

Supplementary Table 1: selected examples of immunological variation among common inbred strains of laboratory mice.

Gene	Function	Strain with deficits or polymorphisms	Normal expression	Strain-related References
Aryl hydrocarbon receptor (<i>Ahr</i>)	Activation of <i>Cyp450</i> expression, decreased IL22, increased IL17	<i>Ahr^d</i> 129/J, AKR, DBA/2J, NZB, SJL/J, SWR	<i>Ahr^{b-1}</i> <i>Ahr^{b-2}</i> A/J, BALB/c, C57BL/6J, CBA/J, C3H/HeJ, DBA/1J	¹⁻⁴
Cathepsin E (<i>Ctse</i>)	Chemotaxis, cell adhesion, antigen processing for MHC class II presentation	C57BL/6JOlaHsd, C57BL/6	BALB/c, 129S2/SvHsd, CBA/J	⁵
Dedicator of cytokinesis 2 (<i>Dock2</i>)	Chemotaxis, lymphocyte migration, B cell development, plasmacytoid DCs function	C57BL/6NHsd (from specific breeding facilities)	All other C57BL/6 substrains	^{6,7}
Complement factor C5 (<i>Hc</i>)	Complement system, innate immunity, C5a anaphylatoxin with spasmogenic and chemotactic activity, C5b subunit of the membrane attack complex	A/J, AKR/J, DBA/2, FVB/NJ, NOD/ShiLtJ, SWR	C57BL/6, 129 strains, DBA/1, MRL/MpJ, NZB, NZW	^{8,9}
Fc receptor, IgE, low affinity II, alpha polypeptide (<i>Fcer2a</i>)	CD23, low-affinity receptor for IgE, negative regulation of IgE response	<i>Fcer2a^{Hie}</i> (reduced level of expression) NZB, 129P1/ReJ, 129/SvJ	BALB/c, C57BL/6	^{10,11}
Interleukin 2 (<i>Il2</i>)	Clonal expansion of B and T cells, immune cell development and activation, immune tolerance, adaptive cell-mediated immunity	<i>Il2^{m1}</i> (hypoactive) MRL/MpJ, NOD/ShiLtJ, SJL/J	C57BL/6	¹²
Interleukin 12b (<i>Il12b</i>)	Development and activation of T cells and NK cell, promotion of TH1-type immune response	Polymorphism enhances activity SJL/J	C57BL/6, C57BL/10ScCr	¹³⁻¹⁵
MX dynamin-like GTPase 1 and 2 (<i>Mx1</i> and <i>Mx2</i>)	Innate immunity, IFN α and IFN β -mediated response to viral infection	Most inbred strains	Most of the wild mice or wild-derived strains	^{16,17}
Neuronal apoptosis inhibitor protein (<i>Naip5</i>)	Recognition of flagellin, activation of the inflammasome, macrophage pyroptosis	A/J	Most inbred strains	^{18,19}
2'-5' oligoadenylate synthetase 1B (<i>Oas1b</i>)	Activation of RNA nucleases, innate immunity, protection against flaviviruses	Most inbred strains	Most of wild mice or wild-derived strains	²⁰

Gene	Function	Strain with deficits or polymorphisms	Normal expression	Strain-related References
NLR family, pyrin domain containing 1B (<i>Nlrp1b</i>)	Innate immunity, inflammasome, response to PAMPs and DAMPs, susceptibility/resistance to <i>Bacillus anthracis</i> infection and anthrax LT challenge	<i>Nlrp1b</i> ^s BALB/cJ, CBA/J, C3H/HeJ, FVB/NJ	<i>Nlrp1b</i> ^r C57BL/6J, NOD/LtJ, AKR/J, A/J, DBA/2J	²¹
Signal regulatory protein alpha (<i>Sirpa</i>)	T cell immune regulation, "don't eat me" signal	Decreased function polymorphism BALB/c, NOD/ShiLtJ	Most other strains	²²
Signaling lymphocytic activation molecule (SLAM) family	Innate and adaptive immunity, self tolerance	<i>Haplotype 2</i> NOD, MRL/MpJ, NZB, NZW, DBA/2J, 129/SvJ, BALB/cJ, C3H/HeJ, others	<i>Haplotype 1</i> C57BL/6, C3H/HeN	²³⁻²⁵
Solute carrier family-1 1a member 1 (<i>Slc11a1</i>)	Macrophage activation and iron metabolism in macrophages	<i>Slc11a1</i> ^s BALB/c, C57BL/6J, C57BL/10J, DBA/1J, NZW	<i>Slc11a1</i> ^r 129 strains, A/J, AKR/J, CBA/C3H/HeJ, C3H/HeN, DBA/2J, NZB	²⁶
T cell receptor beta variable 8 (<i>Tcrb-V8</i>)	NKT and CD8+NKT, glycolipids in context of CD1d	SJL/J, FVB/NJ, SWR, C57L, C57BR	C57BL/6, 129, BALB/c	²⁷⁻²⁹
Toll-like receptor 4 (<i>Tlr4</i>)	Microbial PAMPs (LPS), endogenous DAMPs	<i>Tlr4</i> ^{Lps-d} , <i>Tlr4</i> ^{Lps-del} C3H/HeJ, C57BL/10ScN	<i>Tlr4</i> ^{Lps-n} 129 strains, C3H/HeN, C3H/HeOuJ, C57BL/6, DBA/2J, FVB, NOD/ShiLtJ, NZB, NZW, SJ/J, SWR	^{30,31}

Abbreviations and acronyms used in the table:

Cyp450: cytochrome p450, DAMPs: damage-associated molecular pattern molecules, DC: dendritic cell, IFN: interferon, IL: interleukin, MHC: major histocompatibility complex, NK: natural killer, NKT: natural killer T, PAMPs: pathogen-associated molecular pattern molecules, LPS: lipopolysaccharide, LT: lethal toxin.

References

1. Poland A, Glover E. Characterization and strain distribution pattern of the murine ah receptor specified by the ahd and ahb-3 alleles. *Mol Pharmacol.* 1990;38(0026-895; 3):306-312.
2. De Souza VR, Cabrera WK, Galvan A, et al. Aryl hydrocarbon receptor polymorphism modulates DMBA-induced inflammation and carcinogenesis in phenotypically selected mice. *Int J Cancer.* 2009;124(6):1478-1482.
3. Esser C, Rannug A, Stockinger B. The aryl hydrocarbon receptor in immunity. *Trends Immunol.* 2009;30(9):447-454.
4. Esser C. The immune phenotype of AhR null mouse mutants: Not a simple mirror of xenobiotic receptor over-activation. *Biochem Pharmacol.* 2009;77:597-607. <http://www.sciencedirect.com/science/article/pii/S0006295208007193>.
5. Tulone C, Tsang J, Prokopowicz Z, Grosvenor N, Chain B. Natural cathepsin E deficiency in the immune system of C57BL/6J mice. . 2007;59(12):927-935.
6. Mahajan VS, Demissie E, Mattoo H, et al. Striking immune phenotypes in gene-targeted mice are driven by a copy-number variant originating from a commercially available C57BL/6 strain. *Cell Rep.* 2016;15(9):1901-1909. Accessed Dec 13, 2017. doi: 10.1016/j.celrep.2016.04.080.
7. Purtha WE, Swiecki M, Colonna M, Diamond MS, Bhattacharya D. Spontaneous mutation of the Dock2 gene in Irf5-/- mice complicates interpretation of type I interferon production and antibody responses. *Proc Natl Acad Sci U S A.* 2012;109(15):898. Accessed Dec 13, 2017. doi: 10.1073/pnas.1118155109.
8. Wetsel RA, Fleischer DT, Haviland DL. Deficiency of the murine fifth complement component (C5). A 2-base pair gene deletion in a 5'-exon. *J Biol Chem.* 1990;265(5):2435-2440.
9. Skerka C, Chen Q, Fremeaux-Bacchi V, Roumenina LT. Complement factor H related proteins (CFHRs). *Mol Immunol.* 2013;56:170-180. <http://www.sciencedirect.com/science/article/pii/S0161589013004185>.
10. Ford JW, Sturgill JL, Conrad DH. 129/SvJ mice have mutated CD23 and hyper IgE. *Cell Immunol.* 2009;254(2):124-134. Accessed Mar 19, 2018. doi: 10.1016/j.cellimm.2008.08.003.
11. Lewis G, Rapsomaniki E, Bouriez T, et al. Hyper IgE in new zealand black mice due to a dominant-negative CD23 mutation. *Immunogenetics.* 2004;56(8):564-571. Accessed Mar 19, 2018. doi: 10.1007/s00251-004-0728-4.

12. Encinas JA, Wicker LS, Peterson LB, et al. QTL influencing autoimmune diabetes and encephalomyelitis map to a 0.15-cM region containing IL2. *Nat Genet.* 1999;21(2):158-160.
13. Javan MR, Shahraki S, Safa A, Zamani MR, Salmanejad A, Aslani S. An interleukin 12 B single nucleotide polymorphism increases IL-12p40 production and is associated with increased disease susceptibility in patients with relapsing-remitting multiple sclerosis. *Neurol Res.* 2017;39(5):435-441.
14. Wen X, Chen S, Li P, et al. Single nucleotide polymorphisms of IL12B are associated with takayasu arteritis in chinese han population. *Rheumatol Int.* 2017;37(4):547-555.
15. Zwiers A, Fuss IJ, Seegers D, et al. A polymorphism in the coding region of IL12b promotes IL-12p70 and IL-23 heterodimer formation. *J Immunol.* 2011;2011/02/16:3572-3580.
16. Staeheli P, Grob R, Meier E, Sutcliffe JG, Haller O. Influenza virus-susceptible mice carry mx genes with a large deletion or a nonsense mutation. *Mol Cell Biol.* 1988;8:4518-4523. <http://mcb.asm.org/content/8/10/4518.abstract>.
17. Verhelst J, Spitaels J, Nurnberger C, et al. Functional comparison of Mx1 from two different mouse species reveals the involvement of loop L4 in the antiviral activity against influenza A viruses. *J Virol.* 2015;89(0022-538; 21):10879-10890.
18. Lightfield KL, Persson J, Brubaker SW, et al. Critical function for Naip5 in inflammasome activation by a conserved carboxy-terminal domain of flagellin. *Nat Immunol.* 2008;9(10):1171-1178.
19. Tenthorey JL, Haloupek N, Lopez-Blanco JR, et al. The structural basis of flagellin detection by NAIP5: A strategy to limit pathogen immune evasion. *Science.* 2017;358(6365):888-893.
20. Courtney SC, Di H, Stockman BM, Liu H, Scherbik SV, Brinton MA. Identification of novel host cell binding partners of Oas1b, the protein conferring resistance to flavivirus-induced disease in mice. *J Virol.* 2012;86(0022-538; 15):7953-7963.
21. Moayeri M, Crown D, Newman ZL, et al. Inflammasome sensor Nlrp1b-dependent resistance to anthrax is mediated by caspase-1, IL-1 signaling and neutrophil recruitment. *PLoS Pathogens.* 2010;6(2):e1001222. <http://www.ncbi.nlm.nih.gov/pubmed/21170303>. doi: 10.1371/journal.ppat.1001222.
22. Kwong LS, Brown MH, Barclay AN, Hatherley D. Signal-regulatory protein alpha from the NOD mouse binds human CD47 with an exceptionally high affinity-- implications for engraftment of human cells. *Immunology.* 2014;143(1):61-67.

23. Keszei M, Latchman YE, Vanguri VK, et al. Auto-antibody production and glomerulonephritis in congenic Slamf1-/- and Slamf2-/- B6.129] but not in Slamf1-/- and Slamf2-/- BALB/c.129] mice. *Int Immunol.* 2011;23(2):149-158.
24. Koh AE, Njoroge SW, Feliu M, et al. The SLAM family member CD48 (Slamf2) protects lupus-prone mice from autoimmune nephritis. *J Autoimmun.* 2011;37:48-57. <http://www.ncbi.nlm.nih.gov/pubmed/21561736>.
25. Crampton SP, Morawski PA, Bolland S. Linking susceptibility genes and pathogenesis mechanisms using mouse models of systemic lupus erythematosus. *Disease Models & Mechanisms.* 2014;7:1033-1046.
<http://www.ncbi.nlm.nih.gov/pmc/articles/PMC4142724/>.
26. Stober CB, Brode S, White JK, Popoff JF, Blackwell JM. Slc11a1, formerly Nramp1, is expressed in dendritic cells and influences major histocompatibility complex class II expression and antigen-presenting cell function. *Infect Immun.* 2007;75(10):5059-5067.
27. Osman GE, Hannibal MC, Anderson JP, Lasky SR, Ladiges WC, Hood L. FVB/N (H2(q)) mouse is resistant to arthritis induction and exhibits a genomic deletion of T-cell receptor V beta gene segments. . 1999;49(10):851-859.
28. Osman GE, Hannibal MC, Anderson JP, et al. T-cell receptor vbeta deletion and valpha polymorphism are responsible for the resistance of SWR mouse to arthritis induction. . 1999;49(9):764-772.
29. Behlke MA, Chou HS, Huppi K, Loh DY. Murine T-cell receptor mutants with deletions of beta-chain variable region genes. *Proc Natl Acad Sci U S A.* 1986;83(3):767-771.
30. Poltorak A, He X, Smirnova I, et al. Defective LPS signaling in C3H/HeJ and C57BL/10ScCr mice: Mutations in Tlr4 gene. *Science.* 1998;282(5396):2085-2088.
31. Poltorak A, Smirnova I, He X, et al. Genetic and physical mapping of the lps locus: Identification of the toll-4 receptor as a candidate gene in the critical region. *Blood Cells Mol Dis.* 1998;24(3):340-355.