proteins belonging to the immunoglobulin superfamily. They mediate protein-carbohydrate interactions by selectively binding to different sialic acid moieties present on glycolipids and glycoproteins. [43] |
| 20 | 9919863-10295382 | NAIP | NLR family apoptosis inhibitory protein | recognizes bacterial type III secretion needle protein for inflammasome activation [44] |
| 21 | 64129449-66380363 | BCL11B | B-cell CLL/lymphoma 11B | required for differentiation and survival of $\alpha\beta$ T lymphocytes [45] |
| 22 | 53695239-54103673 | CCR1 XCR1 CXCR6 | C-C motif chemokine receptor 9 X-C motif chemokine receptor 1 C-X-C motif chemokine receptor 6 C-C motif chemokine | interaction between CCR1 and CCL5 modulates the innate immune response during sepsis [46] an integral component in the development of efficient cytotoxic immunity in vivo [47] adaptive immunity to structurally diverse antigens [48] an essential role for CCR9 in the homing of pDC to |

| | | CCR9 | receptor 9 | the intestine under homeostatic and inflammatory conditions [49] |
|----|-------------------|-------------------------------------------|----------------------------------------------------------------------------------------------------------------------------------------------------------|-------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| 23 | 7627089-7874236 | BAK1 | BCL2 antagonist/killer 1 | The receptor-like kinase SERK3/BAK1 is a central regulator of innate immunity in plants [50] |
| 23 | 25178791-27992880 | MHC <i>BTNL2</i> ABHD16A NFKBIL1 | major histocompatibility complex butyrophilin-like 2 (MHC class II associated) abhydrolase domain containing 16A NFKB inhibitor like 1 | associated with many diseases [51] Inhibit T Cell Activation [52] immunomodulatory lysophosphatidylserines are regulated by ABHD16A and ABHD12 interplay [53] confers resistance to experimental autoimmune arthritis through the regulation of dendritic cell functions [54] |
| 24 | 6659975-7838578 | <i>CD226</i> | cluster of differentiation 226 | regulates human T cell function [55] |
| 25 | 23825593-26337016 | IL4R | interleukin 4 receptor | regulate increased lymphocyte proliferation and survival [56] |
| | | IL21R | interleukin 21 receptor | critical for sustained functionality and control of chronic viral infection [57, 58] |
| | | IL27 | interleukin 27 | expression of IL-27 in murine carcinoma cells produces antitumor effects and induces protective immunity in inoculated host animals [59] |
| | | LAT | linker for activation of T- cells | essential role in T cell development [60] |
| | | CD19 | cluster of differentiation 19 | regulates signal transduction thresholds governing humoral immunity and autoimmunity [61] |
| 26 | 16999066-17465791 | BLNK | B-cell linker | an essential role in human B cell development [62] |
| 29 | 43328607-43839783 | BATF2 | basic leucine zipper ATF- like transcription factor 2 | Batf2/Irf1 induces inflammatory responses in classically activated macrophages, lipopolysaccharides, and mycobacterial infection [63] |

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