

Poster number	Poster session	Presenting author	Title
P-1	P-I	Uzal, Francisco	A retrospective study of <i>Clostridioides difficile</i> infection in horses
P-2	P-I	Barbut, Frédéric	Evaluation of Vivalytic <i>C. difficile</i> assay and Vivalytic noro-, rotavirus, <i>C. difficile</i> assay for the detection of toxigenic <i>Clostridioides difficile</i> strains in stools
P-3	P-I	Barbut, Frédéric	Intra-hospital transmission of <i>Clostridioides difficile</i> using whole-genome sequencing
P-4	P-I	Barbut, Frédéric	Epidemiological trends in <i>Clostridioides difficile</i> infections in France, 2019–2024: TRACE national population-based study
P-5	P-I	Savidge, Tor	Microbiome-informed diagnostics distinguish <i>Clostridioides difficile</i> infection from asymptomatic carriage
P-6	P-I	Wang, Xianjun	Accelerated detection of <i>Clostridioides difficile</i> sequence type 37 by integrating MALDI-TOF mass spectrometry with artificial neural network
P-7	P-I	Corver, Jeroen	Cwp20 as potential biomarker to detect <i>Clostridioides difficile</i> in feces of CDI patients
P-8	P-I	Kimani, Andrew Mwaura	<i>Clostridioides difficile</i> in the dairy farm environment of southwest Australia
P-9	P-I	Anantharajah, Ahalieyah	Exploring the circulation of <i>Clostridioides difficile</i> in Brussels wastewater
P-10	P-I	Anantharajah, Ahalieyah	Distinct clinical and antimicrobial resistance profiles of binary toxin genes–positive <i>Clostridioides difficile</i> isolates circulating in Belgium
P-11	P-I	Mrozinski, Alexandre	Treated wastewater irrigation and pathogen dissemination across the food chain and environment: a One Health study of <i>Clostridioides difficile</i>
P-12	P-I	Kalogeropoulou, Eleni	Increasing burden of <i>Clostridioides difficile</i> infection: a twelve-year surveillance study, 2014–2025
P-13	P-I	Angulo, Frederick	Recent increase in the incidence of <i>Clostridioides difficile</i> infection (CDI) in most European countries which conduct population-based CDI surveillance, 2017–2024
P-14	P-I	Angulo, Frederick	Systematic literature review of the <i>Clostridioides difficile</i> infection (CDI) disease burden in mainland China
P-15	P-I	Angulo, Frederick	Protocol for a prospective multi-hospital study of the <i>Clostridioides difficile</i> infection (CDI) disease burden in five cities in China
P-16	P-I	Guo, Linwen	Investigation on the current status of <i>Clostridioides difficile</i> infection prevention and control in Chinese healthcare facilities: a cross-sectional survey of 2,306 healthcare facilities
P-17	P-I	Bassères, Eugénie	Comparative <i>in vitro</i> killing kinetics of Ibezapolstat and comparator antibiotics in <i>Clostridioides difficile</i> and <i>Enterococcus</i> co-cultures
P-18	P-I	Janezic, Sandra	Diversity of <i>Clostridioides difficile</i> in marine sediments of eastern Adriatic Sea
P-19	P-I	Reichl, Julia	The rise and fall of RT027 – the changing ribotype landscape of <i>Clostridioides difficile</i> in Austria, 2008-2024

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P-20	P-I	Callies, Milena	Integrating antimicrobial consumption and <i>Clostridioides difficile</i> infection surveillance in Belgian hospitals: a decade-long retrospective study
P-21	P-I	McCarthy, Natalie	Rebound in estimated number of <i>Clostridioides difficile</i> infection cases, hospitalizations, and deaths in the United States following the COVID-19 pandemic, 2017–2023
P-22	P-I	Hain-Saunders, Natasza	Community, companions and <i>Clostridioides difficile</i> – a case of <i>C. difficile</i> infection in the domestic environment
P-23	P-I	Henigman, Urška	Detection of <i>Clostridioides difficile</i> in pasteurized poultry products
P-24	P-I	Wan, Wenbo	Epidemiology and risk factors of hospital-onset <i>Clostridioides difficile</i> infection in adult inpatients: a retrospective case–control study
P-25	P-I	Lonkeu Njouboussi, Yelena Fadela	Assessment of contamination of point-of-sale foodstuffs with <i>Clostridioides difficile</i> in the United Kingdom
P-26	P-I	Stanley, Ann Marie	Understanding <i>C. difficile</i> TcdB conservation through surveillance of diverse and global contemporary isolates
P-27	P-I	Stanley, Ann Marie	AZD5148 prevents <i>C. difficile</i> toxin B–induced cytotoxicity in a human colon transwell system
P-28	P-I	Cocchi, Monia	Comparison of <i>Clostridioides difficile</i> ribotypes from wastewater isolates and clinical isolates: preliminary results of IZS VE 02/24 RC research project
P-29	P-I	Jwalane, Sephiri	Clinical and wastewater <i>Clostridioides difficile</i> profiles obtained from Cape Town, South Africa
P-30	P-I	Jin, Dazhi	Molecular epidemiology of <i>Clostridioides difficile</i> colonization in patients with chronic kidney disease in Eastern China
P-31	P-I	Jin, Dazhi	Sodium metabisulfite enhances <i>Clostridioides difficile</i> virulence through activation of the LytR regulator
P-32	P-I	Jin, Dazhi	Nitroreductase Nfp-mediated metronidazole resistance in <i>Clostridioides difficile</i>
P-33	P-I	Boudaher, Elizabeth	<i>Clostridioides difficile</i> in dairy cattle
P-34	P-I	Androga, Grace	Sealing the anaerobe gap in Africa
P-35	P-I	van Prehn, Joffrey	<i>Clostridioides difficile</i> infection sentinel surveillance in the Netherlands: epidemiology, WGS, and ribotyping
P-36	P-I	Cerri, Fabrício	Comparison of four culture media for isolation of <i>Clostridioides difficile</i> from neonatal foal feces
P-37	P-I	Kalkeri, Raj	Novel toxin- and non-toxin-based adjuvanted protein vaccines for <i>C. difficile</i> protect against toxin B and spore challenge in mice and hamsters
P-38	P-I	Müller, Irene	Outpatient <i>C. difficile</i> infections: evidence from a German physician survey on clinical management and medical coding practices
P-39	P-I	Seghrouchni Bent Houssine, Imane	Characterization of the regulatory RNA RCd2 in the physiology and the pathogenesis of <i>Clostridioides difficile</i>

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P-40	P-I	Arbeiter, Judith	Exploring the secondary metabolism of <i>Clostridioides difficile</i>
P-41	P-I	Johnson, Katie	Expanding the role of CRISPR-Cas systems in <i>Clostridioides difficile</i>
P-42	P-I	Lalowski, Piotr	Swarming-associated morphological adaptation of <i>Clostridioides difficile</i> cells
P-43	P-I	Candela, Thomas	Elucidating the pleiotropic effects of <i>Clostridioides difficile</i> clpC deletion proteomic and phenotypic insights
P-44	P-I	Candela, Thomas	Structural similarity and immunological cross-reactivity of lipoteichoic acids from <i>Clostridioides difficile</i> and <i>Peptostreptococcus anaerobius</i>
P-45	P-I	Olivença, Carmen	SH3-mediated substrate recognition directs YabG localization and proteolysis during <i>Clostridioides difficile</i> sporulation
P-46	P-I	Hamrock, Fergal	Spatially resolved analysis of sporulation decision-making in <i>Clostridioides difficile</i>
P-47	P-II	Hinc, Krzysztof	Phi027 prophage shapes <i>Clostridioides difficile</i> virulence through the SinR/SinR' axis and the flagellar switch network
P-48	P-II	Damsbo Jensen, Anna	Dimeric VHH binding proteins mitigate <i>Clostridioides difficile</i> toxin-mediated epithelial damage
P-49	P-II	Rodriguez-Rodriguez, Everardo	<i>C. difficile</i> toxins and their interplay with the healthy gut microbiota: sustaining a healthy gut with functional IgG-derived binding proteins
P-50	P-II	Murthy, Ashlesh	Characterization of genotypic diversity in contemporary <i>Clostridioides difficile</i> isolates from ATLAS 2020–2024
P-51	P-II	Dobрила, Horia	Exogenous butyrate induces <i>C. difficile</i> toxin release and sporulation
P-52	P-II	Petersen, Ida	Characterization of the native autoinducing peptide(s) in <i>C. difficile</i>
P-53	P-II	Petersen, Ida	Functional characterization of the autoinducing peptide(s) in <i>C. difficile</i> – a chemical approach
P-54	P-II	Corver, Jeroen	Dissecting the biosynthetic pathway of the flagellin type A glycan in <i>Clostridioides difficile</i> strain 630Δerm
P-55	P-II	Unnikrishnan, Meera	Smart colon-on-a-chip: a high-fidelity in vitro model for studying <i>C. difficile</i> pathogenesis
P-56	P-II	Lindič, Nataša	Structural basis of secondary polysaccharide biosynthesis in <i>C. difficile</i>
P-57	P-II	Salgado, Paula	S-layer deletion in <i>C. difficile</i> clinically relevant strains
P-58	P-II	Salgado, Paula	Adding functionality to <i>C. difficile</i> 's armour – investigation of CWPs
P-59	P-II	Badilla Lobo, Adriana	D,D-carboxypeptidases control the mode of peptidoglycan transpeptidation in <i>C. difficile</i>
P-60	P-II	Moreira Cerri, Fabricio	Prevalence of multidrug-resistant <i>Clostridioides difficile</i> ribotypes isolated from foals in Brazil
P-61	P-II	Alves, Frederico	Defining tetracycline ECOFF and resistance determinants in <i>Clostridioides difficile</i> across One Health reservoirs

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P-62	P-II	Balíková Novotná, Gabriela	A new antibiotic resistance enzyme for a classic target: MrmA redirects radical SAM methylation to 23S rRNA A2058 in <i>C. difficile</i>
P-63	P-II	Heejung, Kim	Long-term trends in antimicrobial resistance and ribotype distribution of <i>C. difficile</i> isolates from two Korean hospitals, 2018–2025
P-64	P-II	Krutova, Marcela	The prediction of antimicrobial resistance in <i>Clostridioides difficile</i> based on genomic data
P-65	P-II	Natarajan, Saran	Beyond the binding site hidden drivers of fidaxomicin resistance
P-66	P-II	Assaf, Mizrahi	Differential impact of vancomycin and fidaxomicin on digestive clearance of <i>Clostridioides difficile</i>
P-67	P-II	Smit, Louise F. E.	Absence of resistance to anti-CDI antibiotics in patients with recurrent <i>Clostridioides difficile</i> infection in the Netherlands
P-68	P-II	Le Monnier, Alban	<i>Clostridioides difficile</i> immunity during pregnancy and passive antibody transfer to neonates from cord blood and breast milk
P-69	P-II	Le Monnier, Alban	High seroprevalence of anti-toxin antibodies to <i>Clostridioides difficile</i> : a multicenter prospective study across three French regions
P-70	P-II	Rashid, Andy	Functional profiling of microbiome recovery reveals patient-specific restoration of colonization resistance to <i>Clostridioides difficile</i>
P-71	P-II	Javorská, Daniela	Prognostic significance of leukocyte differential counts in <i>Clostridioides difficile</i> infection
P-72	P-II	Javorská, Daniela	Uncovering <i>Clostridioides difficile</i> infection risk in COVID-19 patients through predictive modeling
P-73	P-II	Murdan, Sudaxshina	Sublingual immunisation against <i>C. difficile</i> enhances intestinal IgA while subcutaneous immunisation boosts serum IgG
P-74	P-II	Terveer, Elisabeth M.	Biovigilance in faecal microbiota transplantation: a 7-year cohort study and framework for microbiological assessments of infectious adverse events
P-75	P-II	Lidén, Stina	Do CDI outcomes improve after the introduction of a FMT stool bank? A single-center retrospective cohort study
P-76	P-II	Louie, Thomas	Ingestion of deodorized, double-washed, alcohol-shocked gut microbes without encapsulation or bowel preparation as FMT for young pediatric patients with multiple recurrent <i>C. difficile</i> infection
P-77	P-II	Péchiné, Séverine	<i>Faecalibacterium prausnitzii</i> EXL-01 modulates <i>Clostridioides difficile</i> growth in vitro
P-78	P-II	Paddison Rees, Nia	Individual vs pooled gut microbiome responses to simulated <i>C. difficile</i> PCR ribotype 955 infection
P-79	P-II	Abuzar, Sharif MD	Preclinical safety profile of AJ-024, a novel oral antibiotic for <i>C. difficile</i> infection
P-80	P-II	Birk, Christopher	Evaluating ASOs as antibacterial agents for treatment of <i>C. difficile</i> infection

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P-81	P-II	Meffre, Dorian	Amphiphilic cationic copolypeptoids: novel antibacterial agents targeting <i>Clostridioides difficile</i>
P-82	P-II	Hurdle, Julian G.	Microbiological and physiological analysis of fitness costs of fidaxomicin resistance in <i>C. difficile</i>
P-83	P-II	Jarvis, Thale	Structural insights into the potency, selectivity and fitness cost of resistance to CRS3123, a clinical candidate for treatment of <i>C. difficile</i> infections
P-84	P-II	Shealy, Nicolas	Gut maturation state differentially influences <i>Clostridioides difficile</i> TCS repertoire
P-85	P-II	Riley, Thomas	Prevalence of <i>Clostridioides difficile</i> in meat and meat products in Perth, western Australia
P-86	P-II	Riley, Thomas	Prevalence of <i>Clostridioides difficile</i> in households in Perth, western Australia