

Richard P Lifton

List of Publications by Citations

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

318
papers

54,103
citations

109
h-index

230
g-index

334
ext. papers

63,116
ext. citations

16
avg, IF

6.69
L-index

| # | Paper | IF | Citations |
|-----|--|------|-----------|
| 318 | Genetic studies of body mass index yield new insights for obesity biology. <i>Nature</i> , 2015 , 518, 197-206 | 50.4 | 2687 |
| 317 | Molecular basis of human hypertension: role of angiotensinogen. <i>Cell</i> , 1992 , 71, 169-80 | 56.2 | 1569 |
| 316 | De novo mutations revealed by whole-exome sequencing are strongly associated with autism. <i>Nature</i> , 2012 , 485, 237-41 | 50.4 | 1470 |
| 315 | High bone density due to a mutation in LDL-receptor-related protein 5. <i>New England Journal of Medicine</i> , 2002 , 346, 1513-21 | 59.2 | 1357 |
| 314 | Molecular mechanisms of human hypertension. <i>Cell</i> , 2001 , 104, 545-56 | 56.2 | 1323 |
| 313 | Human hypertension caused by mutations in WNK kinases. <i>Science</i> , 2001 , 293, 1107-12 | 33.3 | 1160 |
| 312 | Liddle β syndrome: heritable human hypertension caused by mutations in the beta subunit of the epithelial sodium channel. <i>Cell</i> , 1994 , 79, 407-14 | 56.2 | 1093 |
| 311 | Autoantibodies against type I IFNs in patients with life-threatening COVID-19. <i>Science</i> , 2020 , 370, | 33.3 | 1090 |
| 310 | Inborn errors of type I IFN immunity in patients with life-threatening COVID-19. <i>Science</i> , 2020 , 370, | 33.3 | 994 |
| 309 | A chimaeric 11 beta-hydroxylase/aldosterone synthase gene causes glucocorticoid-remediable aldosteronism and human hypertension. <i>Nature</i> , 1992 , 355, 262-5 | 50.4 | 992 |
| 308 | Genetic diagnosis by whole exome capture and massively parallel DNA sequencing. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009 , 106, 19096-101 | 11.5 | 971 |
| 307 | Gitelman β variant of Bartter β syndrome, inherited hypokalaemic alkalosis, is caused by mutations in the thiazide-sensitive Na-Cl cotransporter. <i>Nature Genetics</i> , 1996 , 12, 24-30 | 36.3 | 960 |
| 306 | Paracellin-1, a renal tight junction protein required for paracellular Mg ²⁺ resorption. <i>Science</i> , 1999 , 285, 103-6 | 33.3 | 941 |
| 305 | Multiple recurrent de novo CNVs, including duplications of the 7q11.23 Williams syndrome region, are strongly associated with autism. <i>Neuron</i> , 2011 , 70, 863-85 | 13.9 | 932 |
| 304 | New genetic loci link adipose and insulin biology to body fat distribution. <i>Nature</i> , 2015 , 518, 187-196 | 50.4 | 920 |
| 303 | Exome sequencing identifies recurrent somatic RAC1 mutations in melanoma. <i>Nature Genetics</i> , 2012 , 44, 1006-14 | 36.3 | 887 |
| 302 | Sequence variants in SLITRK1 are associated with Tourette β syndrome. <i>Science</i> , 2005 , 310, 317-20 | 33.3 | 776 |

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|-----|---|------|-----|
| 301 | BartterB syndrome, hypokalaemic alkalosis with hypercalciuria, is caused by mutations in the Na-K-2Cl cotransporter NKCC2. <i>Nature Genetics</i> , 1996 , 13, 183-8 | 36.3 | 734 |
| 300 | Mutations in the chloride channel gene, CLCNKB, cause BartterB syndrome type III. <i>Nature Genetics</i> , 1997 , 17, 171-8 | 36.3 | 718 |
| 299 | K+ channel mutations in adrenal aldosterone-producing adenomas and hereditary hypertension. <i>Science</i> , 2011 , 331, 768-72 | 33.3 | 707 |
| 298 | Genetic heterogeneity of BartterB syndrome revealed by mutations in the K+ channel, ROMK. <i>Nature Genetics</i> , 1996 , 14, 152-6 | 36.3 | 670 |
| 297 | Mutations in subunits of the epithelial sodium channel cause salt wasting with hyperkalaemic acidosis, pseudohypoaldosteronism type 1. <i>Nature Genetics</i> , 1996 , 12, 248-53 | 36.3 | 667 |
| 296 | Hypertension caused by a truncated epithelial sodium channel gamma subunit: genetic heterogeneity of Liddle syndrome. <i>Nature Genetics</i> , 1995 , 11, 76-82 | 36.3 | 648 |
| 295 | Rare independent mutations in renal salt handling genes contribute to blood pressure variation. <i>Nature Genetics</i> , 2008 , 40, 592-599 | 36.3 | 639 |
| 294 | De novo mutations in histone-modifying genes in congenital heart disease. <i>Nature</i> , 2013 , 498, 220-3 | 50.4 | 591 |
| 293 | Mutations in the gene encoding B1 subunit of H+-ATPase cause renal tubular acidosis with sensorineural deafness. <i>Nature Genetics</i> , 1999 , 21, 84-90 | 36.3 | 551 |
| 292 | Activating mineralocorticoid receptor mutation in hypertension exacerbated by pregnancy. <i>Science</i> , 2000 , 289, 119-23 | 33.3 | 521 |
| 291 | Genomic analysis of non-NF2 meningiomas reveals mutations in TRAF7, KLF4, AKT1, and SMO. <i>Science</i> , 2013 , 339, 1077-80 | 33.3 | 508 |
| 290 | Evidence for a gene influencing blood pressure on chromosome 17. Genome scan linkage results for longitudinal blood pressure phenotypes in subjects from the framingham heart study. <i>Hypertension</i> , 2000 , 36, 477-83 | 8.5 | 483 |
| 289 | LRP6 mutation in a family with early coronary disease and metabolic risk factors. <i>Science</i> , 2007 , 315, 1278-82 | 33.3 | 465 |
| 288 | Mutations in kelch-like 3 and cullin 3 cause hypertension and electrolyte abnormalities. <i>Nature</i> , 2012 , 482, 98-102 | 50.4 | 440 |
| 287 | The Genetic Basis of Mendelian Phenotypes: Discoveries, Challenges, and Opportunities. <i>American Journal of Human Genetics</i> , 2015 , 97, 199-215 | 11 | 432 |
| 286 | Molecular cytogenetic analysis and resequencing of contactin associated protein-like 2 in autism spectrum disorders. <i>American Journal of Human Genetics</i> , 2008 , 82, 165-73 | 11 | 427 |
| 285 | Genome-wide association study identifies susceptibility loci for IgA nephropathy. <i>Nature Genetics</i> , 2011 , 43, 321-7 | 36.3 | 414 |
| 284 | Somatic and germline CACNA1D calcium channel mutations in aldosterone-producing adenomas and primary aldosteronism. <i>Nature Genetics</i> , 2013 , 45, 1050-4 | 36.3 | 410 |

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|-----|--|------|-----|
| 283 | De novo mutations in congenital heart disease with neurodevelopmental and other congenital anomalies. <i>Science</i> , 2015 , 350, 1262-6 | 33.3 | 406 |
| 282 | Whole-exome sequencing identifies recessive WDR62 mutations in severe brain malformations. <i>Nature</i> , 2010 , 467, 207-10 | 50.4 | 395 |
| 281 | A single-gene cause in 29.5% of cases of steroid-resistant nephrotic syndrome. <i>Journal of the American Society of Nephrology: JASN</i> , 2015 , 26, 1279-89 | 12.7 | 378 |
| 280 | Recessive mutations in DGKE cause atypical hemolytic-uremic syndrome. <i>Nature Genetics</i> , 2013 , 45, 531-6 | 36.3 | 357 |
| 279 | Seizures, sensorineural deafness, ataxia, mental retardation, and electrolyte imbalance (SeSAME syndrome) caused by mutations in KCNJ10. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009 , 106, 5842-7 | 11.5 | 356 |
| 278 | Contribution of rare inherited and de novo variants in 2,871 congenital heart disease probands. <i>Nature Genetics</i> , 2017 , 49, 1593-1601 | 36.3 | 348 |
| 277 | Absence of linkage between the angiotensin converting enzyme locus and human essential hypertension. <i>Nature Genetics</i> , 1992 , 1, 72-5 | 36.3 | 343 |
| 276 | Molecular pathogenesis of inherited hypertension with hyperkalemia: the Na-Cl cotransporter is inhibited by wild-type but not mutant WNK4. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2003 , 100, 680-4 | 11.5 | 342 |
| 275 | Mutation of NLRC4 causes a syndrome of enterocolitis and autoinflammation. <i>Nature Genetics</i> , 2014 , 46, 1135-1139 | 36.3 | 337 |
| 274 | Mutations in the mineralocorticoid receptor gene cause autosomal dominant pseudohypoaldosteronism type I. <i>Nature Genetics</i> , 1998 , 19, 279-81 | 36.3 | 335 |
| 273 | Discovery of new risk loci for IgA nephropathy implicates genes involved in immunity against intestinal pathogens. <i>Nature Genetics</i> , 2014 , 46, 1187-96 | 36.3 | 325 |
| 272 | Mutations in ATP6N1B, encoding a new kidney vacuolar proton pump 116-kD subunit, cause recessive distal renal tubular acidosis with preserved hearing. <i>Nature Genetics</i> , 2000 , 26, 71-5 | 36.3 | 319 |
| 271 | WNK4 regulates the balance between renal NaCl reabsorption and K ⁺ secretion. <i>Nature Genetics</i> , 2003 , 35, 372-6 | 36.3 | 318 |
| 270 | Wnk4 controls blood pressure and potassium homeostasis via regulation of mass and activity of the distal convoluted tubule. <i>Nature Genetics</i> , 2006 , 38, 1124-32 | 36.3 | 298 |
| 269 | Roles of the cation-chloride cotransporters in neurological disease. <i>Nature Clinical Practice Neurology</i> , 2008 , 4, 490-503 | | 294 |
| 268 | Hereditary hypertension caused by chimaeric gene duplications and ectopic expression of aldosterone synthase. <i>Nature Genetics</i> , 1992 , 2, 66-74 | 36.3 | 288 |
| 267 | Multilocus linkage identifies two new loci for a mendelian form of stroke, cerebral cavernous malformation, at 7p15-13 and 3q25.2-27. <i>Human Molecular Genetics</i> , 1998 , 7, 1851-8 | 5.6 | 287 |
| 266 | A cluster of metabolic defects caused by mutation in a mitochondrial tRNA. <i>Science</i> , 2004 , 306, 1190-4 | 33.3 | 282 |

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|-----|---|------|-----|
| 265 | Exome sequencing links mutations in PARN and RTEL1 with familial pulmonary fibrosis and telomere shortening. <i>Nature Genetics</i> , 2015 , 47, 512-7 | 36.3 | 279 |
| 264 | Exome sequencing identifies recurrent mutations in NF1 and RASopathy genes in sun-exposed melanomas. <i>Nature Genetics</i> , 2015 , 47, 996-1002 | 36.3 | 261 |
| 263 | IgA nephropathy, the most common cause of glomerulonephritis, is linked to 6q22-23. <i>Nature Genetics</i> , 2000 , 26, 354-7 | 36.3 | 250 |
| 262 | Genomic landscape of cutaneous T cell lymphoma. <i>Nature Genetics</i> , 2015 , 47, 1011-9 | 36.3 | 247 |
| 261 | L-histidine decarboxylase and Tourette's syndrome. <i>New England Journal of Medicine</i> , 2010 , 362, 1901-8 | 59.2 | 242 |
| 260 | The activity of the epithelial sodium channel is regulated by clathrin-mediated endocytosis. <i>Journal of Biological Chemistry</i> , 1997 , 272, 25537-41 | 5.4 | 235 |
| 259 | Genome-wide association study of intracranial aneurysm identifies three new risk loci. <i>Nature Genetics</i> , 2010 , 42, 420-5 | 36.3 | 234 |
| 258 | ADCK4 mutations promote steroid-resistant nephrotic syndrome through CoQ10 biosynthesis disruption. <i>Journal of Clinical Investigation</i> , 2013 , 123, 5179-89 | 15.9 | 231 |
| 257 | Geographic differences in genetic susceptibility to IgA nephropathy: GWAS replication study and geospatial risk analysis. <i>PLoS Genetics</i> , 2012 , 8, e1002765 | 6 | 231 |
| 256 | Mutational landscape of MCPyV-positive and MCPyV-negative Merkel cell carcinomas with implications for immunotherapy. <i>Oncotarget</i> , 2016 , 7, 3403-15 | 3.3 | 229 |
| 255 | Characterization of the mutational landscape of anaplastic thyroid cancer via whole-exome sequencing. <i>Human Molecular Genetics</i> , 2015 , 24, 2318-29 | 5.6 | 225 |
| 254 | Mutations in SEC63 cause autosomal dominant polycystic liver disease. <i>Nature Genetics</i> , 2004 , 36, 575-7 | 36.3 | 224 |
| 253 | Gitelman's syndrome revisited: an evaluation of symptoms and health-related quality of life. <i>Kidney International</i> , 2001 , 59, 710-7 | 9.9 | 224 |
| 252 | Susceptibility loci for intracranial aneurysm in European and Japanese populations. <i>Nature Genetics</i> , 2008 , 40, 1472-7 | 36.3 | 222 |
| 251 | Landscape of somatic single-nucleotide and copy-number mutations in uterine serous carcinoma. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013 , 110, 2916-21 | 11.5 | 221 |
| 250 | A founder mutation as a cause of cerebral cavernous malformation in Hispanic Americans. <i>New England Journal of Medicine</i> , 1996 , 334, 946-51 | 59.2 | 219 |
| 249 | Skint1, the prototype of a newly identified immunoglobulin superfamily gene cluster, positively selects epidermal gammadelta T cells. <i>Nature Genetics</i> , 2008 , 40, 656-62 | 36.3 | 216 |
| 248 | Sites of regulated phosphorylation that control K-Cl cotransporter activity. <i>Cell</i> , 2009 , 138, 525-36 | 56.2 | 215 |

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|-----|--|------|-----|
| 247 | Hypertension with or without adrenal hyperplasia due to different inherited mutations in the potassium channel KCNJ5. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012 , 109, 2533-8 | 11.5 | 215 |
| 246 | A novel form of human mendelian hypertension featuring nonglucocorticoid-remediable aldosteronism. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2008 , 93, 3117-23 | 5.6 | 211 |
| 245 | Recurrent gain of function mutation in calcium channel CACNA1H causes early-onset hypertension with primary aldosteronism. <i>ELife</i> , 2015 , 4, e06315 | 8.9 | 203 |
| 244 | Angiotensin II signaling increases activity of the renal Na-Cl cotransporter through a WNK4-SPAK-dependent pathway. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009 , 106, 4384-9 | 11.5 | 200 |
| 243 | Nonvalidation of reported genetic risk factors for acute coronary syndrome in a large-scale replication study. <i>JAMA - Journal of the American Medical Association</i> , 2007 , 297, 1551-61 | 27.4 | 199 |
| 242 | Multilocus linkage of familial hyperkalaemia and hypertension, pseudohypoaldosteronism type II, to chromosomes 1q31-42 and 17p11-q21. <i>Nature Genetics</i> , 1997 , 16, 202-5 | 36.3 | 197 |
| 241 | Rare copy number variations in congenital heart disease patients identify unique genes in left-right patterning. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011 , 108, 2915-20 | 11.5 | 192 |
| 240 | A translocation causing increased alpha-klotho level results in hypophosphatemic rickets and hyperparathyroidism. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008 , 105, 3455-60 | 11.5 | 188 |
| 239 | Molecular physiology of the WNK kinases. <i>Annual Review of Physiology</i> , 2008 , 70, 329-55 | 23.1 | 184 |
| 238 | Aberrant IgA1 glycosylation is inherited in familial and sporadic IgA nephropathy. <i>Journal of the American Society of Nephrology: JASN</i> , 2008 , 19, 1008-14 | 12.7 | 183 |
| 237 | WNK3 modulates transport of Cl ⁻ in and out of cells: implications for control of cell volume and neuronal excitability. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005 , 102, 16783-8 | 11.5 | 178 |
| 236 | Regulation of NKCC2 by a chloride-sensing mechanism involving the WNK3 and SPAK kinases. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008 , 105, 8458-63 | 11.5 | 174 |
| 235 | Kelch-like 3 and Cullin 3 regulate electrolyte homeostasis via ubiquitination and degradation of WNK4. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013 , 110, 7838-43 | 11.5 | 167 |
| 234 | Recurrent activating mutation in PRKACA in cortisol-producing adrenal tumors. <i>Nature Genetics</i> , 2014 , 46, 613-7 | 36.3 | 165 |
| 233 | Copy-number disorders are a common cause of congenital kidney malformations. <i>American Journal of Human Genetics</i> , 2012 , 91, 987-97 | 11 | 161 |
| 232 | WNK3 kinase is a positive regulator of NKCC2 and NCC, renal cation-Cl ⁻ cotransporters required for normal blood pressure homeostasis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005 , 102, 16777-82 | 11.5 | 156 |
| 231 | WNK4 regulates apical and basolateral Cl ⁻ flux in extrarenal epithelia. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2004 , 101, 2064-9 | 11.5 | 151 |
| 230 | Intracranial aneurysm and hemorrhagic stroke in glucocorticoid-remediable aldosteronism. <i>Hypertension</i> , 1998 , 31, 445-50 | 8.5 | 144 |

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| 229 | Paracellular Cl ⁻ permeability is regulated by WNK4 kinase: insight into normal physiology and hypertension. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2004 , 101, 14877-82 | 11.5 | 143 |
| 228 | Mutations in the Na-Cl cotransporter reduce blood pressure in humans. <i>Hypertension</i> , 2001 , 37, 1458-64 | 8.5 | 142 |
| 227 | Exome Sequencing Identifies Biallelic MSH3 Germline Mutations as a Recessive Subtype of Colorectal Adenomatous Polyposis. <i>American Journal of Human Genetics</i> , 2016 , 99, 337-51 | 11 | 139 |
| 226 | Disruption of contactin 4 (CNTN4) results in developmental delay and other features of 3p deletion syndrome. <i>American Journal of Human Genetics</i> , 2004 , 74, 1286-93 | 11 | 135 |
| 225 | Phosphoregulation of the Na-K-2Cl and K-Cl cotransporters by the WNK kinases. <i>Biochimica Et Biophysica Acta - Molecular Basis of Disease</i> , 2010 , 1802, 1150-8 | 6.9 | 134 |
| 224 | KANK deficiency leads to podocyte dysfunction and nephrotic syndrome. <i>Journal of Clinical Investigation</i> , 2015 , 125, 2375-84 | 15.9 | 133 |
| 223 | An SGK1 site in WNK4 regulates Na ⁺ channel and K ⁺ channel activity and has implications for aldosterone signaling and K ⁺ homeostasis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007 , 104, 4025-9 | 11.5 | 131 |
| 222 | Comprehensive re-sequencing of adrenal aldosterone producing lesions reveal three somatic mutations near the KCNJ5 potassium channel selectivity filter. <i>PLoS ONE</i> , 2012 , 7, e41926 | 3.7 | 128 |
| 221 | Molecular and cellular reorganization of neural circuits in the human lineage. <i>Science</i> , 2017 , 358, 1027-1032 | 9.3 | 127 |
| 220 | Mutational landscape of uterine and ovarian carcinosarcomas implicates histone genes in epithelial-mesenchymal transition. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016 , 113, 12238-12243 | 11.5 | 123 |
| 219 | Mineralocorticoid receptor phosphorylation regulates ligand binding and renal response to volume depletion and hyperkalemia. <i>Cell Metabolism</i> , 2013 , 18, 660-71 | 24.6 | 122 |
| 218 | WNK1, a kinase mutated in inherited hypertension with hyperkalemia, localizes to diverse Cl ⁻ -transporting epithelia. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2003 , 100, 663-8 | 11.5 | 120 |
| 217 | CLCN2 chloride channel mutations in familial hyperaldosteronism type II. <i>Nature Genetics</i> , 2018 , 50, 349-354 | 36.4 | 117 |
| 216 | Two locus inheritance of non-syndromic midline craniosynostosis via rare and common alleles. <i>ELife</i> , 2016 , 5, | 8.9 | 115 |
| 215 | The B1-subunit of the H ⁺ ATPase is required for maximal urinary acidification. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005 , 102, 13616-21 | 11.5 | 113 |
| 214 | Mutations in sphingosine-1-phosphate lyase cause nephrosis with ichthyosis and adrenal insufficiency. <i>Journal of Clinical Investigation</i> , 2017 , 127, 912-928 | 15.9 | 112 |
| 213 | Identification of somatic mutations in parathyroid tumors using whole-exome sequencing. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2012 , 97, E1774-81 | 5.6 | 112 |
| 212 | Recessive loss of function of the neuronal ubiquitin hydrolase UCHL1 leads to early-onset progressive neurodegeneration. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013 , 110, 3489-94 | 11.5 | 111 |

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| 211 | Whole-exome sequencing characterizes the landscape of somatic mutations and copy number alterations in adrenocortical carcinoma. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2015 , 100, E493-502 | 5.6 | 110 |
| 210 | Genetic landscape of metastatic and recurrent head and neck squamous cell carcinoma. <i>Journal of Clinical Investigation</i> , 2016 , 126, 169-80 | 15.9 | 110 |
| 209 | Mutations in nuclear pore genes NUP93, NUP205 and XPO5 cause steroid-resistant nephrotic syndrome. <i>Nature Genetics</i> , 2016 , 48, 457-65 | 36.3 | 109 |
| 208 | Skint-1 is a highly specific, unique selecting component for epidermal T cells. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011 , 108, 3330-5 | 11.5 | 109 |
| 207 | Regression of Chemotherapy-Resistant Polymerase β (POLE) Ultra-Mutated and MSH6 Hyper-Mutated Endometrial Tumors with Nivolumab. <i>Clinical Cancer Research</i> , 2016 , 22, 5682-5687 | 12.9 | 109 |
| 206 | KCNJ10 determines the expression of the apical Na-Cl cotransporter (NCC) in the early distal convoluted tubule (DCT1). <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014 , 111, 11864-9 | 11.5 | 107 |
| 205 | WNK4 regulates activity of the epithelial Na ⁺ channel in vitro and in vivo. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007 , 104, 4020-4 | 11.5 | 106 |
| 204 | Mutations in KEOPS-complex genes cause nephrotic syndrome with primary microcephaly. <i>Nature Genetics</i> , 2017 , 49, 1529-1538 | 36.3 | 105 |
| 203 | KRIT1, a gene mutated in cerebral cavernous malformation, encodes a microtubule-associated protein. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2002 , 99, 10677-82 | 11.5 | 104 |
| 202 | Whole Exome Sequencing of Patients with Steroid-Resistant Nephrotic Syndrome. <i>Clinical Journal of the American Society of Nephrology: CJASN</i> , 2018 , 13, 53-62 | 6.9 | 103 |
| 201 | Frequency and phenotypic spectrum of germline mutations in POLE and seven other polymerase genes in 266 patients with colorectal adenomas and carcinomas. <i>International Journal of Cancer</i> , 2015 , 137, 320-31 | 7.5 | 102 |
| 200 | Autosomal dominant pseudohypoaldosteronism type 1: mechanisms, evidence for neonatal lethality, and phenotypic expression in adults. <i>Journal of the American Society of Nephrology: JASN</i> , 2006 , 17, 1429-36 | 12.7 | 101 |
| 199 | WNK3 bypasses the tonicity requirement for K-Cl cotransporter activation via a phosphatase-dependent pathway. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2006 , 103, 1976-81 | 11.5 | 100 |
| 198 | Insights into genetics, human biology and disease gleaned from family based genomic studies. <i>Genetics in Medicine</i> , 2019 , 21, 798-812 | 8.1 | 100 |
| 197 | WNK protein kinases modulate cellular Cl ⁻ flux by altering the phosphorylation state of the Na-K-Cl and K-Cl cotransporters. <i>Physiology</i> , 2006 , 21, 326-35 | 9.8 | 98 |
| 196 | Early and multiple origins of metastatic lineages within primary tumors. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016 , 113, 2140-5 | 11.5 | 95 |
| 195 | Mutations in DSTYK and dominant urinary tract malformations. <i>New England Journal of Medicine</i> , 2013 , 369, 621-9 | 59.2 | 95 |
| 194 | The Centers for Mendelian Genomics: a new large-scale initiative to identify the genes underlying rare Mendelian conditions. <i>American Journal of Medical Genetics, Part A</i> , 2012 , 158A, 1523-5 | 2.5 | 92 |

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|-----|--|------|----|
| 193 | Monogenic causes of chronic kidney disease in adults. <i>Kidney International</i> , 2019 , 95, 914-928 | 9.9 | 92 |
| 192 | Autoantibodies neutralizing type I IFNs are present in 4% of uninfected individuals over 70 years old and account for 20% of COVID-19 deaths. <i>Science Immunology</i> , 2021 , 6, | 28 | 91 |
| 191 | Novel somatic mutations in primary hyperaldosteronism are related to the clinical, radiological and pathological phenotype. <i>Clinical Endocrinology</i> , 2015 , 83, 779-89 | 3.4 | 88 |
| 190 | A form of the metabolic syndrome associated with mutations in DYRK1B. <i>New England Journal of Medicine</i> , 2014 , 370, 1909-1919 | 59.2 | 86 |
| 189 | Multilineage somatic activating mutations in HRAS and NRAS cause mosaic cutaneous and skeletal lesions, elevated FGF23 and hypophosphatemia. <i>Human Molecular Genetics</i> , 2014 , 23, 397-407 | 5.6 | 85 |
| 188 | Recessive LAMC3 mutations cause malformations of occipital cortical development. <i>Nature Genetics</i> , 2011 , 43, 590-4 | 36.3 | 85 |
| 187 | Molecular cloning and characterization of Atp6n1b: a novel fourth murine vacuolar H ⁺ -ATPase α -subunit gene. <i>Journal of Biological Chemistry</i> , 2001 , 276, 42382-8 | 5.4 | 84 |
| 186 | Genetic Drivers of Kidney Defects in the DiGeorge Syndrome. <i>New England Journal of Medicine</i> , 2017 , 376, 742-754 | 59.2 | 83 |
| 185 | Genomic characterization of sarcomatoid transformation in clear cell renal cell carcinoma. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016 , 113, 2170-5 | 11.5 | 83 |
| 184 | Common variant near the endothelin receptor type A (EDNRA) gene is associated with intracranial aneurysm risk. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011 , 108, 19707-12 | 11.5 | 82 |
| 183 | Localization of a gene for autosomal recessive distal renal tubular acidosis with normal hearing (rdRTA2) to 7q33-34. <i>American Journal of Human Genetics</i> , 1999 , 65, 1656-65 | 11 | 80 |
| 182 | Decreased ENaC expression compensates the increased NCC activity following inactivation of the kidney-specific isoform of WNK1 and prevents hypertension. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010 , 107, 18109-14 | 11.5 | 79 |
| 181 | Isolation of specific RNAs using DNA covalently linked to diazobenzoyloxymethyl cellulose or paper. <i>Methods in Enzymology</i> , 1979 , 68, 206-20 | 1.7 | 78 |
| 180 | Whole exome sequencing frequently detects a monogenic cause in early onset nephrolithiasis and nephrocalcinosis. <i>Kidney International</i> , 2018 , 93, 204-213 | 9.9 | 77 |
| 179 | Whole-Exome Sequencing Identifies Causative Mutations in Families with Congenital Anomalies of the Kidney and Urinary Tract. <i>Journal of the American Society of Nephrology: JASN</i> , 2018 , 29, 2348-2361 | 12.7 | 75 |
| 178 | Isolated polycystic liver disease genes define effectors of polycystin-1 function. <i>Journal of Clinical Investigation</i> , 2017 , 127, 1772-1785 | 15.9 | 74 |
| 177 | Finding genetic contributions to sporadic disease: a recessive locus at 12q24 commonly contributes to patent ductus arteriosus. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2002 , 99, 15054-9 | 11.5 | 73 |
| 176 | Evaluation of the dexamethasone suppression test for the diagnosis of glucocorticoid-remediable aldosteronism. <i>Journal of Clinical Endocrinology and Metabolism</i> , 1997 , 82, 3570-3 | 5.6 | 70 |

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|-----|---|------|----|
| 175 | FAT1 mutations cause a glomerulotubular nephropathy. <i>Nature Communications</i> , 2016 , 7, 10822 | 17.4 | 69 |
| 174 | The Psychiatric Cell Map Initiative: A Convergent Systems Biological Approach to Illuminating Key Molecular Pathways in Neuropsychiatric Disorders. <i>Cell</i> , 2018 , 174, 505-520 | 56.2 | 69 |
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