

Heiko Martin Reutter

List of Publications by Year in descending order

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Version: 2024-02-01

157
papers

4,832
citations

126907

33
h-index

118850

62
g-index

157
all docs

157
docs citations

157
times ranked

5105
citing authors

#	ARTICLE	IF	CITATIONS
1	Key susceptibility locus for nonsyndromic cleft lip with or without cleft palate on chromosome 8q24. <i>Nature Genetics</i> , 2009, 41, 473-477.	21.4	415
2	Genome-wide association study identifies two susceptibility loci for nonsyndromic cleft lip with or without cleft palate. <i>Nature Genetics</i> , 2010, 42, 24-26.	21.4	379
3	Genome-wide meta-analyses of nonsyndromic cleft lip with or without cleft palate identify six new risk loci. <i>Nature Genetics</i> , 2012, 44, 968-971.	21.4	311
4	Mutations in 12 known dominant disease-causing genes clarify many congenital anomalies of the kidney and urinary tract. <i>Kidney International</i> , 2014, 85, 1429-1433.	5.2	203
5	The Exstrophy-epispadias complex. <i>Orphanet Journal of Rare Diseases</i> , 2009, 4, 23.	2.7	198
6	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.	6.1	147
7	Yunis-Varã³n Syndrome Is Caused by Mutations in FIG4, Encoding a Phosphoinositide Phosphatase. <i>American Journal of Human Genetics</i> , 2013, 92, 781-791.	6.2	124
8	Whole-exome resequencing reveals recessive mutations in TRAP1 in individuals with CAKUT and VACTERL association. <i>Kidney International</i> , 2014, 85, 1310-1317.	5.2	106
9	Loss of Function of GALNT2 Lowers High-Density Lipoproteins in Humans, Nonhuman Primates, and Rodents. <i>Cell Metabolism</i> , 2016, 24, 234-245.	16.2	103
10	Mild Recessive Mutations in Six Fraser Syndrome-Related Genes Cause Isolated Congenital Anomalies of the Kidney and Urinary Tract. <i>Journal of the American Society of Nephrology: JASN</i> , 2014, 25, 1917-1922.	6.1	97
11	Sequencing the GRHL3 Coding Region Reveals Rare Truncating Mutations and a Common Susceptibility Variant for Nonsyndromic Cleft Palate. <i>American Journal of Human Genetics</i> , 2016, 98, 755-762.	6.2	92
12	Muscarinic Acetylcholine Receptor M3 Mutation Causes Urinary Bladder Disease and a Prune-Belly-like Syndrome. <i>American Journal of Human Genetics</i> , 2011, 89, 668-674.	6.2	89
13	An Approach to the Identification of Anomalies and Etiologies in Neonates with Identified or Suspected VACTERL (Vertebral Defects, Anal Atresia, Tracheo-Esophageal Fistula with Esophageal) Tj ETQq1 1 0.784314 rgBT/Overlo 2014. 164. 451-457.e1.	1.8	83
14	Clinical geneticists' views of VACTERL/VATER association. <i>American Journal of Medical Genetics, Part A</i> , 2012, 158A, 3087-3100.	1.2	78
15	Mutations in TBX18 Cause Dominant Urinary Tract Malformations via Transcriptional Dysregulation of Ureter Development. <i>American Journal of Human Genetics</i> , 2015, 97, 291-301.	6.2	72
16	Mutations of the SLIT2-ROBO2 pathway genes SLIT2 and SRGAP1 confer risk for congenital anomalies of the kidney and urinary tract. <i>Human Genetics</i> , 2015, 134, 905-916.	3.8	62
17	Research perspectives in the etiology of congenital anorectal malformations using data of the International Consortium on Anorectal Malformations: evidence for risk factors across different populations. <i>Pediatric Surgery International</i> , 2010, 26, 1093-1099.	1.4	58
18	De novo microduplication at 22q11.21 in a patient with VACTERL association. <i>European Journal of Medical Genetics</i> , 2011, 54, 9-13.	1.3	56

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19	Seven new cases of familial isolated bladder exstrophy and epispadias complex (BEEC) and review of the literature. <i>American Journal of Medical Genetics Part A</i> , 2003, 120A, 215-221.	2.4	54
20	Concordance analyses of twins with bladder exstrophyâ€“epispadias complex suggest genetic etiology. <i>American Journal of Medical Genetics, Part A</i> , 2007, 143A, 2751-2756.	1.2	53
21	Bladder exstrophyâ€“epispadias complex. <i>Birth Defects Research Part A: Clinical and Molecular Teratology</i> , 2009, 85, 509-522.	1.6	53
22	Rare Variants in BNC2 Are Implicated in Autosomal-Dominant Congenital Lower Urinary-Tract Obstruction. <i>American Journal of Human Genetics</i> , 2019, 104, 994-1006.	6.2	47
23	Targeted Resequencing of 29 Candidate Genes and Mouse Expression Studies Implicate <i>ZIC3</i> and <i>FOXF1</i> in Human VATER/VACTERL Association. <i>Human Mutation</i> , 2015, 36, 1150-1154.	2.5	46
24	Gender-associated differences in the psychosocial and developmental outcome in patients affected with the bladder exstrophy-epispadias complex. <i>BJU International</i> , 2006, 97, 349-353.	2.5	45
25	Microduplications at 22q11.21 are associated with non-syndromic classic bladder exstrophy. <i>European Journal of Medical Genetics</i> , 2010, 53, 55-60.	1.3	45
26	Melatonin Concentrations and Antioxidative Capacity of Human Breast Milk According to Gestational Age and the Time of Day. <i>Journal of Human Lactation</i> , 2016, 32, NP105-NP110.	1.6	45
27	Infancy-Onset T1DM, Short Stature, and Severe Immunodysregulation in Two Siblings With a Homozygous LRBA Mutation. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2016, 101, 898-904.	3.6	43
28	Histone H3.3 beyond cancer: Germline mutations in <i>Histone 3 Family 3A and 3B</i> cause a previously unidentified neurodegenerative disorder in 46 patients. <i>Science Advances</i> , 2020, 6, .	10.3	43
29	Genome-wide Association Study and Meta-Analysis Identify <i>ISL1</i> as Genome-wide Significant Susceptibility Gene for Bladder Exstrophy. <i>PLoS Genetics</i> , 2015, 11, e1005024.	3.5	41
30	<i>ISL1</i> is a major susceptibility gene for classic bladder exstrophy and a regulator of urinary tract development. <i>Scientific Reports</i> , 2017, 7, 42170.	3.3	41
31	Genome-wide linkage scan of nonsyndromic orofacial clefting in 91 families of central European origin. <i>American Journal of Medical Genetics, Part A</i> , 2009, 149A, 2680-2694.	1.2	38
32	De novo microduplications at 1q41, 2q37.3, and 8q24.3 in patients with VATER/VACTERL association. <i>European Journal of Human Genetics</i> , 2013, 21, 1377-1382.	2.8	38
33	Familial occurrence of the VATER/VACTERL association. <i>Pediatric Surgery International</i> , 2012, 28, 725-729.	1.4	36
34	Genetics of Bladder-Exstrophy-Epispadias Complex (BEEC): Systematic Elucidation of Mendelian and Multifactorial Phenotypes. <i>Current Genomics</i> , 2015, 17, 4-13.	1.6	36
35	Overlapping <i>SETBP1</i> gain-of-function mutations in Schinzel-Giedion syndrome and hematologic malignancies. <i>PLoS Genetics</i> , 2017, 13, e1006683.	3.5	35
36	<i>TGFB3</i> displays parent-of-origin effects among central Europeans with nonsyndromic cleft lip and palate. <i>Journal of Human Genetics</i> , 2008, 53, 656-661.	2.3	34

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37	Underlying genetic factors of the VATER/VACTERL association with special emphasis on the "Renal" phenotype. <i>Pediatric Nephrology</i> , 2016, 31, 2025-2033.	1.7	34
38	Circulating microRNAs are associated with Pulmonary Hypertension and Development of Chronic Lung Disease in Congenital Diaphragmatic Hernia. <i>Scientific Reports</i> , 2018, 8, 10735.	3.3	34
39	Phenotype Severity in the Bladder Exstrophy-Epispadias Complex: Analysis of Genetic and Nongenetic Contributing Factors in 441 Families from North America and Europe. <i>Journal of Pediatrics</i> , 2011, 159, 825-831.e1.	1.8	33
40	De novo 13q deletions in two patients with mild anorectal malformations as part of VATER/VACTERL and VATER/VACTERL-like association and analysis of <i>EFNB2</i> in patients with anorectal malformations. <i>American Journal of Medical Genetics, Part A</i> , 2013, 161, 3035-3041.	1.2	32
41	Sexual function in adult patients with classic bladder exstrophy: A multicenter study. <i>Journal of Pediatric Urology</i> , 2015, 11, 125.e1-125.e6.	1.1	30
42	Inheritance of the VATER/VACTERL association. <i>Pediatric Surgery International</i> , 2012, 28, 681-685.	1.4	29
43	Copy number variations in 375 patients with oesophageal atresia and/or tracheoesophageal fistula. <i>European Journal of Human Genetics</i> , 2016, 24, 1715-1723.	2.8	27
44	VATER/VACTERL association. <i>Clinical Dysmorphology</i> , 2012, 21, 191-195.	0.3	26
45	Array-based molecular karyotyping in 115 VATER/VACTERL and VATER/VACTERL-like patients identifies disease-causing copy number variations. <i>Birth Defects Research</i> , 2017, 109, 1063-1069.	1.5	26
46	Mutations of the Transcriptional Corepressor ZMYM2 Cause Syndromic Urinary Tract Malformations. <i>American Journal of Human Genetics</i> , 2020, 107, 727-742.	6.2	25
47	Involvement of the WNT and FGF signaling pathways in non-isolated anorectal malformations: Sequencing analysis of WNT3A, WNT5A, WNT11, DACT1, FGF10, FGFR2 and the T gene. <i>International Journal of Molecular Medicine</i> , 2012, 30, 1459-1464.	4.0	24
48	Whole-Exome Sequencing in Nine Monozygotic Discordant Twins. <i>Twin Research and Human Genetics</i> , 2016, 19, 60-65.	0.6	24
49	Array-based molecular karyotyping in fetuses with isolated brain malformations identifies disease-causing CNVs. <i>Journal of Neurodevelopmental Disorders</i> , 2016, 8, 11.	3.1	24
50	WNT3 involvement in human bladder exstrophy and cloaca development in zebrafish. <i>Human Molecular Genetics</i> , 2015, 24, 5069-5078.	2.9	23
51	Investigation of FGF10 as a candidate gene in patients with anorectal malformations and exstrophy of the cloaca. <i>Pediatric Surgery International</i> , 2008, 24, 893-897.	1.4	22
52	Assisted Reproductive Techniques and Risk of Exstrophy-Epispadias Complex: A German Case-Control Study. <i>Journal of Urology</i> , 2013, 189, 1524-1529.	0.4	22
53	The Challenges of the European Anorectal Malformations-Net Registry. <i>European Journal of Pediatric Surgery</i> , 2015, 25, 481-487.	1.3	22
54	Lowe syndrome/Dent-2 disease: A comprehensive review of known and novel aspects. <i>Journal of Pediatric Genetics</i> , 2015, 02, 053-068.	0.7	22

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55	SLC20A1 Is Involved in Urinary Tract and Urorectal Development. <i>Frontiers in Cell and Developmental Biology</i> , 2020, 8, 567.	3.7	22
56	Biallelic and monoallelic variants in PLXNA1 are implicated in a novel neurodevelopmental disorder with variable cerebral and eye anomalies. <i>Genetics in Medicine</i> , 2021, 23, 1715-1725.	2.4	22
57	Classic bladder exstrophy: Frequent 22q11.21 duplications and definition of a 414 kb phenocritical region. <i>Birth Defects Research Part A: Clinical and Molecular Teratology</i> , 2014, 100, 512-517.	1.6	21
58	Exome sequencing in syndromic brain malformations identifies novel mutations in <i>ACTB</i> , and <i>SLC9A6</i> , and suggests <i>BAZ1A</i> as a new candidate gene. <i>Birth Defects Research</i> , 2018, 110, 587-597.	1.5	21
59	<i>ANKRD11</i> variants: KBG syndrome and beyond. <i>Clinical Genetics</i> , 2021, 100, 187-200.	2.0	21
60	Genome-wide linkage scan for bladder exstrophy-epispadias complex. <i>Birth Defects Research Part A: Clinical and Molecular Teratology</i> , 2009, 85, 174-178.	1.6	20
61	De novo duplication of 18p11.21-18q12.1 in a female with anorectal malformation. <i>American Journal of Medical Genetics, Part A</i> , 2011, 155, 445-449.	1.2	20
62	Family-Based Association Study of the MTHFR Polymorphism C677T in Patients with Nonsyndromic Cleft Lip and Palate from Central Europe. <i>Cleft Palate-Craniofacial Journal</i> , 2008, 45, 267-271.	0.9	19
63	p63 (TP73L) a key player in embryonic urogenital development with significant dysregulation in human bladder exstrophy tissue. <i>International Journal of Molecular Medicine</i> , 2010, 26, 861-7.	4.0	19
64	Genome-wide expression profiling of urinary bladder implicates desmosomal and cytoskeletal dysregulation in the bladder exstrophy-epispadias complex. <i>International Journal of Molecular Medicine</i> , 2011, 27, 755-65.	4.0	19
65	Genome-wide association study and mouse expression data identify a highly conserved 32 kb intergenic region between WNT3 and WNT9b as possible susceptibility locus for isolated classic exstrophy of the bladder. <i>Human Molecular Genetics</i> , 2014, 23, 5536-5544.	2.9	19
66	Quality of Life after Surgical Treatment for Esophageal Atresia: Long-Term Outcome of 154 Patients. <i>European Journal of Pediatric Surgery</i> , 2017, 27, 443-448.	1.3	19
67	Isolated bladder exstrophy associated with a de novo 0.9 Mb microduplication on chromosome 19p13.12. <i>Birth Defects Research Part A: Clinical and Molecular Teratology</i> , 2013, 97, 133-139.	1.6	18
68	Genome-wide mapping of copy number variations in patients with both anorectal malformations and central nervous system abnormalities. <i>Birth Defects Research Part A: Clinical and Molecular Teratology</i> , 2015, 103, 235-242.	1.6	18
69	CAKUT and Autonomic Dysfunction Caused by Acetylcholine Receptor Mutations. <i>American Journal of Human Genetics</i> , 2019, 105, 1286-1293.	6.2	18
70	Nine new twin pairs with esophageal atresia: A review of the literature and performance of a twin study of the disorder. <i>Birth Defects Research Part A: Clinical and Molecular Teratology</i> , 2012, 94, 182-186.	1.6	17
71	A Genetics-First Approach Revealed Monogenic Disorders in Patients With ARM and VACTERL Anomalies. <i>Frontiers in Pediatrics</i> , 2020, 8, 310.	1.9	17
72	DNA Methylation and Bladder Cancer: Where Genotype does not Predict Phenotype. <i>Current Genomics</i> , 2020, 21, 34-36.	1.6	17

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73	Bladder exstrophyâ€“epispadias complex: Investigation of suppressor of variegation, enhancer of zeste and Trithorax (SET) as a candidate gene in a large cohort of patients. <i>Scandinavian Journal of Urology and Nephrology</i> , 2006, 40, 221-224.	1.4	16
74	Autosomal-dominant non-syndromic anal atresia: sequencing of candidate genes, array-based molecular karyotyping, and review of the literature. <i>European Journal of Pediatrics</i> , 2011, 170, 741-746.	2.7	16
75	Early postnatal echocardiographic assessment of pulmonary blood flow in newborns with congenital diaphragmatic hernia. <i>Journal of Perinatal Medicine</i> , 2018, 46, 735-743.	1.4	16
76	Targeted sequencing of 96 renal developmental microRNAs in 1213 individuals from 980 families with congenital anomalies of the kidney and urinary tract. <i>Nephrology Dialysis Transplantation</i> , 2016, 31, 1280-1283.	0.7	15
77	CNV analysis in 169 patients with bladder exstrophy-epispadias complex. <i>BMC Medical Genetics</i> , 2016, 17, 35.	2.1	15
78	<i>HSPA6</i> : A new autosomal recessive candidate gene for the VATER/VACTERL malformation spectrum. <i>Birth Defects Research</i> , 2019, 111, 591-597.	1.5	15
79	Heterozygous <i>FGF8</i> mutations in patients presenting cryptorchidism and multiple VATER/VACTERL features without limb anomalies. <i>Birth Defects Research Part A: Clinical and Molecular Teratology</i> , 2014, 100, 750-759.	1.6	14
80	More than fetal urine: enteral uptake of amniotic fluid as a major predictor for fetal growth during late gestation. <i>European Journal of Pediatrics</i> , 2016, 175, 825-831.	2.7	14
81	Esophageal Atresia with or without Tracheoesophageal Fistula (EA/TEF): Association of Different EA/TEF Subtypes with Specific Co-occurring Congenital Anomalies and Implications for Diagnostic Workup. <i>European Journal of Pediatric Surgery</i> , 2018, 28, 176-182.	1.3	14
82	MTHFR 677 TT genotype in a mother and her child with Down syndrome, atrioventricular canal and exstrophy of the bladder: implications of a mutual genetic risk factor?. <i>European Journal of Pediatrics</i> , 2006, 165, 566-568.	2.7	13
83	Parental risk factors of anorectal malformations: Analysis with a regional populationâ€“based control group. <i>Birth Defects Research Part A: Clinical and Molecular Teratology</i> , 2016, 106, 133-141.	1.6	13
84	Arrayâ€“based molecular karyotyping in fetal brain malformations: Identification of novel candidate genes and chromosomal regions. <i>Birth Defects Research Part A: Clinical and Molecular Teratology</i> , 2016, 106, 16-26.	1.6	13
85	Parental risk factors for congenital diaphragmatic hernia â€“ a large German case-control study. <i>BMC Pediatrics</i> , 2021, 21, 278.	1.7	13
86	Genome-wide analysis for micro-aberrations in familial exstrophy of the bladder using array-based comparative genomic hybridization. <i>BJU International</i> , 2007, 100, 646-650.	2.5	12
87	Second study on the recurrence risk of isolated esophageal atresia with or without tracheaâ€“esophageal fistula among firstâ€“degree relatives: No evidence for increased risk of recurrence of EA/TEF or for malformations of the VATER/VACTERL association spectrum. <i>Birth Defects Research Part A: Clinical and Molecular Teratology</i> , 2013, 97, 786-791.	1.6	12
88	Antioxidative status and oxidative stress in the fetal circulation at birth: the effects of time of delivery and presence of labor. <i>Early Human Development</i> , 2015, 91, 119-124.	1.8	12
89	Unraveling the Genetics of Congenital Diaphragmatic Hernia: An Ongoing Challenge. <i>Frontiers in Pediatrics</i> , 2021, 9, 800915.	1.9	12
90	Bladder exstrophy and Epstein type congenital macrothrombocytopenia: Evidence for a common cause?. <i>American Journal of Medical Genetics, Part A</i> , 2006, 140A, 2251-2253.	1.2	11

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91	Evaluation of two commercially available ELISA kits for the determination of melatonin concentrations in amniotic fluid throughout pregnancy. <i>Annals of Clinical Biochemistry</i> , 2017, 54, 107-112.	1.6	11
92	Nonsyndromic cleft palate: An association study at GWAS candidate loci in a multiethnic sample. <i>Birth Defects Research</i> , 2018, 110, 871-882.	1.5	11
93	A Classic Twin Study of Isolated Gastroschisis. <i>Fetal and Pediatric Pathology</i> , 2012, 31, 324-330.	0.7	10
94	Evidence for annular pancreas as an associated anomaly in the VATER/VACTERL association and investigation of the gene encoding pancreas specific transcription factor 1A as a candidate gene. <i>American Journal of Medical Genetics, Part A</i> , 2014, 164, 1611-1613.	1.2	10
95	Epidemiologic analysis of families with isolated anorectal malformations suggests high prevalence of autosomal dominant inheritance. <i>Orphanet Journal of Rare Diseases</i> , 2017, 12, 180.	2.7	10
96	Classic bladder exstrophy and adenocarcinoma of the bladder: Methylome analysis provide no evidence for underlying disease-mechanisms of this association. <i>Cancer Genetics</i> , 2019, 235-236, 18-20.	0.4	10
97	Currarino syndrome: a comprehensive genetic review of a rare congenital disorder. <i>Orphanet Journal of Rare Diseases</i> , 2021, 16, 167.	2.7	10
98	Bias in patient series with VACTERL association. <i>American Journal of Medical Genetics, Part A</i> , 2011, 155, 2039-2041.	1.2	9
99	CNV Analysis in Monozygotic Twin Pairs Discordant for Urorectal Malformations. <i>Twin Research and Human Genetics</i> , 2013, 16, 802-807.	0.6	9
100	Targeted Resequencing of Putative Growth-Related Genes Using Whole Exome Sequencing in Patients with Severe Primary IGF-I Deficiency. <i>Hormone Research in Paediatrics</i> , 2017, 88, 408-417.	1.8	9
101	Sexual Function and Quality of Life in Adult Male Individuals with Exstrophy-Epispadias Complex—a Survey of the German CURE-Network. <i>Urology</i> , 2018, 112, 215-221.	1.0	9
102	Association Between Exstrophy-epispadias Complex And Congenital Anomalies: A German Multicenter Study. <i>Urology</i> , 2019, 123, 210-220.	1.0	9
103	Human exome and mouse embryonic expression data implicate ZFX3, TRPS1, and CHD7 in human esophageal atresia. <i>PLoS ONE</i> , 2020, 15, e0234246.	2.5	9
104	Candidate gene association study implicates <i>p63</i> in the etiology of nonsyndromic bladder exstrophy-epispadias complex. <i>Birth Defects Research Part A: Clinical and Molecular Teratology</i> , 2013, 97, 759-763.	1.6	8
105	Lessons Learned from CNV Analysis of Major Birth Defects. <i>International Journal of Molecular Sciences</i> , 2020, 21, 8247.	4.1	8
106	The Genomic Architecture of Bladder Exstrophy Epispadias Complex. <i>Genes</i> , 2021, 12, 1149.	2.4	8
107	Low maternal folate concentrations and maternal MTHFR C677T polymorphism are associated with an increased risk for neural tube defects in offspring: a case-control study among Pakistani case and control mothers. <i>Asia Pacific Journal of Clinical Nutrition</i> , 2018, 27, 253-260.	0.4	8
108	Embryonic expression of the cysteine rich protein 61 (<i>CYR61</i>) gene: A candidate for the development of human epispadias. <i>Birth Defects Research Part A: Clinical and Molecular Teratology</i> , 2010, 88, 546-550.	1.6	7

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109	Evidence for linkage of the bladder exstrophy-epispadias complex on chromosome 4q31.21-22 and 19q13.31-41 from a consanguineous Iranian family. <i>Birth Defects Research Part A: Clinical and Molecular Teratology</i> , 2010, 88, 757-761.	1.6	7
110	Evaluation of sexual function in females with exstrophy-epispadias-complex: a survey of the multicenter German CURE-Net. <i>Journal of Pediatric Urology</i> , 2016, 13, 183.e1-183.e6.	1.1	7
111	Congenital intrahepatic portocaval shunts and hypoglycemia due to secondary hyperinsulinism: a case report and review of the literature. <i>Journal of Medical Case Reports</i> , 2018, 12, 336.	0.8	7
112	Male infant with paternal uniparental diploidy mosaicism and a 46,XX/46,XY karyotype. <i>American Journal of Medical Genetics, Part A</i> , 2019, 179, 2252-2256.	1.2	7
113	Family-based association study of the MTHFR polymorphism C677T in the bladder-exstrophy-epispadias-complex. <i>American Journal of Medical Genetics, Part A</i> , 2006, 140A, 2506-2509.	1.2	6
114	A family-based association study in Central Europeans: No evidence for the cystathionine beta-synthase c.844ins68 gene variant as a risk factor for non-syndromic cleft lip and palate. <i>American Journal of Medical Genetics, Part A</i> , 2007, 143A, 205-207.	1.2	6
115	Scimitar syndrome in a case with VACTERL association. <i>Cardiology in the Young</i> , 2015, 25, 606-609.	0.8	6
116	Towards a Central Role of ISL1 in the Bladder Exstrophy-epispadias Complex (BEEC): Computational Characterization of Genetic Variants and Structural Modelling. <i>Genes</i> , 2018, 9, 609.	2.4	6
117	A classic twin study of lower urinary tract obstruction: Report of 3 cases and literature review. <i>LUTS: Lower Urinary Tract Symptoms</i> , 2019, 11, O85-O88.	1.3	6
118	Treatment Strategies and Outcome of the Exstrophy-epispadias Complex in Germany: Data From the German CURE-Net. <i>Frontiers in Pediatrics</i> , 2020, 8, 174.	1.9	6
119	Exome survey of individuals affected by VATER / VACTERL with renal phenotypes identifies phenocopies and novel candidate genes. <i>American Journal of Medical Genetics, Part A</i> , 2021, 185, 3784-3792.	1.2	6
120	Re-sequencing of candidate genes <i>FOXF1</i> , <i>HSPA6</i> , <i>HAAO</i> , and <i>KYNU</i> in 522 individuals with VATER/VACTERL, VACTERL, VACTERL/VACTERL-like association, and isolated anorectal malformation. <i>Birth Defects Research</i> , 2022, 114, 478-486.	1.5	6
121	Definition, diagnosis and clinical management of non-obstructive kidney dysplasia: a consensus statement by the ERKNet Working Group on Kidney Malformations. <i>Nephrology Dialysis Transplantation</i> , 2022, 37, 2351-2362.	0.7	6
122	Possible association of Down syndrome and exstrophy-epispadias complex: report of two new cases and review of the literature. <i>European Journal of Pediatrics</i> , 2009, 168, 881-883.	2.7	5
123	Primary pulmonary hypertension, congenital heart defect, central nervous system malformations, hypo- and aplastic toes: Another case of Yunis-Varian syndrome or report of a new entity. <i>European Journal of Medical Genetics</i> , 2012, 55, 27-31.	1.3	5
124	Clinical and echocardiographic risk factors for extubation failure in infants with congenital diaphragmatic hernia. <i>Paediatric Anaesthesia</i> , 2018, 28, 864-872.	1.1	5
125	The Role of De Novo Variants in Patients with Congenital Diaphragmatic Hernia. <i>Genes</i> , 2021, 12, 1405.	2.4	5
126	Successful ECMO therapy in a child with COVID-19-associated ARDS and acute lymphoblastic leukemia. <i>Pediatric Blood and Cancer</i> , 2021, 68, e29100.	1.5	4

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127	Genome-Wide Survey for Microdeletions or -Duplications in 155 Patients with Lower Urinary Tract Obstructions (LUTO). <i>Genes</i> , 2021, 12, 1449.	2.4	4
128	A Prevalence Estimation of Exstrophy and Epispadias in Germany From Public Health Insurance Data. <i>Frontiers in Pediatrics</i> , 2021, 9, 648414.	1.9	4
129	First genome-wide association study of esophageal atresia identifies three genetic risk loci at CTNNA3, FOXF1/FOXC2/FOXL1, and HNF1B. <i>Human Genetics and Genomics Advances</i> , 2022, 3, 100093.	1.7	4
130	Transforming growth factor-beta receptor type 1 (TGFB1) is not associated with non-syndromic cleft lip with or without cleft palate in patients of Central European descent. <i>International Journal of Pediatric Otorhinolaryngology</i> , 2009, 73, 1334-1338.	1.0	3
131	Mutations in PTF1A are not a common cause for human VATER/VACTERL association or neural tube defects mirroring Danforth's short tail mouse. <i>Molecular Medicine Reports</i> , 2015, 12, 1579-1583.	2.4	3
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