

Publikationen mit Beteiligung NENT 2017-2021 für die ARAMIS-Datenbank
(*Listeria monocytogenes*, STEC, Salmonella)

Nüesch-Inderbinen, M., Bloemberg, G., Müller, A., Stevens, M., Cernela, N., Kollöffel, B., Stephan, R. (2021). Listeriosis caused by persistence of *Listeria monocytogenes* serotype 4b Sequence Type 6 in cheese production environment. *Emerging Infectious Diseases* 1,284-288. doi:10.3201/eid2701.203266.

Nüesch-Inderbinen, M., Stevens, M.J.A., Cernela, N., Müller, A., Biggel, M. Stephan, R. (2021). Distribution of virulence factors, antimicrobial resistance genes and phylogenetic relatedness among Shiga toxin-producing *Escherichia coli* serogroup O91 strains from human infections. *International Journal of Medical Microbiology* 8, 151541. <https://doi.org/10.1016/j.ijmm.2021.151541>

Sah, R., Donovan, S., Seth-Smith, HMB, **Bloemberg, G., Wüthrich, D., Stephan, R., Kataria, S., Kumar, M., Singla, S., Deswal, V., Kaur, A., Neumayr, A., Hinic, V., Egli, A., Kuenzli, E.** (2020). A novel lineage of ceftriaxone-resistant *Salmonella* Typhi from India that is closely related to XDR *S. Typhi* found in Pakistan. *Clinical Infectious Diseases* 71(5),1327-1330. doi: 10.1093/cid/ciz1204

Nüesch-Inderbinen, M., Cernela, N., Wüthrich, C., Egli, A., Stephan, R. (2018). Genetic characterization of Shiga toxin producing *Escherichia coli* belonging to the emerging hybrid pathotype O80:H2 isolated from humans 2010-2017 in Switzerland. *International Journal of Medical Microbiology* 308, 534-538. DOI: 10.1016/j.ijmm.2018.05.007

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