

Toshiyuki Yamamoto

List of Publications by Year in descending order

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papers

2,623
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docs citations

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times ranked

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| # | ARTICLE | IF | CITATIONS |
|----|--|-----|-----------|
| 1 | A Japanese patient with a 2p25.3 terminal deletion presented with early-onset obesity, intellectual disability and diabetes mellitus: A case report. <i>Journal of Diabetes Investigation</i> , 2022, 13, 391-396. | 1.1 | 1 |
| 2 | Clinical and genetic diagnosis of thirteen Japanese patients with hereditary spherocytosis. <i>Human Genome Variation</i> , 2022, 9, 1. | 0.4 | 10 |
| 3 | Congenital anemia reveals distinct targeting mechanisms for master transcription factor GATA1. <i>Blood</i> , 2022, 139, 2534-2546. | 0.6 | 14 |
| 4 | Induced pluripotent stem cells established from a female patient with Xq22 deletion confirm that BEX2 escapes from X-chromosome inactivation. <i>Congenital Anomalies (discontinued)</i> , 2021, 61, 63-67. | 0.3 | 2 |
| 5 | MCT8 deficiency in a patient with a novel frameshift variant in the SLC16A2 gene. <i>Human Genome Variation</i> , 2021, 8, 10. | 0.4 | 0 |
| 6 | Deep intronic deletion in intron 3 of PLP1 is associated with a severe phenotype of Pelizaeus-Merzbacher disease. <i>Human Genome Variation</i> , 2021, 8, 14. | 0.4 | 4 |
| 7 | A recurrent de novo ZSWIM6 variant in a Japanese patient with severe neurodevelopmental delay and frequent vomiting. <i>Human Genome Variation</i> , 2021, 8, 16. | 0.4 | 2 |
| 8 | Clinical spectrum of individuals with de novo <i>EBF3</i> variants or deletions. <i>American Journal of Medical Genetics, Part A</i> , 2021, 185, 2913-2921. | 0.7 | 7 |
| 9 | <i>HECW2</i> -related disorder in four Japanese patients. <i>American Journal of Medical Genetics, Part A</i> , 2021, 185, 2895-2902. | 0.7 | 6 |
| 10 | Genomic Aberrations Associated with the Pathophysiological Mechanisms of Neurodevelopmental Disorders. <i>Cells</i> , 2021, 10, 2317. | 1.8 | 6 |
| 11 | Recurrent de novo pathogenic variant of WASF1 in a Japanese patient with neurodevelopmental disorder with absent language and variable seizures. <i>Human Genome Variation</i> , 2021, 8, 43. | 0.4 | 4 |
| 12 | Establishment of a simple and rapid method to detect MECP2 duplications using digital polymerase chain reaction. <i>Congenital Anomalies (discontinued)</i> , 2020, 60, 10-14. | 0.3 | 4 |
| 13 | Primrose syndrome associated with unclassified immunodeficiency and a novel <i>ZBTB20</i> mutation. <i>American Journal of Medical Genetics, Part A</i> , 2020, 182, 521-526. | 0.7 | 5 |
| 14 | Coffin-Siris syndrome with bilateral macular dysplasia caused by a novel exonic deletion in ARID1B. <i>Congenital Anomalies (discontinued)</i> , 2020, 60, 189-193. | 0.3 | 2 |
| 15 | Novel LAMA2 variants identified in a patient with white matter abnormalities. <i>Human Genome Variation</i> , 2020, 7, 16. | 0.4 | 1 |
| 16 | De novo SMARCA2 variants clustered outside the helicase domain cause a new recognizable syndrome with intellectual disability and blepharophimosis distinct from Nicolaides-Baraitser syndrome. <i>Genetics in Medicine</i> , 2020, 22, 1838-1850. | 1.1 | 31 |
| 17 | Application of induced pluripotent stem cells in epilepsy. <i>Molecular and Cellular Neurosciences</i> , 2020, 108, 103535. | 1.0 | 13 |
| 18 | Molecular Profiles of Breast Cancer in a Single Institution. <i>Anticancer Research</i> , 2020, 40, 4567-4570. | 0.5 | 2 |

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|----|--|-----|-----------|
| 19 | Breakpoint junction analysis for complex genomic rearrangements with the caldera volcano-like pattern. <i>Human Mutation</i> , 2020, 41, 2119-2127. | 1.1 | 2 |
| 20 | Novel COL4A1 mutations identified in infants with congenital hemolytic anemia in association with brain malformations. <i>Human Genome Variation</i> , 2020, 7, 42. | 0.4 | 2 |
| 21 | Analyses of breakpoint junctions of complex genomic rearrangements comprising multiple consecutive microdeletions by nanopore sequencing. <i>Journal of Human Genetics</i> , 2020, 65, 735-741. | 1.1 | 8 |
| 22 | Complex chromosomal rearrangements of human chromosome 21 in a patient manifesting clinical features partially overlapped with that of Down syndrome. <i>Human Genetics</i> , 2020, 139, 1555-1563. | 1.8 | 5 |
| 23 | The involvement of U-type dicentric chromosomes in the formation of terminal deletions with or without adjacent inverted duplications. <i>Human Genetics</i> , 2020, 139, 1417-1427. | 1.8 | 12 |
| 24 | Gitelman syndrome caused by a novel hemiallelic missense mutation in SLC12A3 revealed by 16q12.2q21 microdeletion. <i>Human Genome Variation</i> , 2020, 7, 17. | 0.4 | 0 |
| 25 | Long-term natural history of an adult patient with distal 22q11.2 deletion from low copy repeat to E. <i>Congenital Anomalies (discontinued)</i> , 2019, 59, 102-103. | 0.3 | 0 |
| 26 | Advantages of ddPCR in detection of PLP1 duplications. <i>Intractable and Rare Diseases Research</i> , 2019, 8, 198-202. | 0.3 | 6 |
| 27 | PRRT2 mutations in Japanese patients with benign infantile epilepsy and paroxysmal kinesigenic dyskinesia. <i>Seizure: the Journal of the British Epilepsy Association</i> , 2019, 71, 1-5. | 0.9 | 16 |
| 28 | Genomic backgrounds of Japanese patients with undiagnosed neurodevelopmental disorders. <i>Brain and Development</i> , 2019, 41, 776-782. | 0.6 | 36 |
| 29 | Elucidation of the pathogenic mechanism and potential treatment strategy for a female patient with spastic paraplegia derived from a single-nucleotide deletion in PLP1. <i>Journal of Human Genetics</i> , 2019, 64, 665-671. | 1.1 | 9 |
| 30 | PRRT2 Mutation in a Sporadic Case of Paroxysmal Kinesigenic Dyskinesia. <i>Juntendo Medical Journal</i> , 2019, 65, 95-98. | 0.1 | 0 |
| 31 | Narrowing down the region responsible for 1q23.3q24.1 microdeletion by identifying the smallest deletion. <i>Human Genome Variation</i> , 2019, 6, 47. | 0.4 | 1 |
| 32 | Three Japanese patients with 3p13 microdeletions involving FOXP1. <i>Brain and Development</i> , 2019, 41, 257-262. | 0.6 | 4 |
| 33 | Phenotypic features of 1q41q42 microdeletion including WDR26 and FBXO28 are clinically recognizable: The first case from Japan. <i>Brain and Development</i> , 2019, 41, 452-455. | 0.6 | 3 |
| 34 | Natural histories of patients with Wolf-Hirschhorn syndrome derived from variable chromosomal abnormalities. <i>Congenital Anomalies (discontinued)</i> , 2019, 59, 169-173. | 0.3 | 6 |
| 35 | De novo 1p35.2 microdeletion including PUM1 identified in a patient with sporadic West syndrome. <i>Congenital Anomalies (discontinued)</i> , 2019, 59, 193-194. | 0.3 | 2 |
| 36 | Infantile spasms related to a 5q31.2-q31.3 microdeletion including PURA. <i>Human Genome Variation</i> , 2018, 5, 18007. | 0.4 | 6 |

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|----|---|-----|-----------|
| 37 | Somatic mosaic deletions involving <i>SCN1A</i> cause Dravet syndrome. <i>American Journal of Medical Genetics, Part A</i> , 2018, 176, 657-662. | 0.7 | 17 |
| 38 | A 10q21.3q22.2 microdeletion identified in a patient with severe developmental delay and multiple congenital anomalies including congenital heart defects. <i>Congenital Anomalies (discontinued)</i> , 2018, 58, 36-38. | 0.3 | 3 |
| 39 | Novel A178P mutation in <i>SLC16A2</i> in a patient with Allan-Herndon-Dudley syndrome. <i>Congenital Anomalies (discontinued)</i> , 2018, 58, 143-144. | 0.3 | 3 |
| 40 | Identification of a rare homozygous <i>SZT2</i> variant due to uniparental disomy in a patient with a neurodevelopmental disorder. <i>Intractable and Rare Diseases Research</i> , 2018, 7, 245-250. | 0.3 | 11 |
| 41 | Early-Onset Diabetes Mellitus in a Patient With a Chromosome 13q34qter Microdeletion Including IRS2. <i>Journal of the Endocrine Society</i> , 2018, 2, 1207-1213. | 0.1 | 5 |
| 42 | Novel compound heterozygous <i>EPC5</i> mutations consisted with a missense mutation and a microduplication in the exon 1 region identified in a Japanese patient with Vici syndrome. <i>American Journal of Medical Genetics, Part A</i> , 2018, 176, 2803-2807. | 0.7 | 3 |
| 43 | Interstitial deletion within 7q31.1q31.3 in a woman with mild intellectual disability and schizophrenia. <i>Neuropsychiatric Disease and Treatment</i> , 2018, Volume 14, 1773-1778. | 1.0 | 6 |
| 44 | A novel MLH1 mutation in a Japanese family with Lynch syndrome associated with small bowel cancer. <i>Human Genome Variation</i> , 2018, 5, 13. | 0.4 | 12 |
| 45 | Independent occurrence of de novo HSPD1 and HIP1 variants in brothers with different neurological disorders – leukodystrophy and autism. <i>Human Genome Variation</i> , 2018, 5, 18. | 0.4 | 6 |
| 46 | An episode of acute encephalopathy with biphasic seizures and late reduced diffusion followed by hemiplegia and intractable epilepsy observed in a patient with a novel frameshift mutation in HNRNPU. <i>Brain and Development</i> , 2018, 40, 813-818. | 0.6 | 13 |
| 47 | A novel PLP1 mutation F240L identified in a patient with congenital type Pelizaeus-Merzbacher disease. <i>Human Genome Variation</i> , 2017, 4, 16044. | 0.4 | 3 |
| 48 | Possible genes responsible for developmental delay observed in patients with rare 2q23q24 microdeletion syndrome: Literature review and description of an additional patient. <i>Congenital Anomalies (discontinued)</i> , 2017, 57, 109-113. | 0.3 | 5 |
| 49 | A 7q31.33q32.1 microdeletion including LRRC4 and GRM8 is associated with severe intellectual disability and characteristics of autism. <i>Human Genome Variation</i> , 2017, 4, 17001. | 0.4 | 19 |
| 50 | <i>ARHGEF9</i> disease. <i>Neurology: Genetics</i> , 2017, 3, e148. | 0.9 | 35 |
| 51 | Aspartylglucosaminuria caused by a novel homozygous mutation in the AGA gene was identified by an exome-first approach in a patient from Japan. <i>Brain and Development</i> , 2017, 39, 422-425. | 0.6 | 3 |
| 52 | An Xq22.1q22.2 nullisomy in a male patient with severe neurological impairment. <i>American Journal of Medical Genetics, Part A</i> , 2017, 173, 1124-1127. | 0.7 | 5 |
| 53 | A novel COL1A1 mutation in a family with osteogenesis imperfecta associated with phenotypic variabilities. <i>Human Genome Variation</i> , 2017, 4, 17007. | 0.4 | 3 |
| 54 | MED13L haploinsufficiency syndrome: A de novo frameshift and recurrent intragenic deletions due to parental mosaicism. , 2017, 173, 1264-1269. | | 23 |

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|----|--|-----|-----------|
| 55 | Familial 9q33q34 microduplication in siblings with developmental disorders and macrocephaly. <i>European Journal of Medical Genetics</i> , 2017, 60, 650-654. | 0.7 | 3 |
| 56 | Characteristics of rare and private deletions identified in phenotypically normal individuals. <i>Human Genome Variation</i> , 2017, 4, 17037. | 0.4 | 1 |
| 57 | A novel TUBB4A mutation G96R identified in a patient with hypomyelinating leukodystrophy onset beyond adolescence. <i>Human Genome Variation</i> , 2017, 4, 17035. | 0.4 | 15 |
| 58 | Mutations in NSD1 and NFIX in Three Patients with Clinical Features of Sotos Syndrome and Malan Syndrome. <i>Journal of Pediatric Genetics</i> , 2017, 06, 234-237. | 0.3 | 5 |
| 59 | A novel DARS2 mutation in a Japanese patient with leukoencephalopathy with brainstem and spinal cord involvement but no lactate elevation. <i>Human Genome Variation</i> , 2017, 4, 17051. | 0.4 | 6 |
| 60 | Abdominal paraganglioma in a young woman with 1p36 deletion syndrome. <i>American Journal of Medical Genetics, Part A</i> , 2017, 173, 495-500. | 0.7 | 2 |
| 61 | A case of Dravet syndrome with cortical myoclonus indicated by jerk-locked back-averaging of electroencephalogram data. <i>Brain and Development</i> , 2017, 39, 75-79. | 0.6 | 5 |
| 62 | The smallest de novo 20q11.2 microdeletion causing intellectual disability and dysmorphic features. <i>Human Genome Variation</i> , 2017, 4, 17050. | 0.4 | 3 |
| 63 | A novel <i>CASK</i> mutation identified in siblings exhibiting developmental disorders with/without microcephaly. <i>Intractable and Rare Diseases Research</i> , 2017, 6, 177-182. | 0.3 | 18 |
| 64 | A novel <i>PGK1</i> mutation associated with neurological dysfunction and the absence of episodes of hemolytic anemia or myoglobinuria. <i>Intractable and Rare Diseases Research</i> , 2017, 6, 132-136. | 0.3 | 14 |
| 65 | A 15q14 microdeletion involving MEIS2 identified in a patient with autism spectrum disorder. <i>Human Genome Variation</i> , 2017, 4, 17029. | 0.4 | 11 |
| 66 | Novel <i>SLC16A2</i> mutations in patients with Allan-Herndon-Dudley syndrome. <i>Intractable and Rare Diseases Research</i> , 2016, 5, 214-217. | 0.3 | 14 |
| 67 | A 16q12.2q21 deletion identified in a patient with developmental delay, epilepsy, short stature, and distinctive features. <i>Congenital Anomalies (discontinued)</i> , 2016, 56, 253-255. | 0.3 | 5 |
| 68 | Tattonâ€“Brownâ€“Rahman syndrome due to 2p23 microdeletion. <i>American Journal of Medical Genetics, Part A</i> , 2016, 170, 1339-1342. | 0.7 | 23 |
| 69 | A novel <i>TUBB3</i> mutation in a sporadic patient with asymmetric cortical dysplasia. <i>American Journal of Medical Genetics, Part A</i> , 2016, 170, 1076-1079. | 0.7 | 19 |
| 70 | Pancreatic developmental defect evaluated by celiac artery angiography in a patient with MODY5. <i>Human Genome Variation</i> , 2016, 3, 16022. | 0.4 | 6 |
| 71 | Challenges in detecting genomic copy number aberrations using next-generation sequencing data and the eXome Hidden Markov Model: a clinical exome-first diagnostic approach. <i>Human Genome Variation</i> , 2016, 3, 16025. | 0.4 | 38 |
| 72 | A 12p13 GRIN2B deletion is associated with developmental delay and macrocephaly. <i>Human Genome Variation</i> , 2016, 3, 16029. | 0.4 | 12 |

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|----|--|-----|-----------|
| 73 | Loss-of-function mutations and global rearrangements in GPC3 in patients with Simpsonâ€“Golabiâ€“Behmel syndrome. <i>Human Genome Variation</i> , 2016, 3, 16033. | 0.4 | 7 |
| 74 | 7p22.1 microdeletions involving ACTB associated with developmental delay, short stature, and microcephaly. <i>European Journal of Medical Genetics</i> , 2016, 59, 502-506. | 0.7 | 10 |
| 75 | Concurrent occurrence of an inherited 16p13.11 microduplication and a de novo 19p13.3 microdeletion involving MAP2K2 in a patient with developmental delay, distinctive facial features, and lambdoid synostosis. <i>European Journal of Medical Genetics</i> , 2016, 59, 559-563. | 0.7 | 5 |
| 76 | A de novo microdeletion in a patient with inner ear abnormalities suggests that the 10q26.13 region contains the responsible gene. <i>Human Genome Variation</i> , 2016, 3, 16008. | 0.4 | 8 |
| 77 | Use of targeted nextâ€“generation sequencing for molecular diagnosis of craniosynostosis: Identification of a novel <i>de novo</i> mutation of <i>EFNB1</i> . <i>Congenital Anomalies (discontinued)</i> , 2016, 56, 91-93. | 0.3 | 9 |
| 78 | Loss-of-function mutations of STXBP1 in patients with epileptic encephalopathy. <i>Brain and Development</i> , 2016, 38, 280-284. | 0.6 | 15 |
| 79 | Focal seizures and epileptic spasms in a child with Down syndrome from a family with a PRRT2 mutation. <i>Brain and Development</i> , 2016, 38, 597-600. | 0.6 | 7 |
| 80 | White matter abnormalities in an adult patient with l-2-hydroxyglutaric aciduria. <i>Brain and Development</i> , 2016, 38, 142-144. | 0.6 | 6 |
| 81 | The Novel Missense Mutation of GATA1 Caused Red Cell Adenosine Deaminase Overproduction Associated with Congenital Hemolytic Anemia. <i>Blood</i> , 2016, 128, 400-400. | 0.6 | 1 |
| 82 | Characteristics of 2p15â€“p16.1 microdeletion syndrome: Review and description of two additional patients. <i>Congenital Anomalies (discontinued)</i> , 2015, 55, 125-132. | 0.3 | 13 |
| 83 | Recurrent occurrences of CDKL5 mutations in patients with epileptic encephalopathy. <i>Human Genome Variation</i> , 2015, 2, 15042. | 0.4 | 8 |
| 84 | A novel MED12 mutation associated with non-specific X-linked intellectual disability. <i>Human Genome Variation</i> , 2015, 2, 15018. | 0.4 | 18 |
| 85 | Novel PLA2G6 mutations associated with an exonic deletion due to non-allelic homologous recombination in a patient with infantile neuroaxonal dystrophy. <i>Human Genome Variation</i> , 2015, 2, 15048. | 0.4 | 8 |
| 86 | An association of 19p13.2 microdeletions with Malan syndrome and Chiari malformation. <i>American Journal of Medical Genetics, Part A</i> , 2015, 167, 724-730. | 0.7 | 16 |
| 87 | Single Nucleotide Variations in CLCN6 Identified in Patients with Benign Partial Epilepsies in Infancy and/or Febrile Seizures. <i>PLoS ONE</i> , 2015, 10, e0118946. | 1.1 | 13 |
| 88 | A de novo microdeletion involving PAFAH1B (LIS1) related to lissencephaly phenotype. <i>Data in Brief</i> , 2015, 4, 488-491. | 0.5 | 5 |
| 89 | Microarray analysis of 50 patients reveals the critical chromosomal regions responsible for 1p36 deletion syndrome-related complications. <i>Brain and Development</i> , 2015, 37, 515-526. | 0.6 | 43 |
| 90 | Mutations in the genes encoding eukaryotic translation initiation factor 2B in Japanese patients with vanishing white matter disease. <i>Brain and Development</i> , 2015, 37, 960-966. | 0.6 | 10 |

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|-----|---|-----|-----------|
| 91 | Megalencephalic leukoencephalopathy with subcortical cysts caused by compound heterozygous mutations in MLC1 , in patients with and without subcortical cysts in the brain. Journal of the Neurological Sciences, 2015, 351, 211-213. | 0.3 | 7 |
| 92 | Epilepsy and Other Symptoms Associated with Chromosome 9q34.11 Microdeletion. Journal of Pediatric Epilepsy, 2015, 04, 023-029. | 0.1 | 2 |
| 93 | Growth profiles of 34 patients with Wolf-Hirschhorn syndrome. Journal of Pediatric Genetics, 2015, 01, 033-037. | 0.3 | 3 |
| 94 | Epilepsy in 1p36 Deletion Syndrome Is Not Associated with Deletion Size. Journal of Pediatric Epilepsy, 2015, 04, 004-007. | 0.1 | 0 |
| 95 | Periventricular heterotopia and white matter abnormalities in a girl with mosaic ring chromosome 6. Molecular Cytogenetics, 2015, 8, 54. | 0.4 | 8 |
| 96 | Comment on "Delayed myelination is not a constant feature of Allan-Herndon-Dudley syndrome: Report of a new case and review of the literature" by Azzolini S et al. Brain & Development 2014;36:716-720. Brain and Development, 2015, 37, 988-989. | 0.6 | 3 |
| 97 | Leukoencephalopathy associated with 11q24 deletion involving the gene encoding hepatic and glial cell adhesion molecule in two patients. European Journal of Medical Genetics, 2015, 58, 492-496. | 0.7 | 18 |
| 98 | Characteristics of patients with benign partial epilepsy in infancy without PRRT2 mutations. Epilepsy Research, 2015, 118, 10-13. | 0.8 | 8 |
| 99 | CHCHD2 is down-regulated in neuronal cells differentiated from iPS cells derived from patients with lissencephaly. Genomics, 2015, 106, 196-203. | 1.3 | 12 |
| 100 | A de novo TUBB4A mutation in a patient with hypomyelination mimicking Pelizaeus-Merzbacher disease. Brain and Development, 2015, 37, 281-285. | 0.6 | 13 |
| 101 | An emerging phenotype of Xq22 microdeletions in females with severe intellectual disability, hypotonia and behavioral abnormalities. Journal of Human Genetics, 2014, 59, 300-306. | 1.1 | 29 |
| 102 | Mild developmental delay and obesity in two patients with mosaic 1p36 deletion syndrome. American Journal of Medical Genetics, Part A, 2014, 164, 415-420. | 0.7 | 5 |
| 103 | Narrowing of the responsible region for severe developmental delay and autistic behaviors in WAGR syndrome down to 1.6 Mb including <i>PAX6</i> , <i>WT1</i> , and <i>PRRG4</i> . American Journal of Medical Genetics, Part A, 2014, 164, 634-638. | 0.7 | 31 |
| 104 | Neuropsychological profiles of patients with 2q37.3 deletion associated with developmental dyspraxia. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2014, 165, 684-690. | 1.1 | 10 |
| 105 | Clinical Course and Images of Four Familial Cases of Allan-Herndon-Dudley Syndrome With a Novel Monocarboxylate Transporter 8 Gene Mutation. Pediatric Neurology, 2014, 51, 414-416. | 1.0 | 13 |
| 106 | Growth patterns of patients with 1p36 deletion syndrome. Congenital Anomalies (discontinued), 2014, 54, 82-86. | 0.3 | 4 |
| 107 | Whole-exome sequencing identifies a de novo TUBA1A mutation in a patient with sporadic malformations of cortical development: a case report. BMC Research Notes, 2014, 7, 465. | 0.6 | 19 |
| 108 | Overlapping microdeletions involving 15q22.2 narrow the critical region for intellectual disability to NARG2 and RORA. European Journal of Medical Genetics, 2014, 57, 163-168. | 0.7 | 11 |

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|-----|--|-----|-----------|
| 109 | Novel compound heterozygous mutations of POLR3A revealed by whole-exome sequencing in a patient with hypomyelination. <i>Brain and Development</i> , 2014, 36, 315-321. | 0.6 | 20 |
| 110 | Pontine Malformation, Undecussated Pyramidal Tracts, and Regional Polymicrogyria: A New Syndrome. <i>Pediatric Neurology</i> , 2014, 50, 384-388. | 1.0 | 6 |
| 111 | MLC1 mutations in Japanese patients with megalencephalic leukoencephalopathy with subcortical cysts. <i>Human Genome Variation</i> , 2014, 1, 14019. | 0.4 | 8 |
| 112 | A novel KCNT1 mutation in a Japanese patient with epilepsy of infancy with migrating focal seizures. <i>Human Genome Variation</i> , 2014, 1, 14027. | 0.4 | 15 |
| 113 | SLC16A2 mutations in two Japanese patients with Allan-Herndon-Dudley syndrome. <i>Human Genome Variation</i> , 2014, 1, 14010. | 0.4 | 6 |
| 114 | Clinical impacts of genomic copy number gains at Xq28. <i>Human Genome Variation</i> , 2014, 1, 14001. | 0.4 | 30 |
| 115 | De novo triplication of 11q12.3 in a patient with developmental delay and distinctive facial features. <i>Molecular Cytogenetics</i> , 2013, 6, 15. | 0.4 | 5 |
| 116 | MECP2 duplication syndrome in both genders. <i>Brain and Development</i> , 2013, 35, 411-419. | 0.6 | 48 |
| 117 | Interstitial Duplication of 2q32.1-q33.3 in a Patient With Epilepsy, Developmental Delay, and Autistic Behavior. <i>American Journal of Medical Genetics, Part A</i> , 2013, 161, 1078-1084. | 0.7 | 26 |
| 118 | Clinical manifestations of Xq28 functional disomy involving <i>MECP2</i> in one female and two male patients. <i>American Journal of Medical Genetics, Part A</i> , 2013, 161, 1779-1785. | 0.7 | 21 |
| 119 | Microdeletions of 5.5 Mb (4q13.2-q13.3) and 4.1 Mb (7p15.3-p21.1) associated with a saethre-like phenotype, severe intellectual disability, and autism. <i>American Journal of Medical Genetics, Part A</i> , 2013, 161, 2078-2083. | 0.7 | 12 |
| 120 | Lissencephaly with marked ventricular dilation, agenesis of corpus callosum, and cerebellar hypoplasia caused by TUBA1A mutation. <i>Brain and Development</i> , 2013, 35, 274-279. | 0.6 | 34 |
| 121 | A novel homozygous mutation of GJC2 derived from maternal uniparental disomy in a female patient with Pelizaeus-Merzbacher-like disease. <i>Journal of the Neurological Sciences</i> , 2013, 330, 123-126. | 0.3 | 12 |
| 122 | Microdeletions of 3p21.31 characterized by developmental delay, distinctive features, elevated serum creatine kinase levels, and white matter involvement. <i>American Journal of Medical Genetics, Part A</i> , 2013, 161, 3049-3056. | 0.7 | 12 |
| 123 | PRRT2 mutation in Japanese children with benign infantile epilepsy. <i>Brain and Development</i> , 2013, 35, 641-646. | 0.6 | 31 |
| 124 | A recurrent KCNT1 mutation in two sporadic cases with malignant migrating partial seizures in infancy. <i>Gene</i> , 2013, 531, 467-471. | 1.0 | 80 |
| 125 | A cryptic microdeletion including <i>MBD5</i> occurring within the breakpoint of a reciprocal translocation between chromosomes 2 and 5 in a patient with developmental delay and obesity. <i>American Journal of Medical Genetics, Part A</i> , 2013, 161, 850-855. | 0.7 | 14 |
| 126 | 109 kb deletion of chromosome 4p16.3 in a patient with mild phenotype of Wolf-Hirschhorn syndrome. <i>American Journal of Medical Genetics, Part A</i> , 2013, 161, 1465-1469. | 0.7 | 19 |

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|-----|---|-----|-----------|
| 127 | <scp>P</scp>elizaesusâ€“<scp>M</scp>erzbacher disease as a chromosomal disorder. Congenital Anomalies (discontinued), 2013, 53, 3-8. | 0.3 | 22 |
| 128 | Challenges in genetic counseling because of intraâ€“familial phenotypic variation of oralâ€“facialâ€“digital syndrome type 1. Congenital Anomalies (discontinued), 2013, 53, 155-159. | 0.3 | 3 |
| 129 | Wholeâ€“exome sequencing of a unique brain malformation with periventricular heterotopia, cingulate polymicrogyria and midbrain tectal hyperplasia. Neuropathology, 2013, 33, 553-560. | 0.7 | 7 |
| 130 | Novel nucleotide mutation leading to a recurrent amino acid alteration in <i>SH3BP2</i> in a patient with cherubism. Congenital Anomalies (discontinued), 2013, 53, 166-169. | 0.3 | 6 |
| 131 | Pelizaeus-Merzbacher disease caused by a duplication-inverted triplication-duplication in chromosomal segments including the PLP1 region. European Journal of Medical Genetics, 2012, 55, 400-403. | 0.7 | 28 |
| 132 | De novo microdeletion of 5q14.3 excluding <i>MEF2C</i> in a patient with infantile spasms, microcephaly, and agenesis of the corpus callosum. American Journal of Medical Genetics, Part A, 2012, 158A, 2272-2276. | 0.7 | 22 |
| 133 | Subtelomeric deletions of 1q43q44 and severe brain impairment associated with delayed myelination. Journal of Human Genetics, 2012, 57, 593-600. | 1.1 | 20 |
| 134 | Reduced PLP1 expression in induced pluripotent stem cells derived from a Pelizaeusâ€“Merzbacher disease patient with a partial PLP1 duplication. Journal of Human Genetics, 2012, 57, 580-586. | 1.1 | 19 |
| 135 | An unmasked mutation of <i>EIF2B2</i> due to submicroscopic deletion of 14q24.3 in a patient with vanishing white matter disease. American Journal of Medical Genetics, Part A, 2012, 158A, 1771-1777. | 0.7 | 11 |
| 136 | Jacobsen syndrome due to an unbalanced translocation between 11q23 and 22q11.2 identified at age 40 years. American Journal of Medical Genetics, Part A, 2012, 158A, 220-223. | 0.7 | 5 |
| 137 | Tandem configurations of variably duplicated segments of 22q11.2 confirmed by fiber-FISH analysis. Journal of Human Genetics, 2011, 56, 810-812. | 1.1 | 9 |
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