# Betty P Tsao

List of Publications by Year
in descending order



| 140 | 10,699 | 32410 | 37326 |
|---|---|---|---|
| papers | citations | 55 h-index | 100 g-index |

| 163 | 163 | 163 | 11944 |
|---|---|---|---|
| all docs | docs citations | times ranked | citing authors |

| # | Article | IF | Citations |
|---|---|---|---|
| 1 | Human SLE variant NCF1-R90H promotes kidney damage and murine lupus through enhanced Tfh2 responses induced by defective efferocytosis of macrophages. Annals of the Rheumatic Diseases, 2022, 81, 255-267. | 0.5 | 14 |
| 2 | <scp>Upâ€Regulated</scp> Interleukinâ€10 Induced by <scp>E2F</scp> Transcription Factor 2â€“<scp>MicroRNA</scp>â€17â€5p Circuitry in Extrafollicular Effector B Cells Contributes to Autoantibody Production in Systemic Lupus Erythematosus. Arthritis and Rheumatology, 2022, 74, 496-507. | 2.9 | 12 |
| 3 | Prediction models of treatment response in lupus nephritis. Kidney International, 2022, 101, 379-389. | 2.6 | 18 |
| 4 | RNASE2 Mediates Age-Associated B Cell Expansion Through Monocyte Derived IL-10 in Patients With Systemic Lupus Erythematosus. Frontiers in Immunology, 2022, 13, 752189. | 2.2 | 9 |
| 5 | Complement C4, the Major Histocompatibility Complex, and Autoimmunity. Arthritis and Rheumatology, 2022, 74, 1318-1320. | 2.9 | 4 |
| 6 | Genes and genetics in human SLE. , 2021, , 85-96. | | 2 |
| 7 | IFNL4 Genotype Does Not Associate with CD4 T-Cell Recovery in People Living with Human Immunodeficiency Virus. AIDS Research and Human Retroviruses, 2021, 37, 184-188. | 0.5 | 2 |
| 8 | Lupus susceptibility genes. , 2021, , 25-33. | | 0 |
| 9 | Deep sequencing reveals a DAP1 regulatory haplotype that potentiates autoimmunity in systemic lupus erythematosus. Genome Biology, 2020, 21, 281. | 3.8 | 8 |
| 10 | Reduced Let-7f in Bone Marrow-Derived Mesenchymal Stem Cells Triggers Treg/Th17 Imbalance in Patients With Systemic Lupus Erythematosus. Frontiers in Immunology, 2020, 11, 233. | 2.2 | 30 |
| 11 | Rigorous Plasma Microbiome Analysis Method Enables Disease Association Discovery in Clinic. Frontiers in Microbiology, 2020, 11, 613268. | 1.5 | 12 |
| 12 | Amino acid signatures of HLA Class-I and II molecules are strongly associated with SLE susceptibility and autoantibody production in Eastern Asians. PLoS Genetics, 2019, 15, e1008092. | 1.5 | 36 |
| 13 | 180â€…Examining the transcriptional impact of liganded estrogen receptor alpha in the inflammatory milieu of systemic lupus erythematosus. , 2019, , . | | 0 |
| 14 | Genetics of Human SLE. , 2019, , 54-68. | | 5 |
| 15 | A plausibly causal functional lupus-associated risk variant in the STAT1â€“STAT4 locus. Human Molecular Genetics, 2018, 27, 2392-2404. | 1.4 | 34 |
| 16 | GG-07â€…Regulatory polymorphisms in EMSY gene are associated with autoantibodies in healthy individuals. , 2018, , . | | 0 |
| 17 | Genetic contributions to lupus nephritis in a multi-ethnic cohort of systemic lupus erythematous patients. PLoS ONE, 2018, 13, e0199003. | 1.1 | 46 |
| 18 | Transcription Factor SOX5 Promotes the Migration and Invasion of Fibroblast-Like Synoviocytes in Part by Regulating MMP-9 Expression in Collagen-Induced Arthritis. Frontiers in Immunology, 2018, 9, 749. | 2.2 | 33 |

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| 19 | Genetic variants in systemic lupus erythematosus susceptibility loci, XKR6 and GLT1D1 are associated with childhood-onset SLE in a Korean cohort. Scientific Reports, 2018, 8, 9962. | 1.6 | 25 |
| 20 | Genetic fine mapping of systemic lupus erythematosus MHC associations in Europeans and African Americans. Human Molecular Genetics, 2018, 27, 3813-3824. | 1.4 | 43 |
| 21 | A missense variant in NCF1 is associated with susceptibility to multiple autoimmune diseases. Nature Genetics, 2017, 49, 433-437. | 9.4 | 143 |
| 22 | Updates in Lupus Genetics. Current Rheumatology Reports, 2017, 19, 68. | 2.1 | 99 |
| 23 | Transancestral mapping and genetic load in systemic lupus erythematosus. Nature Communications, 2017, 8, 16021. | 5.8 | 314 |
| 24 | X Chromosome Dose and Sex Bias in Autoimmune Diseases: Increased Prevalence of 47,XXX in Systemic Lupus Erythematosus and SjÃ¶gren's Syndrome. Arthritis and Rheumatology, 2016, 68, 1290-1300. | 2.9 | 114 |
| 25 | Identification of a Systemic Lupus Erythematosus Risk Locus Spanning ATG16L2, FCHSD2, and P2RY2 in Koreans. Arthritis and Rheumatology, 2016, 68, 1197-1209. | 2.9 | 89 |
| 26 | Regulatory polymorphisms modulate the expression of HLA class II molecules and promote autoimmunity. ELife, 2016, 5, . | 2.8 | 113 |
| 27 | Genes and Genetics in Human Systemic Lupus Erythematosus. , 2016, , 69-76. | | 1 |
| 28 | Genomeâ€Wide Association Study in an Amerindian Ancestry Population Reveals Novel Systemic Lupus Erythematosus Risk Loci and the Role of European Admixture. Arthritis and Rheumatology, 2016, 68, 932-943. | 2.9 | 138 |
| 29 | Modulation of IL-6 induced RANKL expression in arthritic synovium by a transcription factor SOX5. Scientific Reports, 2016, 6, 32001. | 1.6 | 41 |
| 30 | CD3Zhypermethylation is associated with severe clinical manifestations in systemic lupus erythematosus and reduces CD3Î¶-chain expression in T cells. Rheumatology, 2016, 56, kew405. | 0.9 | 12 |
| 31 | Decreased SMG7 expression associates with lupus-risk variants and elevated antinuclear antibody production. Annals of the Rheumatic Diseases, 2016, 75, 2007-2013. | 0.5 | 16 |
| 32 | Preferential association of a functional variant in complement receptor 2 with antibodies to double-stranded DNA. Annals of the Rheumatic Diseases, 2016, 75, 242-252. | 0.5 | 10 |
| 33 | Focused transcription from the human CR2/CD21 core promoter is regulated by synergistic activity of TATA and Initiator elements in mature B cells. Cellular and Molecular Immunology, 2016, 13, 119-131. | 4.8 | 3 |
| 34 | Lupus risk variants in the PXK locus alter B-cell receptor internalization. Frontiers in Genetics, 2015, 5, 450. | 1.1 | 25 |
| 35 | Lupus Risk Variant Increases pSTAT1 Binding and Decreases ETS1 Expression. American Journal of Human Genetics, 2015, 96, 731-739. | 2.6 | 36 |
| 36 | Genetic associations of leptin-related polymorphisms with systemic lupus erythematosus. Clinical Immunology, 2015, 161, 157-162. | 1.4 | 10 |

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| 37 | The IRF5â€“TNPO3 association with systemic lupus erythematosus has two components that other autoimmune disorders variably share. Human Molecular Genetics, 2015, 24, 582-596. | 1.4 | 74 |
| 38 | Identification of interferon-inducible genes as diagnostic biomarker for systemic lupus erythematosus. Clinical Rheumatology, 2015, 34, 71-79. | 1.0 | 43 |
| 39 | Restored Immunosuppressive Effect of Mesenchymal Stem Cells on B Cells After Olfactory 1/Early B Cell Factorâ€“Associated Zincâ€Finger Protein Downâ€Regulation in Patients With Systemic Lupus Erythematosus. Arthritis and Rheumatology, 2014, 66, 3413-3423. | 2.9 | 35 |
| 40 | Genetics of systemic lupus erythematosus: immune responses and end organ resistance to damage. Current Opinion in Immunology, 2014, 31, 87-96. | 2.4 | 47 |
| 41 | Advances in lupus genetics and epigenetics. Current Opinion in Rheumatology, 2014, 26, 482-492. | 2.0 | 104 |
| 42 | Endâ€Stage Renal Disease in African Americans With Lupus Nephritis Is Associated With APOL1. Arthritis and Rheumatology, 2014, 66, 390-396. | 2.9 | 242 |
| 43 | Two Functional Lupus-Associated BLK Promoter Variants Control Cell-Type- and Developmental-Stage-Specific Transcription. American Journal of Human Genetics, 2014, 94, 586-598. | 2.6 | 59 |
| 44 | Lupus Nephritis Susceptibility Loci in Women with Systemic Lupus Erythematosus. Journal of the American Society of Nephrology: JASN, 2014, 25, 2859-2870. | 3.0 | 117 |
| 45 | Transcription factor Ikaros Represses Protein Phosphatase 2A (PP2A) Expression through an Intronic Binding Site. Journal of Biological Chemistry, 2014, 289, 13751-13757. | 1.6 | 20 |
| 46 | Systemic Lupus Erythematosus, Genetics. , 2014, , 1171-1178. | | 0 |
| 47 | Plasma levels of osteopontin identify patients at risk for organ damage in systemic lupus erythematosus. Arthritis Research and Therapy, 2013, 15, R18. | 1.6 | 32 |
| 48 | Brief Report: Singleâ€nucleotide polymorphisms in VKORC1 are risk factors for systemic lupus erythematosus in Asians. Arthritis and Rheumatism, 2013, 65, 211-215. | 6.7 | 10 |
| 49 | Variable Association of Reactive Intermediate Genes with Systemic Lupus Erythematosus in Populations with Different African Ancestry. Journal of Rheumatology, 2013, 40, 842-849. | 1.0 | 15 |
| 50 | Preferential Binding to Elk-1 by SLE-Associated IL10 Risk Allele Upregulates IL10 Expression. PLoS Genetics, 2013, 9, e1003870. | 1.5 | 36 |
| 51 | Admixture Mapping in Lupus Identifies Multiple Functional Variants within IFIH1 Associated with Apoptosis, Inflammation, and Autoantibody Production. PLoS Genetics, 2013, 9, e1003222. | 1.5 | 107 |
| 52 | Trans-Ancestral Studies Fine Map the SLE-Susceptibility Locus TNFSF4. PLoS Genetics, 2013, 9, e1003554. | 1.5 | 50 |
| 53 | MicroRNA-3148 Modulates Allelic Expression of Toll-Like Receptor 7 Variant Associated with Systemic Lupus Erythematosus. PLoS Genetics, 2013, 9, e1003336. | 1.5 | 107 |
| 54 | Fine mapping of Xq28: bothMECP2 and IRAK1contribute to risk for systemic lupus erythematosus in multiple ancestral groups. Annals of the Rheumatic Diseases, 2013, 72, 437-444. | 0.5 | 97 |

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| 55 | Recent insights into the genetic basis of systemic lupus erythematosus. Annals of the Rheumatic Diseases, 2013, 72, ii56-ii61. | 0.5 | 117 |
| 56 | ABIN1 Dysfunction as a Genetic Basis for Lupus Nephritis. Journal of the American Society of Nephrology: JASN, 2013, 24, 1743-1754. | 3.0 | 70 |
| 57 | Genetics of Human SLE. , 2013, , 35-45. | | 4 |
| 58 | PTPN22 Association in Systemic Lupus Erythematosus (SLE) with Respect to Individual Ancestry and Clinical Sub-Phenotypes. PLoS ONE, 2013, 8, e69404. | 1.1 | 57 |
| 59 | Novel identification of the IRF7 region as an anticentromere autoantibody propensity locus in systemic sclerosis. Annals of the Rheumatic Diseases, 2012, 71, 114-119. | 0.5 | 62 |
| 60 | Analysis of autosomal genes reveals geneâ€“sex interactions and higher total genetic risk in men with systemic lupus erythematosus. Annals of the Rheumatic Diseases, 2012, 71, 694-699. | 0.5 | 87 |
| 61 | Impact of genetic ancestry and sociodemographic status on the clinical expression of systemic lupus erythematosus in American Indianâ€“European populations. Arthritis and Rheumatism, 2012, 64, 3687-3694. | 6.7 | 70 |
| 62 | Association of two independent functional risk haplotypes in TNIP1 with systemic lupus erythematosus. Arthritis and Rheumatism, 2012, 64, 3695-3705. | 6.7 | 69 |
| 63 | Variation in the ICAM1â€“ICAM4â€“ICAM5 locus is associated with systemic lupus erythematosus susceptibility in multiple ancestries. Annals of the Rheumatic Diseases, 2012, 71, 1809-1814. | 0.5 | 60 |
| 64 | Evaluation of TRAF6 in a large multiancestral lupus cohort. Arthritis and Rheumatism, 2012, 64, 1960-1969. | 6.7 | 51 |
| 65 | Identification of IRF8, TMEM39A, and IKZF3-ZPBP2 as Susceptibility Loci for Systemic Lupus Erythematosus in a Large-Scale Multiracial Replication Study. American Journal of Human Genetics, 2012, 90, 648-660. | 2.6 | 161 |
| 66 | Transcriptional effects of a lupus-associated polymorphism in the 5â€² untranslated region (UTR) of human complement receptor 2 (CR2/CD21). Molecular Immunology, 2012, 52, 165-173. | 1.0 | 12 |
| 67 | Evidence for geneâ€“gene epistatic interactions among susceptibility loci for systemic lupus erythematosus. Arthritis and Rheumatism, 2012, 64, 485-492. | 6.7 | 53 |
| 68 | Inhibition of Aberrant Circulating Tfh Cell Proportions by Corticosteroids in Patients with Systemic Lupus Erythematosus. PLoS ONE, 2012, 7, e51982. | 1.1 | 91 |
| 69 | Differential Genetic Associations for Systemic Lupus Erythematosus Based on Antiâ€“dsDNA Autoantibody Production. PLoS Genetics, 2011, 7, e1001323. | 1.5 | 206 |
| 70 | Association of a functional variant downstream of TNFAIP3 with systemic lupus erythematosus. Nature Genetics, 2011, 43, 253-258. | 9.4 | 242 |
| 71 | Identification of a Systemic Lupus Erythematosus Susceptibility Locus at 11p13 between PDHX and CD44 in a Multiethnic Study. American Journal of Human Genetics, 2011, 88, 83-91. | 2.6 | 72 |
| 72 | Association of a functional IRF7 variant with systemic lupus erythematosus. Arthritis and Rheumatism, 2011, 63, 749-754. | 6.7 | 118 |

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| 73 | Genetic analyses of interferon pathway-related genes reveal multiple new loci associated with systemic lupus erythematosus. Arthritis and Rheumatism, 2011, 63, 2049-2057. | 6.7 | 45 |
| 74 | Association of PPP2CA polymorphisms with systemic lupus erythematosus susceptibility in multiple ethnic groups. Arthritis and Rheumatism, 2011, 63, 2755-2763. | 6.7 | 36 |
| 75 | Identification of novel genetic susceptibility loci in African American lupus patients in a candidate gene association study. Arthritis and Rheumatism, 2011, 63, 3493-3501. | 6.7 | 109 |
| 76 | A Functional Variant in MicroRNA-146a Promoter Modulates Its Expression and Confers Disease Risk for Systemic Lupus Erythematosus. PLoS Genetics, 2011, 7, e1002128. | 1.5 | 241 |
| 77 | Risk Alleles for Systemic Lupus Erythematosus in a Large Case-Control Collection and Associations with Clinical Subphenotypes. PLoS Genetics, 2011, 7, e1001311. | 1.5 | 154 |
| 78 | A Comprehensive Analysis of Shared Loci between Systemic Lupus Erythematosus (SLE) and Sixteen Autoimmune Diseases Reveals Limited Genetic Overlap. PLoS Genetics, 2011, 7, e1002406. | 1.5 | 148 |
| 79 | Phenotypic associations of genetic susceptibility loci in systemic lupus erythematosus. Annals of the Rheumatic Diseases, 2011, 70, 1752-1757. | 0.5 | 110 |
| 80 | Constitutive Genes and Lupus. , 2011, , 47-61. | | 4 |
| 81 | Association of Genetic Variants in Complement Factor H and Factor H-Related Genes with Systemic Lupus Erythematosus Susceptibility. PLoS Genetics, 2011, 7, e1002079. | 1.5 | 181 |
| 82 | A functional RANKL polymorphism associated with younger age at onset of rheumatoid arthritis. Arthritis and Rheumatism, 2010, 62, 2864-2875. | 6.7 | 35 |
| 83 | Association of IRF5 polymorphisms with activation of the interferon Î± pathway. Annals of the Rheumatic Diseases, 2010, 69, 611-617. | 0.5 | 54 |
| 84 | Sex-specific association of X-linked Toll-like receptor 7 (TLR7) with male systemic lupus erythematosus. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 15838-15843. | 3.3 | 324 |
| 85 | Male-only Systemic Lupus. Journal of Rheumatology, 2010, 37, 1480-1487. | 1.0 | 13 |
| 86 | Treatment with apolipoprotein A-1 mimetic peptide reduces lupus-like manifestations in a murine lupus model of accelerated atherosclerosis. Arthritis Research and Therapy, 2010, 12, R93. | 1.6 | 47 |
| 87 | Genetic susceptibility to systemic lupus erythematosus in the genomic era. Nature Reviews Rheumatology, 2010, 6, 683-692. | 3.5 | 319 |
| 88 | Olf1/EBF associated zinc finger protein interfered with antinuclear antibody production in patients with systemic lupus erythematosus. Arthritis Research and Therapy, 2010, 12, R59. | 1.6 | 8 |
| 89 | European population substructure is associated with mucocutaneous manifestations and autoantibody production in systemic lupus erythematosus. Arthritis and Rheumatism, 2009, 60, 2448-2456. | 6.7 | 27 |
| 90 | Plasmin immunization preferentially induces potentially prothrombotic IgG anticardiolipin antibodies in MRL/MpJ mice. Arthritis and Rheumatism, 2009, 60, 3108-3117. | 6.7 | 2 |

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| 91 | Pathogenesis of Systemic Lupus Erythematosus. , 2009, , 1233-1262. | | 4 |
| 92 | Genome-wide association scan in women with systemic lupus erythematosus identifies susceptibility variants in ITGAM, PXK, KIAA1542 and other loci. Nature Genetics, 2008, 40, 204-210. | 9.4 | 1,192 |
| 93 | A loss-of-function variant of PTPN22 is associated with reduced risk of systemic lupus erythematosus. Human Molecular Genetics, 2008, 18, 569-579. | 1.4 | 106 |
| 94 | A genome wide association study of systemic lupus erythematosus (SLE) by SLEGEN, the International SLE Genetics Consortium.. FASEB Journal, 2008, 22, 850.1. | 0.2 | 0 |
| 95 | Association of a common complement receptor 2 haplotype with increased risk of systemic lupus erythematosus. Proceedings of the National Academy of Sciences of the United States of America, 2007, 104, 3961-3966. | 3.3 | 62 |
| 96 | ApoEâ^'/â^'Fasâ^'/â^' C57BL/6 mice: a novel murine model simultaneously exhibits lupus nephritis, atherosclerosis, and osteopenia. Journal of Lipid Research, 2007, 48, 794-805. | 2.0 | 62 |
| 97 | Gene Copy-Number Variation and Associated Polymorphisms of Complement Component C4 in Human Systemic Lupus Erythematosus (SLE): Low Copy Number Is a Risk Factor for and High Copy Number Is a Protective Factor against SLE Susceptibility in European Americans. American Journal of Human Genetics, 2007, 80, 1037-1054. | 2.6 | 411 |
| 98 | Current topics in human SLE genetics. Seminars in Immunopathology, 2006, 28, 97-107. | 4.0 | 61 |
| 99 | Association of tumor necrosis factor Î± polymorphism, but not the shared epitope, with increased radiographic progression in a seropositive rheumatoid arthritis inception cohort. Arthritis and Rheumatism, 2006, 54, 1105-1116. | 6.7 | 49 |
| 100 | Association of increased interferon-inducible gene expression with disease activity and lupus nephritis in patients with systemic lupus erythematosus. Arthritis and Rheumatism, 2006, 54, 2951-2962. | 6.7 | 404 |
| 101 | Association analysis of the R620W polymorphism of protein tyrosine phosphatase PTPN22 in systemic lupus erythematosus families: Increased t allele frequency in systemic lupus erythematosus patients with autoimmune thyroid disease. Arthritis and Rheumatism, 2005, 52, 2396-2402. | 6.7 | 80 |
| 102 | Maternal HLA class II compatibility in men with systemic lupus erythematosus. Arthritis and Rheumatism, 2005, 52, 2768-2773. | 6.7 | 36 |
| 103 | A stop codon polymorphism of Toll-like receptor 5 is associated with resistance to systemic lupus erythematosus. Proceedings of the National Academy of Sciences of the United States of America, 2005, 102, 10593-10597. | 3.3 | 144 |
| 104 | CD72 polymorphisms associated with alternative splicing modify susceptibility to human systemic lupus erythematosus through epistatic interaction with FCGR2B. Human Molecular Genetics, 2004, 13, 2907-2917. | 1.4 | 43 |
| 105 | Current advances in the human lupus genetics. Current Rheumatology Reports, 2004, 6, 391-398. | 2.1 | 35 |
| 106 | Association of Fc? receptor IIA, but not IIB and IIIA, polymorphisms with systemic lupus erythematosus: A family-based association study in Caucasians. Arthritis and Rheumatism, 2004, 50, 671-673. | 6.7 | 34 |
| 107 | Systemic lupus erythematosus genome scan: Support for linkage at 1q23, 2q33, 16q12-13, and 17q21-23 and novel evidence at 3p24, 10q23-24, 13q32, and 18q22-23. Arthritis and Rheumatism, 2004, 50, 3203-3210. | 6.7 | 66 |
| 108 | Interaction betweenRANKL andHLA-DRB1 genotypes may contribute to younger age at onset of seropositive rheumatoid arthritis in an inception cohort. Arthritis and Rheumatism, 2004, 50, 3093-3103. | 6.7 | 42 |

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| 109 | Update on human systemic lupus erythematosus genetics. Current Opinion in Rheumatology, 2004, 16, 513-521. | 2.0 | 139 |
| 110 | Identification and characterization of SmD183-119-reactive T cells that provide T cell help for pathogenic anti-double-stranded DNA antibodies. Arthritis and Rheumatism, 2003, 48, 475-485. | 6.7 | 216 |
| 111 | Identification and characterization of a peptide mimetic that may detect a species of disease-associated anticardiolipin antibodies in patients with the antiphospholipid syndrome. Arthritis and Rheumatism, 2003, 48, 737-745. | 6.7 | 14 |
| 112 | The genetics of human systemic lupus erythematosus. Trends in Immunology, 2003, 24, 595-602. | 2.9 | 165 |
| 113 | Familiality and co-occurrence of clinical features of systemic lupus erythematosus. Arthritis and Rheumatism, 2002, 46, 2678-2685. | 6.7 | 51 |
| 114 | Linkage and interaction of loci on 1q23 and 16q12 may contribute to susceptibility to systemic lupus erythematosus. Arthritis and Rheumatism, 2002, 46, 2928-2936. | 6.7 | 55 |
| 115 | An update on genetic studies of systemic lupus erythematosus. Current Rheumatology Reports, 2002, 4, 359-367. | 2.1 | 46 |
| 116 | FcÎ³ receptor IIIA polymorphism in Korean patients with systemic lupus erythematosus. Rheumatology International, 2002, 21, 222-226. | 1.5 | 17 |
| 117 | Genetics and systemic lupus erythematosus. Current Rheumatology Reports, 2001, 3, 183-190. | 2.1 | 22 |
| 118 | Treatment with a consensus peptide based on amino acid sequences in autoantibodies prevents T cell activation by autoantigens and delays disease onset in murine lupus. Arthritis and Rheumatism, 2001, 44, 432-441. | 6.7 | 103 |
| 119 | Poly(ADP)â€ribose polymerase and susceptibility to systemic lupus erythematosus and primary antiphospholipid syndrome: Comment on the article by Delrieu et al. Arthritis and Rheumatism, 2000, 43, 1421-1422. | 6.7 | 10 |
| 120 | Genetics and systemic lupus erythematosus. Current Rheumatology Reports, 2000, 2, 13-18. | 2.1 | 14 |
| 121 | Single-nucleotide polymorphisms of T cell receptor ? chain in patients with systemic lupus erythematosus. Arthritis and Rheumatism, 1999, 42, 2601-2605. | 6.7 | 29 |
| 122 | PARP alleles within the linked chromosomal region are associated with systemic lupus erythematosus. Journal of Clinical Investigation, 1999, 103, 1135-1140. | 3.9 | 99 |
| 123 | Autoantibodies as a Source of Peptides That Regulate Autoantibody Production. , 1999, , 371-388. | | 0 |
| 124 | Abnormal distribution of Fc? receptor type IIa polymorphisms in Korean patients with systemic lupus erythematosus. Arthritis and Rheumatism, 1998, 41, 421-426. | 6.7 | 92 |
| 125 | Altered Immune Responses in Interleukin 10 Transgenic Mice. Journal of Experimental Medicine, 1997, 185, 2101-2110. | 4.2 | 261 |
| 126 | Commentary: Genetics of systemic lupus erythematosus. Current Opinion in Rheumatology, 1997, 9, 377-379. | 2.0 | 10 |

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| 127 | Autoimmunity and Tolerance in Ig-Transgenic Mice: Murine SLE as a Model to Study B Cell Tolerance. International Reviews of Immunology, 1994, 11, 305-320. | 1.5 | 0 |
| 128 | Comparison of pathogenic and non-pathogenic murine antibodies to DNA: antigen binding and structural characteristics. International Immunology, 1994, 6, 817-830. | 1.8 | 145 |
| 129 | A peptide derived from an autoantibody can stimulate t cells in the (nzb Ã— nzw)f1 mouse model of systemic lupus erythematosus. Arthritis and Rheumatism, 1993, 36, 355-364. | 6.7 | 70 |
| 130 | B cells are anergic in transgenic mice that express IgM anti-DNA antibodies. European Journal of Immunology, 1993, 23, 2332-2339. | 1.6 | 48 |
| 131 | Contribution of Major Histocompatibility Complex (MHC) to Upregulation of Anti-DNA Antibody in Transgenic Mice. Journal of Autoimmunity, 1993, 6, 1-9. | 3.0 | 3 |
| 132 | Ig-transgenic mice as models for studying the regulation and role of anti-DNA antibodies in murine lupus. ImmunoMethods, 1992, 1, 185-190. | 0.8 | 2 |
| 133 | T cell up-regulation of B cells via their idiotypes contributing to the development of systemic lupus erythematosus. American Journal of Medicine, 1988, 85, 32-34. | 0.6 | 8 |
| 134 | Idiotype selection is an immunoregulatory mechanism which contributes to the pathogenesis of systemic lupus erythematosus. Journal of Autoimmunity, 1988, 1, 673-681. | 3.0 | 1 |
| 135 | In vivofunctional analysis ofin vitroprotein binding sites in the immunoglobulin heavy chain enhancer. Nucleic Acids Research, 1988, 16, 3239-3253. | 6.5 | 64 |
| 136 | The role of cytoplasmic free calcium concentration in B-cell tolerance. Cellular Immunology, 1987, 108, 335-342. | 1.4 | 7 |
| 137 | Macrophage-derived soluble factors mediate suppression induced by 2,4-dinitrophenyl-conjugated mouse IgG in hybridoma cells. Cellular Immunology, 1985, 91, 362-374. | 1.4 | 0 |
| 138 | Central suppression of monoclonal B cells: DNP-MGG suppresses proliferation and immunoglobulin synthesis in anti-DNP-secreting hybridoma and myeloma. Cellular Immunology, 1984, 88, 96-108. | 1.4 | 6 |
| 139 | Evidence that the hydrophobic domain of rat renal Î³-glutamyltransferase spans the brush border membrane. Biochimica Et Biophysica Acta - Biomembranes, 1982, 690, 199-206. | 1.4 | 5 |
| 140 | Membrane association and orientation of rat renal activities capable of degrading glutathione. International Journal of Biochemistry & Cell Biology, 1980, 12, 219-222. | 0.8 | 7 |