Mark R Cookson

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86 282 33,821 181 g-index h-index citations papers 8.7 7.12 39,175 323 L-index avg, IF ext. papers ext. citations

| # | Paper | IF | Citations |
|-----|---|-------------------|-----------|
| 282 | alpha-Synuclein locus triplication causes Parkinson's disease. <i>Science</i> , 2003 , 302, 841 | 33.3 | 3252 |
| 281 | Guidelines for the use and interpretation of assays for monitoring autophagy. Autophagy, 2012, 8, 445- | 5 44 .2 | 2783 |
| 280 | PINK1 is selectively stabilized on impaired mitochondria to activate Parkin. <i>PLoS Biology</i> , 2010 , 8, e1000 | 02 9/8 | 1887 |
| 279 | Genome-wide association study reveals genetic risk underlying Parkinson's disease. <i>Nature Genetics</i> , 2009 , 41, 1308-12 | 36.3 | 1469 |
| 278 | The Parkinson's disease protein DJ-1 is neuroprotective due to cysteine-sulfinic acid-driven mitochondrial localization. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2004 , 101, 9103-8 | 11.5 | 888 |
| 277 | Abundant quantitative trait loci exist for DNA methylation and gene expression in human brain. <i>PLoS Genetics</i> , 2010 , 6, e1000952 | 6 | 612 |
| 276 | Kinase activity is required for the toxic effects of mutant LRRK2/dardarin. <i>Neurobiology of Disease</i> , 2006 , 23, 329-41 | 7.5 | 608 |
| 275 | Common genetic variants influence human subcortical brain structures. <i>Nature</i> , 2015 , 520, 224-9 | 50.4 | 601 |
| 274 | Identification of novel risk loci, causal insights, and heritable risk for Parkinson's disease: a meta-analysis of genome-wide association studies. <i>Lancet Neurology, The</i> , 2019 , 18, 1091-1102 | 24.1 | 562 |
| 273 | The biochemistry of Parkinson's disease. <i>Annual Review of Biochemistry</i> , 2005 , 74, 29-52 | 29.1 | 536 |
| 272 | Parkin protects against the toxicity associated with mutant alpha-synuclein: proteasome dysfunction selectively affects catecholaminergic neurons. <i>Neuron</i> , 2002 , 36, 1007-19 | 13.9 | 506 |
| 271 | An in vitro model of Parkinson's disease: linking mitochondrial impairment to altered alpha-synuclein metabolism and oxidative damage. <i>Journal of Neuroscience</i> , 2002 , 22, 7006-15 | 6.6 | 490 |
| 270 | Genetic variability in the regulation of gene expression in ten regions of the human brain. <i>Nature Neuroscience</i> , 2014 , 17, 1418-1428 | 25.5 | 483 |
| 269 | Guidelines for the use and interpretation of assays for monitoring autophagy (4th edition). <i>Autophagy</i> , 2021 , 17, 1-382 | 10.2 | 440 |
| 268 | Alpha-synuclein implicated in Parkinson's disease is present in extracellular biological fluids, including human plasma. <i>FASEB Journal</i> , 2003 , 17, 1945-7 | 0.9 | 436 |
| 267 | The role of leucine-rich repeat kinase 2 (LRRK2) in Parkinson's disease. <i>Nature Reviews Neuroscience</i> , 2010 , 11, 791-7 | 13.5 | 400 |
| 266 | Identification of common variants influencing risk of the tauopathy progressive supranuclear palsy. Nature Genetics, 2011 , 43, 699-705 | 36.3 | 386 |

(2007-2010)

| 265 | Evidence for natural antisense transcript-mediated inhibition of microRNA function. <i>Genome Biology</i> , 2010 , 11, R56 | 18.3 | 375 |
|-----|--|-------|-----|
| 264 | A soluble Esynuclein construct forms a dynamic tetramer. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011 , 108, 17797-802 | 11.5 | 354 |
| 263 | DJ-1 acts in parallel to the PINK1/parkin pathway to control mitochondrial function and autophagy. <i>Human Molecular Genetics</i> , 2011 , 20, 40-50 | 5.6 | 346 |
| 262 | The expression of DJ-1 (PARK7) in normal human CNS and idiopathic Parkinson's disease. <i>Brain</i> , 2004 , 127, 420-30 | 11.2 | 341 |
| 261 | Mutations in the Matrin 3 gene cause familial amyotrophic lateral sclerosis. <i>Nature Neuroscience</i> , 2014 , 17, 664-666 | 25.5 | 319 |
| 260 | Distinct DNA methylation changes highly correlated with chronological age in the human brain. <i>Human Molecular Genetics</i> , 2011 , 20, 1164-72 | 5.6 | 312 |
| 259 | Mutations in PTEN-induced putative kinase 1 associated with recessive parkinsonism have differential effects on protein stability. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005 , 102, 5703-8 | 11.5 | 303 |
| 258 | Unbiased screen for interactors of leucine-rich repeat kinase 2 supports a common pathway for sporadic and familial Parkinson disease. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014 , 111, 2626-31 | 11.5 | 282 |
| 257 | Mitochondrial function and morphology are impaired in parkin-mutant fibroblasts. <i>Annals of Neurology</i> , 2008 , 64, 555-65 | 9.4 | 280 |
| 256 | Intersecting pathways to neurodegeneration in Parkinson's disease: effects of the pesticide rotenone on DJ-1, alpha-synuclein, and the ubiquitin-proteasome system. <i>Neurobiology of Disease</i> , 2006 , 22, 404-20 | 7.5 | 262 |
| 255 | alpha-Synuclein and neuronal cell death. <i>Molecular Neurodegeneration</i> , 2009 , 4, 9 | 19 | 257 |
| 254 | Genetics of Parkinson's disease and parkinsonism. <i>Annals of Neurology</i> , 2006 , 60, 389-98 | 9.4 | 252 |
| 253 | Genome-wide association study of obsessive-compulsive disorder. <i>Molecular Psychiatry</i> , 2013 , 18, 788-9 | 815.1 | 244 |
| 252 | Pink1 forms a multiprotein complex with Miro and Milton, linking Pink1 function to mitochondrial trafficking. <i>Biochemistry</i> , 2009 , 48, 2045-52 | 3.2 | 238 |
| 251 | The Parkinson disease-associated leucine-rich repeat kinase 2 (LRRK2) is a dimer that undergoes intramolecular autophosphorylation. <i>Journal of Biological Chemistry</i> , 2008 , 283, 16906-14 | 5.4 | 236 |
| 250 | Lysosomal impairment in Parkinson's disease. <i>Movement Disorders</i> , 2013 , 28, 725-32 | 7 | 228 |
| 249 | Deletion at ITPR1 underlies ataxia in mice and spinocerebellar ataxia 15 in humans. <i>PLoS Genetics</i> , 2007 , 3, e108 | 6 | 221 |
| 248 | The R1441C mutation of LRRK2 disrupts GTP hydrolysis. <i>Biochemical and Biophysical Research Communications</i> , 2007 , 357, 668-71 | 3.4 | 219 |

| 247 | Major Shifts in Glial Regional Identity Are a Transcriptional Hallmark of Human Brain Aging. <i>Cell Reports</i> , 2017 , 18, 557-570 | 10.6 | 217 |
|-----|---|------|-----|
| 246 | Leucine-rich repeat kinase 2 mutations and Parkinson's disease: three questions. <i>ASN Neuro</i> , 2009 , 1, | 5.3 | 216 |
| 245 | Formation of a stabilized cysteine sulfinic acid is critical for the mitochondrial function of the parkinsonism protein DJ-1. <i>Journal of Biological Chemistry</i> , 2009 , 284, 6476-85 | 5.4 | 210 |
| 244 | Excessive burden of lysosomal storage disorder gene variants in Parkinson's disease. <i>Brain</i> , 2017 , 140, 3191-3203 | 11.2 | 209 |
| 243 | A two-stage meta-analysis identifies several new loci for Parkinson's disease. <i>PLoS Genetics</i> , 2011 , 7, e1002142 | 6 | 209 |
| 242 | Cytoplasmic Pink1 activity protects neurons from dopaminergic neurotoxin MPTP. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008 , 105, 1716-21 | 11.5 | 206 |
| 241 | Mitochondrial dysfunction in a cell culture model of familial amyotrophic lateral sclerosis. <i>Brain</i> , 2002 , 125, 1522-33 | 11.2 | 205 |
| 240 | Structure of the ROC domain from the Parkinson's disease-associated leucine-rich repeat kinase 2 reveals a dimeric GTPase. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008 , 105, 1499-504 | 11.5 | 198 |
| 239 | LRRK2 modulates vulnerability to mitochondrial dysfunction in Caenorhabditis elegans. <i>Journal of Neuroscience</i> , 2009 , 29, 9210-8 | 6.6 | 196 |
| 238 | Converging pathways in neurodegeneration, from genetics to mechanisms. <i>Nature Neuroscience</i> , 2018 , 21, 1300-1309 | 25.5 | 183 |
| 237 | L166P mutant DJ-1, causative for recessive Parkinson's disease, is degraded through the ubiquitin-proteasome system. <i>Journal of Biological Chemistry</i> , 2003 , 278, 36588-95 | 5.4 | 181 |
| 236 | Novel genetic loci associated with hippocampal volume. <i>Nature Communications</i> , 2017 , 8, 13624 | 17.4 | 173 |
| 235 | RNA binding activity of the recessive parkinsonism protein DJ-1 supports involvement in multiple cellular pathways. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008 , 105, 10244-9 | 11.5 | 171 |
| 234 | DYT16, a novel young-onset dystonia-parkinsonism disorder: identification of a segregating mutation in the stress-response protein PRKRA. <i>Lancet Neurology, The</i> , 2008 , 7, 207-15 | 24.1 | 159 |
| 233 | The chaperone activity of heat shock protein 90 is critical for maintaining the stability of leucine-rich repeat kinase 2. <i>Journal of Neuroscience</i> , 2008 , 28, 3384-91 | 6.6 | 158 |
| 232 | Oxidative stress and motor neurone disease. <i>Brain Pathology</i> , 1999 , 9, 165-86 | 6 | 156 |
| 231 | Mitochondrial alterations in PINK1 deficient cells are influenced by calcineurin-dependent dephosphorylation of dynamin-related protein 1. <i>PLoS ONE</i> , 2009 , 4, e5701 | 3.7 | 150 |
| 230 | Novel genetic loci underlying human intracranial volume identified through genome-wide association. <i>Nature Neuroscience</i> , 2016 , 19, 1569-1582 | 25.5 | 147 |

(2008-2004)

| 229 | A strategy for designing inhibitors of alpha-synuclein aggregation and toxicity as a novel treatment for Parkinson's disease and related disorders. <i>FASEB Journal</i> , 2004 , 18, 1315-7 | 0.9 | 146 |
|-----|--|------|-----|
| 228 | LRRK2 phosphorylates membrane-bound Rabs and is activated by GTP-bound Rab7L1 to promote recruitment to the trans-Golgi network. <i>Human Molecular Genetics</i> , 2018 , 27, 385-395 | 5.6 | 143 |
| 227 | Cell systems and the toxic mechanism(s) of alpha-synuclein. Experimental Neurology, 2008, 209, 5-11 | 5.7 | 141 |
| 226 | Caspase-1 causes truncation and aggregation of the Parkinson's disease-associated protein Bynuclein. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016 , 113, 9587-92 | 11.5 | 139 |
| 225 | Parkinson's disease: insights from pathways. <i>Human Molecular Genetics</i> , 2010 , 19, R21-7 | 5.6 | 137 |
| 224 | MAPT expression and splicing is differentially regulated by brain region: relation to genotype and implication for tauopathies. <i>Human Molecular Genetics</i> , 2012 , 21, 4094-103 | 5.6 | 134 |
| 223 | Metabolic activity determines efficacy of macroautophagic clearance of pathological oligomeric alpha-synuclein. <i>American Journal of Pathology</i> , 2009 , 175, 736-47 | 5.8 | 127 |
| 222 | The Parkinson's disease kinase LRRK2 autophosphorylates its GTPase domain at multiple sites. <i>Biochemical and Biophysical Research Communications</i> , 2009 , 389, 449-54 | 3.4 | 125 |
| 221 | Evolutionary and functional relationships within the DJ1 superfamily. <i>BMC Evolutionary Biology</i> , 2004 , 4, 6 | 3 | 120 |
| 220 | DJ-1 regulation of mitochondrial function and autophagy through oxidative stress. <i>Autophagy</i> , 2011 , 7, 531-2 | 10.2 | 118 |
| 219 | Reelin and stk25 have opposing roles in neuronal polarization and dendritic Golgi deployment. <i>Cell</i> , 2010 , 143, 826-36 | 56.2 | 117 |
| 218 | Co-ordinate transcriptional regulation of dopamine synthesis genes by alpha-synuclein in human neuroblastoma cell lines. <i>Journal of Neurochemistry</i> , 2003 , 85, 957-68 | 6 | 116 |
| 217 | Mutant LRRK2 toxicity in neurons depends on LRRK2 levels and synuclein but not kinase activity or inclusion bodies. <i>Journal of Neuroscience</i> , 2014 , 34, 418-33 | 6.6 | 115 |
| 216 | Parkinsonism due to mutations in PINK1, parkin, and DJ-1 and oxidative stress and mitochondrial pathways. <i>Cold Spring Harbor Perspectives in Medicine</i> , 2012 , 2, a009415 | 5.4 | 111 |
| 215 | RING finger 1 mutations in Parkin produce altered localization of the protein. <i>Human Molecular Genetics</i> , 2003 , 12, 2957-65 | 5.6 | 111 |
| 214 | Genome-wide screen identifies rs646776 near sortilin as a regulator of progranulin levels in human plasma. <i>American Journal of Human Genetics</i> , 2010 , 87, 890-7 | 11 | 110 |
| 213 | Identification and functional characterization of a novel R621C mutation in the synphilin-1 gene in Parkinson's disease. <i>Human Molecular Genetics</i> , 2003 , 12, 1223-31 | 5.6 | 110 |
| 212 | Emerging pathways in genetic Parkinson's disease: Potential role of ceramide metabolism in Lewy body disease. <i>FEBS Journal</i> , 2008 , 275, 5767-73 | 5.7 | 109 |

| 211 | mRNA expression, splicing and editing in the embryonic and adult mouse cerebral cortex. <i>Nature Neuroscience</i> , 2013 , 16, 499-506 | 25.5 | 107 |
|-----|---|--------------------------------|-----|
| 210 | LRRK2 at the interface of autophagosomes, endosomes and lysosomes. <i>Molecular Neurodegeneration</i> , 2016 , 11, 73 | 19 | 107 |
| 209 | Membrane-associated farnesylated UCH-L1 promotes alpha-synuclein neurotoxicity and is a therapeutic target for Parkinson's disease. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009 , 106, 4635-40 | 11.5 | 102 |
| 208 | Cross-disorder genome-wide analyses suggest a complex genetic relationship between Tourette's syndrome and OCD. <i>American Journal of Psychiatry</i> , 2015 , 172, 82-93 | 11.9 | 101 |
| 207 | Insight into the mode of action of the LRRK2 Y1699C pathogenic mutant. <i>Journal of Neurochemistry</i> , 2011 , 116, 304-15 | 6 | 101 |
| 206 | Integration of GWAS SNPs and tissue specific expression profiling reveal discrete eQTLs for human traits in blood and brain. <i>Neurobiology of Disease</i> , 2012 , 47, 20-8 | 7.5 | 100 |
| 205 | A direct interaction between leucine-rich repeat kinase 2 and specific Eubulin isoforms regulates tubulin acetylation. <i>Journal of Biological Chemistry</i> , 2014 , 289, 895-908 | 5.4 | 98 |
| 204 | The expression of the glial glutamate transporter protein EAAT2 in motor neuron disease: an immunohistochemical study. <i>European Journal of Neuroscience</i> , 1998 , 10, 2481-9 | 3.5 | 97 |
| 203 | A pathway-based analysis provides additional support for an immune-related genetic susceptibility to Parkinson's disease. <i>Human Molecular Genetics</i> , 2013 , 22, 1039-49 | 5.6 | 96 |
| 202 | Mitochondria and dopamine: new insights into recessive parkinsonism. <i>Neuron</i> , 2004 , 43, 301-4 | 13.9 | 94 |
| 201 | Development and characterisation of a glutamate-sensitive motor neurone cell line. <i>Journal of Neurochemistry</i> , 2000 , 74, 1895-902 | 6 | 92 |
| 200 | Age-associated miRNA alterations in skeletal muscle from rhesus monkeys reversed by caloric restriction. <i>Aging</i> , 2013 , 5, 692-703 | 5.6 | 91 |
| 199 | Parkinson disease-associated mutation R1441H in LRRK2 prolongs the "active state" of its GTPase domain. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014 , 111, 405 | 5 ¹ 60 ⁵ | 88 |
| 198 | LRRK2 Pathways Leading to Neurodegeneration. <i>Current Neurology and Neuroscience Reports</i> , 2015 , 15, 42 | 6.6 | 87 |
| 197 | The G2385R variant of leucine-rich repeat kinase 2 associated with Parkinson's disease is a partial loss-of-function mutation. <i>Biochemical Journal</i> , 2012 , 446, 99-111 | 3.8 | 87 |
| 196 | Analysis of the cytosolic proteome in a cell culture model of familial amyotrophic lateral sclerosis reveals alterations to the proteasome, antioxidant defenses, and nitric oxide synthetic pathways. Journal of Biological Chemistry, 2003, 278, 6371-83 | 5.4 | 85 |
| 195 | Genes and parkinsonism. <i>Lancet Neurology, The</i> , 2003 , 2, 221-8 | 24.1 | 85 |
| 194 | Tyrosinase exacerbates dopamine toxicity but is not genetically associated with Parkinson's disease. <i>Journal of Neurochemistry</i> , 2005 , 93, 246-56 | 6 | 84 |

| 193 | LRRK2 promotes the activation of NLRC4 inflammasome during Typhimurium infection. <i>Journal of Experimental Medicine</i> , 2017 , 214, 3051-3066 | 16.6 | 83 | |
|-----|---|---------------|----|--|
| 192 | MKK6 binds and regulates expression of Parkinson's disease-related protein LRRK2. <i>Journal of Neurochemistry</i> , 2010 , 112, 1593-604 | 6 | 83 | |
| 191 | Pathways to Parkinsonism. <i>Neuron</i> , 2003 , 37, 7-10 | 13.9 | 83 | |
| 190 | LRRK2 links genetic and sporadic Parkinson's disease. <i>Biochemical Society Transactions</i> , 2019 , 47, 651-66 | 5 5 .1 | 82 | |
| 189 | LRRK2 kinase activity is dependent on LRRK2 GTP binding capacity but independent of LRRK2 GTP binding. <i>PLoS ONE</i> , 2011 , 6, e23207 | 3.7 | 80 | |
| 188 | Peroxynitrite and hydrogen peroxide induced cell death in the NSC34 neuroblastoma x spinal cord cell line: role of poly (ADP-ribose) polymerase. <i>Journal of Neurochemistry</i> , 1998 , 70, 501-8 | 6 | 78 | |
| 187 | Biochemical characterization of highly purified leucine-rich repeat kinases 1 and 2 demonstrates formation of homodimers. <i>PLoS ONE</i> , 2012 , 7, e43472 | 3.7 | 78 | |
| 186 | The Parkinson's disease associated LRRK2 exhibits weaker in vitro phosphorylation of 4E-BP compared to autophosphorylation. <i>PLoS ONE</i> , 2010 , 5, e8730 | 3.7 | 77 | |
| 185 | Phosphorylation of LRRK2 by casein kinase 1lregulates trans-Golgi clustering via differential interaction with ARHGEF7. <i>Nature Communications</i> , 2014 , 5, 5827 | 17.4 | 76 | |
| 184 | Identification of protein phosphatase 1 as a regulator of the LRRK2 phosphorylation cycle. <i>Biochemical Journal</i> , 2013 , 456, 119-28 | 3.8 | 76 | |
| 183 | Effects of DJ-1 mutations and polymorphisms on protein stability and subcellular localization. <i>Molecular Brain Research</i> , 2005 , 134, 76-83 | | 75 | |
| 182 | Mitochondrial quality control and dynamics in Parkinson's disease. <i>Antioxidants and Redox Signaling</i> , 2012 , 16, 869-82 | 8.4 | 74 | |
| 181 | Mutations in LRRK2/dardarin associated with Parkinson disease are more toxic than equivalent mutations in the homologous kinase LRRK1. <i>Journal of Neurochemistry</i> , 2007 , 102, 93-102 | 6 | 72 | |
| 180 | Finding useful biomarkers for Parkinson's disease. Science Translational Medicine, 2018, 10, | 17.5 | 69 | |
| 179 | Genetic modifiers of risk and age at onset in GBA associated Parkinson's disease and Lewy body dementia. <i>Brain</i> , 2020 , 143, 234-248 | 11.2 | 69 | |
| 178 | Hexokinase activity is required for recruitment of parkin to depolarized mitochondria. <i>Human Molecular Genetics</i> , 2014 , 23, 145-56 | 5.6 | 68 | |
| 177 | DJ-1, PINK1, and their effects on mitochondrial pathways. <i>Movement Disorders</i> , 2010 , 25 Suppl 1, S44-8 | 7 | 67 | |
| 176 | Glial phagocytic clearance in Parkinson's disease. <i>Molecular Neurodegeneration</i> , 2019 , 14, 16 | 19 | 66 | |

| 175 | FUS mutations in sporadic amyotrophic lateral sclerosis. <i>Neurobiology of Aging</i> , 2011 , 32, 550.e1-4 | 5.6 | 64 |
|-----|--|------------------|----|
| 174 | Amyotrophic lateral sclerosis 2-deficiency leads to neuronal degeneration in amyotrophic lateral sclerosis through altered AMPA receptor trafficking. <i>Journal of Neuroscience</i> , 2006 , 26, 11798-806 | 6.6 | 64 |
| 173 | Genetic neuropathology of Parkinson's disease. <i>International Journal of Clinical and Experimental Pathology</i> , 2008 , 1, 217-31 | 1.4 | 63 |
| 172 | Pathways to Parkinsonism Redux: convergent pathobiological mechanisms in genetics of Parkinson's disease. <i>Human Molecular Genetics</i> , 2015 , 24, R32-44 | 5.6 | 61 |
| 171 | Analysis of IFT74 as a candidate gene for chromosome 9p-linked ALS-FTD. <i>BMC Neurology</i> , 2006 , 6, 44 | 3.1 | 61 |
| 170 | Selective loss of neurofilament expression in Cu/Zn superoxide dismutase (SOD1) linked amyotrophic lateral sclerosis. <i>Journal of Neurochemistry</i> , 2002 , 82, 1118-28 | 6 | 60 |
| 169 | Aggregation of Esynuclein in brain samples from subjects with glucocerebrosidase mutations. <i>Molecular Genetics and Metabolism</i> , 2011 , 104, 185-8 | 3.7 | 57 |
| 168 | Post-transcriptional regulation of mRNA associated with DJ-1 in sporadic Parkinson disease. <i>Neuroscience Letters</i> , 2009 , 452, 8-11 | 3.3 | 57 |
| 167 | Mechanisms in dominant parkinsonism: The toxic triangle of LRRK2, alpha-synuclein, and tau. <i>BioEssays</i> , 2010 , 32, 227-235 | 4.1 | 57 |
| 166 | The Endosomal-Lysosomal Pathway Is Dysregulated by Expression. <i>Frontiers in Neuroscience</i> , 2017 , 11, 702 | 5.1 | 56 |
| 165 | How genetics research in Parkinson's disease is enhancing understanding of the common idiopathic forms of the disease. <i>Current Opinion in Neurology</i> , 2005 , 18, 706-11 | 7.1 | 56 |
| 164 | mTOR independent regulation of macroautophagy by Leucine Rich Repeat Kinase 2 via Beclin-1. <i>Scientific Reports</i> , 2016 , 6, 35106 | 4.9 | 54 |
| 163 | Transfected synphilin-1 forms cytoplasmic inclusions in HEK293 cells. <i>Molecular Brain Research</i> , 2001 , 97, 94-102 | | 53 |
| 162 | The role of monogenic genes in idiopathic Parkinson's disease. <i>Neurobiology of Disease</i> , 2019 , 124, 230- | 2 3.9 | 53 |
| 161 | Genes associated with Parkinson's disease: regulation of autophagy and beyond. <i>Journal of Neurochemistry</i> , 2016 , 139 Suppl 1, 91-107 | 6 | 50 |
| 160 | Frequency of Loss of Function Variants in LRRK2 in Parkinson Disease. <i>JAMA Neurology</i> , 2018 , 75, 1416- | 1422 | 50 |
| 159 | Is inhibition of kinase activity the only therapeutic strategy for LRRK2-associated Parkinson's disease?. <i>BMC Medicine</i> , 2012 , 10, 20 | 11.4 | 50 |
| 158 | Parkinson disease-associated mutations in LRRK2 cause centrosomal defects via Rab8a phosphorylation. <i>Molecular Neurodegeneration</i> , 2018 , 13, 3 | 19 | 49 |

| 157 | Genetic comorbidities in Parkinson's disease. Human Molecular Genetics, 2014, 23, 831-41 | 5.6 | 49 |
|-----|---|------|----|
| 156 | Glial cell inclusions and the pathogenesis of neurodegenerative diseases. <i>Neuron Glia Biology</i> , 2004 , 1, 13-21 | | 48 |
| 155 | LRRK2 mediates tubulation and vesicle sorting from lysosomes. Science Advances, 2020, 6, | 14.3 | 48 |
| 154 | Age-associated changes in gene expression in human brain and isolated neurons. <i>Neurobiology of Aging</i> , 2013 , 34, 1199-209 | 5.6 | 44 |
| 153 | Leucine-rich repeat kinase 2 interacts with p21-activated kinase 6 to control neurite complexity in mammalian brain. <i>Journal of Neurochemistry</i> , 2015 , 135, 1242-56 | 6 | 44 |
| 152 | Expression of PINK1 mRNA in human and rodent brain and in Parkinson's disease. <i>Brain Research</i> , 2007 , 1184, 10-6 | 3.7 | 44 |
| 151 | The metalloprotease inhibitor TIMP-3 regulates amyloid precursor protein and apolipoprotein E receptor proteolysis. <i>Journal of Neuroscience</i> , 2007 , 27, 10895-905 | 6.6 | 44 |
| 150 | The roles of kinases in familial Parkinson's disease. <i>Journal of Neuroscience</i> , 2007 , 27, 11865-8 | 6.6 | 43 |
| 149 | Cu/Zn superoxide dismutase (SOD1) mutations associated with familial amyotrophic lateral sclerosis (ALS) affect cellular free radical release in the presence of oxidative stress. <i>Amyotrophic Lateral Sclerosis and Other Motor Neuron Disorders: Official Publication of the World Federation of</i> | | 43 |
| 148 | Neurology, Research Group on Motor Neuron Diseases, 2002, 3, 75-85 Heterogeneity of leucine-rich repeat kinase 2 mutations: genetics, mechanisms and therapeutic implications. Neurotherapeutics, 2014, 11, 738-50 | 6.4 | 42 |
| 147 | The endocytic membrane trafficking pathway plays a major role in the risk of Parkinson's disease. <i>Movement Disorders</i> , 2019 , 34, 460-468 | 7 | 40 |
| 146 | Initial assessment of the pathogenic mechanisms of the recently identified Alzheimer risk Loci. <i>Annals of Human Genetics</i> , 2013 , 77, 85-105 | 2.2 | 40 |
| 145 | Parkin's substrates and the pathways leading to neuronal damage. <i>NeuroMolecular Medicine</i> , 2003 , 3, 1-13 | 4.6 | 40 |
| 144 | Variation in tau isoform expression in different brain regions and disease states. <i>Neurobiology of Aging</i> , 2013 , 34, 1922.e7-1922.e12 | 5.6 | 39 |
| 143 | Differential protein-protein interactions of LRRK1 and LRRK2 indicate roles in distinct cellular signaling pathways. <i>Journal of Neurochemistry</i> , 2014 , 131, 239-50 | 6 | 38 |
| 142 | Resolving the polymorphism-in-probe problem is critical for correct interpretation of expression QTL studies. <i>Nucleic Acids Research</i> , 2013 , 41, e88 | 20.1 | 38 |
| 141 | Alpha-synuclein overexpression increases dopamine toxicity in BE2-M17 cells. <i>BMC Neuroscience</i> , 2010 , 11, 41 | 3.2 | 38 |
| 140 | Detection of endogenous S1292 LRRK2 autophosphorylation in mouse tissue as a readout for kinase activity. <i>Npj Parkinsonm Disease</i> , 2018 , 4, 13 | 9.7 | 37 |

| 139 | Proteomic analysis reveals co-ordinated alterations in protein synthesis and degradation pathways in LRRK2 knockout mice. <i>Human Molecular Genetics</i> , 2018 , 27, 3257-3271 | 5.6 | 37 |
|-----|--|-----|----|
| 138 | The R1441C mutation alters the folding properties of the ROC domain of LRRK2. <i>Biochimica Et Biophysica Acta - Molecular Basis of Disease</i> , 2009 , 1792, 1194-7 | 6.9 | 37 |
| 137 | A comprehensive analysis of SNCA-related genetic risk in sporadic parkinson disease. <i>Annals of Neurology</i> , 2018 , 84, 117-129 | 9.4 | 33 |
| 136 | Arsenite stress down-regulates phosphorylation and 14-3-3 binding of leucine-rich repeat kinase 2 (LRRK2), promoting self-association and cellular redistribution. <i>Journal of Biological Chemistry</i> , 2014 , 289, 21386-400 | 5.4 | 33 |
| 135 | Parkin deficiency disrupts calcium homeostasis by modulating phospholipase C signalling. <i>FEBS Journal</i> , 2009 , 276, 5041-52 | 5.7 | 33 |
| 134 | The expression of the glutamate re-uptake transporter excitatory amino acid transporter 1 (EAAT1) in the normal human CNS and in motor neurone disease: an immunohistochemical study. <i>Neuroscience</i> , 2002 , 109, 27-44 | 3.9 | 32 |
| 133 | Cellular functions of LRRK2 implicate vesicular trafficking pathways in Parkinson's disease. <i>Biochemical Society Transactions</i> , 2016 , 44, 1603-1610 | 5.1 | 32 |
| 132 | AKT signalling selectively regulates PINK1 mitophagy in SHSY5Y cells and human iPSC-derived neurons. <i>Scientific Reports</i> , 2018 , 8, 8855 | 4.9 | 31 |
| 131 | Gene expression in the Parkinson's disease brain. Brain Research Bulletin, 2012, 88, 302-12 | 3.9 | 31 |
| 130 | Cell population-specific expression analysis of human cerebellum. <i>BMC Genomics</i> , 2012 , 13, 610 | 4.5 | 31 |
| 129 | Characterisation of a novel NR4A2 mutation in Parkinson's disease brain. <i>Neuroscience Letters</i> , 2009 , 457, 75-9 | 3.3 | 31 |
| 128 | Superoxide-induced nitric oxide release from cultured glial cells. <i>Brain Research</i> , 2001 , 911, 203-10 | 3.7 | 31 |
| 127 | Transcriptome analysis of LRRK2 knock-out microglia cells reveals alterations of inflammatory- and oxidative stress-related pathways upon treatment with Esynuclein fibrils. <i>Neurobiology of Disease</i> , 2019 , 129, 67-78 | 7.5 | 30 |
| 126 | RNA-binding proteins implicated in neurodegenerative diseases. <i>Wiley Interdisciplinary Reviews RNA</i> , 2017 , 8, e1397 | 9.3 | 30 |
| 125 | PAK6 Phosphorylates 14-3-3[to Regulate Steady State Phosphorylation of LRRK2. <i>Frontiers in Molecular Neuroscience</i> , 2017 , 10, 417 | 6.1 | 30 |
| 124 | Hexokinases link DJ-1 to the PINK1/parkin pathway. <i>Molecular Neurodegeneration</i> , 2017 , 12, 70 | 19 | 29 |
| 123 | Screening of AP endonuclease as a candidate gene for amyotrophic lateral sclerosis (ALS). <i>NeuroReport</i> , 2000 , 11, 1695-7 | 1.7 | 29 |
| 122 | The impact of fraudulent and irreproducible data to the translational research crisis - solutions and implementation. <i>Journal of Neurochemistry</i> , 2016 , 139 Suppl 2, 253-270 | 6 | 28 |

| 121 | The DYT6 Dystonia Protein THAP1 Regulates Myelination within the Oligodendrocyte Lineage. <i>Developmental Cell</i> , 2017 , 42, 52-67.e4 | 10.2 | 28 | |
|-----|---|-------------------|----|--|
| 120 | The role of PTEN-induced kinase 1 in mitochondrial dysfunction and dynamics. <i>International Journal of Biochemistry and Cell Biology</i> , 2009 , 41, 2025-35 | 5.6 | 28 | |
| 119 | The role of Rab GTPases in the pathobiology of Parkinson' disease. <i>Current Opinion in Cell Biology</i> , 2019 , 59, 73-80 | 9 | 27 | |
| 118 | Penetrance of Parkinson's Disease in LRRK2 p.G2019S Carriers Is Modified by a Polygenic Risk Score. <i>Movement Disorders</i> , 2020 , 35, 774-780 | 7 | 27 | |
| 117 | Pathogenic LRRK2 mutations do not alter gene expression in cell model systems or human brain tissue. <i>PLoS ONE</i> , 2011 , 6, e22489 | 3.7 | 27 | |
| 116 | Differential gene expression in a cell culture model of SOD1-related familial motor neurone disease. <i>Human Molecular Genetics</i> , 2002 , 11, 2061-75 | 5.6 | 27 | |
| 115 | Mechanisms of Mutant LRRK2 Neurodegeneration. Advances in Neurobiology, 2017, 14, 227-239 | 2.1 | 26 | |
| 114 | Poly(ADP-ribose) polymerase is found in both the nucleus and cytoplasm of human CNS neurons. Brain Research, 1999 , 834, 182-5 | 3.7 | 26 | |
| 113 | Kinesin-associated protein 3 (KIFAP3) has no effect on survival in a population-based cohort of ALS patients. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010 , 107, 12. | 3 35-8 | 25 | |
| 112 | Identification and characterization of the human parkin gene promoter. <i>Journal of Neurochemistry</i> , 2001 , 78, 1146-52 | 6 | 25 | |
| 111 | Molecular mechanism of olesoxime-mediated neuroprotection through targeting Esynuclein interaction with mitochondrial VDAC. <i>Cellular and Molecular Life Sciences</i> , 2020 , 77, 3611-3626 | 10.3 | 25 | |
| 110 | The G2385R risk factor for Parkinson's disease enhances CHIP-dependent intracellular degradation of LRRK2. <i>Biochemical Journal</i> , 2017 , 474, 1547-1558 | 3.8 | 24 | |
| 109 | The Parkinson's Disease Protein LRRK2 Interacts with the GARP Complex to Promote Retrograde Transport to the trans-Golgi Network. <i>Cell Reports</i> , 2020 , 31, 107614 | 10.6 | 24 | |
| 108 | DNAJC proteins and pathways to parkinsonism. FEBS Journal, 2019, 286, 3080-3094 | 5.7 | 23 | |
| 107 | Comparative Protein Interaction Network Analysis Identifies Shared and Distinct Functions for the Human ROCO Proteins. <i>Proteomics</i> , 2018 , 18, e1700444 | 4.8 | 23 | |
| 106 | Genetic determinants of survival in progressive supranuclear palsy: a genome-wide association study. <i>Lancet Neurology, The</i> , 2021 , 20, 107-116 | 24.1 | 23 | |
| 105 | Leucine-rich repeat kinase 2 controls protein kinase A activation state through phosphodiesterase 4. <i>Journal of Neuroinflammation</i> , 2018 , 15, 297 | 10.1 | 23 | |
| 104 | Transcriptomic profiling of the human brain reveals that altered synaptic gene expression is associated with chronological aging. <i>Scientific Reports</i> , 2017 , 7, 16890 | 4.9 | 22 | |

| 103 | Role of LRRK2 kinase dysfunction in Parkinson disease. <i>Expert Reviews in Molecular Medicine</i> , 2011 , 13, e20 | 6.7 | 22 |
|-----|--|------|----|
| 102 | Nicotinic receptors in the putamen of patients with dementia with Lewy bodies and Parkinson's disease: relation to changes in alpha-synuclein expression. <i>Neuroscience Letters</i> , 2002 , 335, 134-8 | 3.3 | 22 |
| 101 | Downregulation of Pink1 influences mitochondrial fusion-fission machinery and sensitizes to neurotoxins in dopaminergic cells. <i>NeuroToxicology</i> , 2014 , 44, 140-8 | 4.4 | 21 |
| 100 | Mutant torsinA interacts with tyrosine hydroxylase in cultured cells. <i>Neuroscience</i> , 2009 , 164, 1127-37 | 3.9 | 21 |
| 99 | LRRK2 mediates microglial neurotoxicity via NFATc2 in rodent models of synucleinopathies. <i>Science Translational Medicine</i> , 2020 , 12, | 17.5 | 21 |
| 98 | Analysis of macroautophagy related proteins in G2019S LRRK2 Parkinson's disease brains with Lewy body pathology. <i>Brain Research</i> , 2018 , 1701, 75-84 | 3.7 | 20 |
| 97 | Animal models for drug discovery in dystonia. Expert Opinion on Drug Discovery, 2008, 3, 83-97 | 6.2 | 20 |
| 96 | Normal localization of deltaF323-Y328 mutant torsinA in transfected human cells. <i>Neuroscience Letters</i> , 2002 , 327, 75-8 | 3.3 | 20 |
| 95 | Large-scale pathway specific polygenic risk and transcriptomic community network analysis identifies novel functional pathways in Parkinson disease. <i>Acta Neuropathologica</i> , 2020 , 140, 341-358 | 14.3 | 19 |
| 94 | In Vitro CRISPR/Cas9-Directed Gene Editing to Model LRRK2 G2019S Parkinson's Disease in Common Marmosets. <i>Scientific Reports</i> , 2020 , 10, 3447 | 4.9 | 19 |
| 93 | Reduction of PINK1 or DJ-1 impair mitochondrial motility in neurites and alter ER-mitochondria contacts. <i>Journal of Cellular and Molecular Medicine</i> , 2018 , 22, 5439-5449 | 5.6 | 19 |
| 92 | 14-3-3 proteins are promising LRRK2 interactors. <i>Biochemical Journal</i> , 2010 , 430, e5-6 | 3.8 | 19 |
| 91 | The Genetic Architecture of Parkinson Disease in Spain: Characterizing Population-Specific Risk, Differential Haplotype Structures, and Providing Etiologic Insight. <i>Movement Disorders</i> , 2019 , 34, 1851- | 1863 | 18 |
| 90 | The LRRK2 signalling system. <i>Cell and Tissue Research</i> , 2018 , 373, 39-50 | 4.2 | 18 |
| 89 | The function of orthologues of the human Parkinson's disease gene LRRK2 across species: implications for disease modelling in preclinical research. <i>Biochemical Journal</i> , 2016 , 473, 221-32 | 3.8 | 18 |
| 88 | Hero versus antihero: the multiple roles of alpha-synuclein in neurodegeneration. <i>Experimental Neurology</i> , 2006 , 199, 238-42 | 5.7 | 18 |
| 87 | Genetic analysis of amyotrophic lateral sclerosis identifies contributing pathways and cell types. <i>Science Advances</i> , 2021 , 7, | 14.3 | 18 |
| 86 | Preclinical modeling of chronic inhibition of the Parkinson's disease associated kinase LRRK2 reveals altered function of the endolysosomal system in vivo. <i>Molecular Neurodegeneration</i> , 2021 , 16, 17 | 19 | 17 |

(2020-2021)

| 85 | Identification of Candidate Parkinson Disease Genes by Integrating Genome-Wide Association Study, Expression, and Epigenetic Data Sets. <i>JAMA Neurology</i> , 2021 , 78, 464-472 | 17.2 | 17 |
|----|---|------|----|
| 84 | Cellular effects of LRRK2 mutations. <i>Biochemical Society Transactions</i> , 2012 , 40, 1070-3 | 5.1 | 16 |
| 83 | Neurodegeneration: how does parkin prevent Parkinson's disease?. Current Biology, 2003, 13, R522-4 | 6.3 | 16 |
| 82 | Selective loss of neurofilament proteins after exposure of differentiated human IMR-32 neuroblastoma cells to oxidative stress. <i>Brain Research</i> , 1996 , 738, 162-6 | 3.7 | 16 |
| 81 | Parkinson's disease-associated mutations in the GTPase domain of LRRK2 impair its nucleotide-dependent conformational dynamics. <i>Journal of Biological Chemistry</i> , 2019 , 294, 5907-5913 | 5.4 | 15 |
| 8o | Evolution of neurodegeneration. <i>Current Biology</i> , 2012 , 22, R753-61 | 6.3 | 15 |
| 79 | Molecular changes in the absence of severe pathology in the pulvinar in dementia with Lewy bodies. <i>Movement Disorders</i> , 2018 , 33, 982-991 | 7 | 14 |
| 78 | Unaltered alpha-synuclein blood levels in juvenile Parkinsonism with a parkin exon 4 deletion. <i>Neuroscience Letters</i> , 2005 , 374, 189-91 | 3.3 | 14 |
| 77 | Parkin and alpha-synuclein: opponent actions in the pathogenesis of Parkinson's disease. <i>Neuroscientist</i> , 2004 , 10, 63-72 | 7.6 | 14 |
| 76 | Comprehensive promoter level expression quantitative trait loci analysis of the human frontal lobe. <i>Genome Medicine</i> , 2016 , 8, 65 | 14.4 | 14 |
| 75 | Genetic analysis of neurodegenerative diseases in a pathology cohort. <i>Neurobiology of Aging</i> , 2019 , 76, 214.e1-214.e9 | 5.6 | 14 |
| 74 | AgingRNA in development and disease. Wiley Interdisciplinary Reviews RNA, 2012, 3, 133-43 | 9.3 | 13 |
| 73 | The Effects of Variants in the Parkin, PINK1, and DJ-1 Genes along with Evidence for their Pathogenicity. <i>Current Protein and Peptide Science</i> , 2017 , 18, 702-714 | 2.8 | 13 |
| 72 | Sequential screening nominates the Parkinson's disease associated kinase LRRK2 as a regulator of Clathrin-mediated endocytosis. <i>Neurobiology of Disease</i> , 2020 , 141, 104948 | 7.5 | 13 |
| 71 | Extracellular clusterin limits the uptake of Esynuclein fibrils by murine and human astrocytes. <i>Glia</i> , 2021 , 69, 681-696 | 9 | 13 |
| 70 | LAG3 is not expressed in human and murine neurons and does not modulate | 12 | 13 |
| 69 | mTOR independent alteration in ULK1 Ser758 phosphorylation following chronic LRRK2 kinase inhibition. <i>Bioscience Reports</i> , 2018 , 38, | 4.1 | 12 |
| 68 | APOE4 is Associated with Differential Regional Vulnerability to Bioenergetic Deficits in Aged APOE Mice. <i>Scientific Reports</i> , 2020 , 10, 4277 | 4.9 | 11 |

| 67 | Use of cysteine-reactive cross-linkers to probe conformational flexibility of human DJ-1 demonstrates that Glu18 mutations are dimers. <i>Journal of Neurochemistry</i> , 2014 , 130, 839-53 | 6 | 11 |
|----|---|------|----|
| 66 | Post-translational decrease in respiratory chain proteins in the Polg mutator mouse brain. <i>PLoS ONE</i> , 2014 , 9, e94646 | 3.7 | 11 |
| 65 | Dystonia and the nuclear envelope. <i>Neuron</i> , 2005 , 48, 875-7 | 13.9 | 11 |
| 64 | Glutamine synthetase activity and expression are not affected by the development of motor neuronopathy in the G93A SOD-1/ALS mouse. <i>Molecular Brain Research</i> , 2001 , 94, 131-6 | | 11 |
| 63 | The Polg Mutator Phenotype Does Not Cause Dopaminergic Neurodegeneration in DJ-1-Deficient Mice. <i>ENeuro</i> , 2015 , 2, | 3.9 | 11 |
| 62 | Divergent Effects of G2019S and R1441C LRRK2 Mutations on LRRK2 and Rab10 Phosphorylations in Mouse Tissues. <i>Cells</i> , 2020 , 9, | 7.9 | 11 |
| 61 | Assessment of APOE in atypical parkinsonism syndromes. <i>Neurobiology of Disease</i> , 2019 , 127, 142-146 | 7.5 | 10 |
| 60 | Identification of Stk25 as a genetic modifier of Tau phosphorylation in Dab1-mutant mice. <i>PLoS ONE</i> , 2012 , 7, e31152 | 3.7 | 10 |
| 59 | A feedforward loop links Gaucher and Parkinson's diseases?. <i>Cell</i> , 2011 , 146, 9-11 | 56.2 | 10 |
| 58 | Development, characterisation and epitope mapping of novel monoclonal antibodies for DJ-1 (PARK7) protein. <i>Neuroscience Letters</i> , 2005 , 383, 225-30 | 3.3 | 10 |
| 57 | Microarray analysis reveals induction of heat shock proteins mRNAs by the torsion dystonia protein, TorsinA. <i>Neuroscience Letters</i> , 2003 , 343, 5-8 | 3.3 | 10 |
| 56 | Biochemical characterization of torsinB. <i>Molecular Brain Research</i> , 2004 , 127, 1-9 | | 10 |
| 55 | High-frequency head impact causes chronic synaptic adaptation and long-term cognitive impairment in mice. <i>Nature Communications</i> , 2021 , 12, 2613 | 17.4 | 10 |
| 54 | Alpha-synuclein triggers T-cell response. Is Parkinson's disease an autoimmune disorder?. <i>Movement Disorders</i> , 2017 , 32, 1327 | 7 | 9 |
| 53 | Deep sequencing of coding and non-coding RNA in the CNS. Brain Research, 2010, 1338, 146-54 | 3.7 | 9 |
| 52 | Sequence conservation between mouse and human synphilin-1. <i>Neuroscience Letters</i> , 2002 , 322, 9-12 | 3.3 | 9 |
| 51 | Genomewide Association Studies of LRRK2 Modifiers of Parkinson's Disease. <i>Annals of Neurology</i> , 2021 , 90, 76-88 | 9.4 | 9 |
| 50 | Crystallizing ideas about Parkinson's disease. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2003 , 100, 9111-3 | 11.5 | 8 |

| 49 | Tackling neurodegenerative diseases with genomic engineering: A new stem cell initiative from the NIH. <i>Neuron</i> , 2021 , 109, 1080-1083 | 13.9 | 8 |
|----|---|------|---|
| 48 | LRRK2 recruitment, activity, and function in organelles. FEBS Journal, 2021, | 5.7 | 8 |
| 47 | Mutations in LRRK2 linked to Parkinson disease sequester Rab8a to damaged lysosomes and regulate transferrin-mediated iron uptake in microglia <i>PLoS Biology</i> , 2021 , 19, e3001480 | 9.7 | 8 |
| 46 | Parkin-mediated ubiquitination regulates phospholipase C-gamma1. <i>Journal of Cellular and Molecular Medicine</i> , 2009 , 13, 3061-8 | 5.6 | 7 |
| 45 | Cultured glial cells are resistant to the effects of motor neurone disease-associated SOD1 mutations. <i>Neuroscience Letters</i> , 2001 , 302, 146-50 | 3.3 | 7 |
| 44 | Unravelling the role of defective genes. <i>Progress in Brain Research</i> , 2010 , 183, 43-57 | 2.9 | 6 |
| 43 | LRRK2 mediates tubulation and vesicle sorting from membrane damaged lysosomes | | 6 |
| 42 | Investigation of Autosomal Genetic Sex Differences in Parkinson's Disease. <i>Annals of Neurology</i> , 2021 , 90, 35-42 | 9.4 | 6 |
| 41 | Pathogenic mutations in LRRK2 sequester Rab8a to damaged lysosomes and regulate transferrin-mediated iron uptake in microglia | | 5 |
| 40 | Association of Variants in the SPTLC1 Gene With Juvenile Amyotrophic Lateral Sclerosis. <i>JAMA Neurology</i> , 2021 , 78, 1236-1248 | 17.2 | 5 |
| 39 | ADAR2 affects mRNA coding sequence edits with only modest effects on gene expression or splicing in vivo. <i>RNA Biology</i> , 2016 , 13, 15-24 | 4.8 | 4 |
| 38 | Age-modulated association between prefrontal NAA and the BDNF gene. <i>International Journal of Neuropsychopharmacology</i> , 2013 , 16, 1185-93 | 5.8 | 4 |
| 37 | A revised 1.6 Istructure of the GTPase domain of the Parkinson disease-associated protein LRRK2 provides insights into mechanisms | | 4 |
| 36 | Combined Knockout of Lrrk2 and Rab29 Does Not Result in Behavioral Abnormalities in vivo. Journal of Parkinsonm Disease, 2021 , 11, 569-584 | 5.3 | 4 |
| 35 | Transcriptomic changes in brain development. <i>International Review of Neurobiology</i> , 2014 , 116, 233-50 | 4.4 | 3 |
| 34 | Identification of the epitope of a monoclonal antibody to DJ-1. Neuroscience Letters, 2005, 374, 203-6 | 3.3 | 3 |
| 33 | Molecules that cause or prevent Parkinson's disease. <i>PLoS Biology</i> , 2004 , 2, e401 | 9.7 | 3 |
| 32 | Preclinical Modeling of Chronic Inhibition of the Parkinson Disease Associated Kinase LRRK2 Reveals Altered Function of the Endolysosomal System in Vivo | | 3 |

| 31 | Mutations in Auxilin cause parkinsonism via impaired clathrin-mediated trafficking at the Golgi apparatus and synapse | | 3 |
|----|--|-----------------------|---|
| 30 | Identification of sixteen novel candidate genes for late onset Parkinson's disease. <i>Molecular Neurodegeneration</i> , 2021 , 16, 35 | 19 | 3 |
| 29 | Pathways of protein synthesis and degradation in PD pathogenesis. <i>Progress in Brain Research</i> , 2020 , 252, 217-270 | 2.9 | 3 |
| 28 | CoExp: A Web Tool for the Exploitation of Co-expression Networks. <i>Frontiers in Genetics</i> , 2021 , 12, 6301 | 8 4 7 5 | 3 |
| 27 | Coding and Noncoding Variation in LRRK2 and Parkinson's Disease Risk. Movement Disorders, 2021, | 7 | 3 |
| 26 | Proteomics; applications in familial Parkinson's disease. <i>Journal of Neurochemistry</i> , 2019 , 151, 446-458 | 6 | 2 |
| 25 | Structure, function, and leucine-rich repeat kinase 2: On the importance of reproducibility in understanding Parkinson's disease. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016 , 113, 8346-8 | 11.5 | 2 |
| 24 | The persistence of memory. New England Journal of Medicine, 2006, 355, 2697-8 | 59.2 | 2 |
| 23 | RNA sequencing of whole blood reveals early alterations in immune cells and gene expression in Parkinson disease. <i>Nature Aging</i> , 2021 , 1, 734-747 | | 2 |
| 22 | Transcriptome analysis of collagen VI-related muscular dystrophy muscle biopsies. <i>Annals of Clinical and Translational Neurology</i> , 2021 , 8, 2184-2198 | 5.3 | 2 |
| 21 | Combined knockout of Lrrk2 and Rab29 does not result in behavioral abnormalities in vivo | | 2 |
| 20 | Regulation of mitophagy by the NSL complex underlies genetic risk for Parkinson disease at Chr16q11.2 and on the MAPT H1 allele | | 2 |
| 19 | Evidence for connecting multiple neurodegenerative diseases. <i>Brain Communications</i> , 2021 , 3, fcab095 | 4.5 | 2 |
| 18 | Generation of fourteen isogenic cell lines for Parkinson's disease-associated leucine-rich repeat kinase (LRRK2). <i>Stem Cell Research</i> , 2021 , 53, 102354 | 1.6 | 2 |
| 17 | Quality Control Metrics for Whole Blood Transcriptome Analysis in the Parkinson Progression Markers Initiative (PPMI) | | 2 |
| 16 | Proteomics: techniques and applications in neuroscience. <i>Journal of Neurochemistry</i> , 2019 , 151, 394-396 | 66 | 1 |
| 15 | The commercial genetic testing landscape for Parkinson's disease. <i>Parkinsonism and Related Disorders</i> , 2021 , 92, 107-111 | 3.6 | 1 |
| 14 | The Role of LRRK2 Kinase Activity in Cellular PD Models 2008 , 423-431 | | 1 |

LIST OF PUBLICATIONS

| 13 | Rab activation | | 1 | |
|----|---|-----|---|--|
| 12 | Differences in Stability, Activity and Mutation Effects Between Human and Mouse Leucine-Rich Repeat Kinase 2. <i>Neurochemical Research</i> , 2019 , 44, 1446-1459 | 4.6 | 1 | |
| 11 | Directing LRRK2 to membranes of the endolysosomal pathway triggers RAB phosphorylation and JIP4 recruitment <i>Neurobiology of Disease</i> , 2022 , 105769 | 7.5 | 1 | |
| 10 | Generation of iPSC line from a Parkinson patient with PARK7 mutation and CRISPR-edited Gibco human episomal iPSC line to mimic PARK7 mutation. <i>Stem Cell Research</i> , 2021 , 55, 102506 | 1.6 | Ο | |
| 9 | Dominant torsinA mutations in cellular systems. Advances in Neurology, 2004, 94, 73-8 | | 0 | |
| 8 | Evaluation of Current Methods to Detect Cellular Leucine-Rich Repeat Kinase 2 (LRRK2) Kinase Activity. <i>Journal of Parkinson</i> Disease, 2022 , 1-25 | 5.3 | O | |
| 7 | Signatures of disrupted synaptic maintenance in the entorhinal cortex of both pathology-free APOE4 carriers and aged APOE4 mice. <i>Alzheimermand Dementia</i> , 2020 , 16, e046192 | 1.2 | | |
| 6 | Parkinson∃ Disease and Other Synucleinopathies 2018 , 117-143 | | | |
| 5 | Gene Linkage and Systems Biology. <i>Advances in Neurobiology</i> , 2017 , 15, 479-489 | 2.1 | | |
| 4 | Conference Scene: Parkinson disease in the UK: a report from the Parkinson UK Research Conference. <i>Neurodegenerative Disease Management</i> , 2011 , 1, 25-27 | 2.8 | | |
| 3 | Parkinson Disease and Related Disorders 2006 , 199-212 | | | |
| 2 | Transcriptional signatures in iPSC-derived neurons are reproducible across labs when differentiation protocols are closely matched. <i>Stem Cell Research</i> , 2021 , 56, 102558 | 1.6 | | |
| 1 | Assays for Pten-Induced Novel Kinase 1 (PINK1) and Leucine-Rich Repeat Kinase 2 (LRRK2), Kinases Associated with Parkinson Disease. <i>Neuromethods</i> , 2012 , 219-236 | 0.4 | | |