

Veröffentlichungen (Stand: 12/2023)

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Originalveröffentlichungen in Zeitschriften mit wissenschaftlichem Beirat

Gesamtzahl 127, davon 26 primo loco; Impactpunkte > 312_{ISI}

1. Hageman IC, Midrio P, van der Steeg HJJ, Jenetzky E, Iacobelli BD, Morandi A, Sloots CEJ, Schmiedeke E, Broens PMA, Fascetti Leon F, Çavuşoğlu YH, Gorter R, Trajanovska M, King SK, Aminoff D, Schwarzer N, Haanen M, de Blaauw I, van Rooij IALM and ARM-Net Consortium. More than ten years of the European Anorectal Malformation Network (ARM-Net) patient registry: clinical and surgical characteristics of 2619 patients. *British Journal of Surgery*. 2023. accepted (IF₂₀₂₃= 11.122; Hw. A)
2. Berger B, Egert V, Martin D, Jenetzky E. Inpatient Fasting Options for People with T1d in German-Speaking Countries. *J Hum Nutr Food Sci* 2023; 11(3): 1167. <https://doi.org/10.47739/2333-6706/1167> (IF₂₀₂₃=0; Short Communication)
3. Hageman IC, van der Steeg HJJ, Jenetzky E, Trajanovska M, King SK, de Blaauw I, van Rooij IALM; ARM-Net Consortium. A Quality Assessment of the ARM-Net Registry Design and Data Collection. *J Pediatr Surg*. 2023; 58(10):1921-1928. <https://doi.org/10.1016/j.jpedsurg.2023.02.049> . accepted (IF₂₀₂₃= 2.003; Hw. B)
4. Köllges R, Stegmann J, Schneider S, Waffenschmidt L, Fazaal J, Breuer K, Hilger AC, Dworschak GC, Mingardo E, Rösch W, Hofmann A, Neissner C, Ebert AK, Stein R, Younsi N, Hirsch-Koch K, Schmiedeke E, Zwink N, Jenetzky E, Thiele H, Ludwig KU, Reutter H. Exome Survey and Candidate Gene Re-Sequencing Identifies Novel Exstrophy Candidate Genes and Implicates LZTR1 in Disease Formation. *Biomolecules*. 2023 Jul 13;13(7):1117. <https://doi.org/10.3390/biom13071117> IF₂₀₂₂= 5.007; Hw. B)
5. Wrenger J, Martin DD, Jenetzky E. Infants' immunisations, their timing and the risk of allergic diseases (INITIAL): an observational prospective cohort study protocol. *BMJ Open*. 2023 Jun 23;13(6):e072722. <https://doi.org/10.1136/bmjopen-2023-072722> (IF₂₀₂₃= 3.007; Hw. B)
6. Hamideh Kerdar S, Gwiada M, Berger B, Rathjens L, Schwarz S, Jenetzky E, Martin DD. Predictors of sustained use of mobile health applications: Content analysis of user perspectives from a fever management app. *Digit Health*. 2023 Jun 5;9:20552076231180418. <https://doi.org/10.1177/20552076231180418> .
7. Stausberg J, Harkener S, Engel C, Finger R, Heinz C, Jenetzky E, Jersch P, Martin D, Rupp R, Schoenthaler M, Suwelack B, Wegner J. Cross-Registry Benchmarking of Data Quality: Lessons Learned. *Stud Health Technol Inform*. 2023 May 18;302:167-171. <https://doi.org/10.3233/SHTI230096>
8. Möhler R, Jenetzky E, Schwarz S, Gwiada M, Rathjens L, Martin DD. Children's Symptoms with a Febrile Illness and a Positive or Negative Test of SARS-CoV-2 during the Omicron Wave. *Children (Basel)*. 2023 Feb 22;10(3):419. <https://doi.org/10.3390/children10030419>
9. Rathjens L, Gwiada M, Schwarz S, Möhler R, Martin DD, Jenetzky E. Reliability of Data Collected via Ecological Momentary Assessment on the Example of FeverApp

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 12. Fabian J, Dworschak GC, Waffenschmidt L, Schierbaum L, Bendixen C, Heilmann-Heimbach S, Sivalingam S, Buness A, Schwarzer N, Boemers TM, Schmiedeke E, Neser J, Leonhardt J, Kosch F, Weih S, Gielen HM, Hosie S, Kabs C, Palta M, Märzheuser S, Bode LM, Lacher M, Schäfer FM, Stehr M, Knorr C, Ure B, Kleine K, Rolle U, Zaniew M, Phillip G, Zwink N, **Jenetzky E**, Reutter H, Hilger AC. Genome-wide identification of disease-causing copy number variations in 450 individuals with anorectal malformations. *Eur J Hum Genet*. 2023; 31(1):105-111. . <https://doi.org/10.1038/s41431-022-01216-5>
 13. Mingardo E, Beaman G, Grote P, Nordenskjöld A, Newman W, Woolf AS, Eckstein M, Hilger AC, Dworschak GC, Rösch W, Ebert AK, Stein R, Brusco A, Di Grazia M, Tamer A, Torres FM, Hernandez JL, Erben P, Maj C, Olmos JM, Riancho JA, Valero C, Hostettler IC, Houlden H, Werring DJ, Schumacher J, Gehlen J, Giel AS, Buerfent BC, Arkani S, Åkesson E, Rotstein E, Ludwig M, Holmdahl G, Giorgio E, Berettini A, Keene D, Cervellione RM, Younsi N, Ortlieb M, Oswald J, Haid B, Promm M, Neissner C, Hirsch K, Stehr M, Schäfer FM, Schmiedeke E, Boemers TM, van Rooij IALM, Feitz WFJ, Marcelis CLM, Lacher M, Nelson J, Ure B, Fortmann C, Gale DP, Chan MMY, Ludwig KU, Nöthen MM, Heilmann S, Zwink N, **Jenetzky E**, Odermatt B, Knapp M, Reutter H. A genome-wide association study with tissue transcriptomics identifies genetic drivers for classic bladder exstrophy. *Commun Biol*. 2022 Nov 9;5(1):1203. <https://doi.org/10.1038/s42003-022-04092-3>. (IF = 6.548)
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 15. Rathjens L, Fingerhut I, Martin, D, Hamideh Kerdar S, Gwiasda M, Schwarz S, **Jenetzky E**. Data Completeness and Concordance in the FeverApp Registry: Comparative Study. *JMIR Pediatr Parent*. 2022; 5(4): e35510. <https://doi.org/10.2196/35510> (IF=4.0)
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