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Beatrice Margherita Bonazzi, *Microbiology Unit, IRCSS Azienda Ospedaliero-Universitaria di Bologna, Italy / Alma Mater Studiorum University of Bologna, Italy*

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Jody E Phelan, *Department of Infection Biology, London School of Hygiene and Tropical Medicine, London, UK / Statens Serum Institut, Copenhagen, Denmark*

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Thierry Wirth, *EPHE, PSL University, Paris, France / Institut de Systématique, Evolution, Biodiversité, ISYEB, Muséum national d'Histoire naturelle, CNRS, Sorbonne Université, EPHE, Université des Antilles, Paris, France*

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Sara Truden, *MSc*

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Laima Vasiliauskaite, *Vilnius University, Faculty of Medicine, Institute of Biomedical Sciences, Department of Physiology, Biochemistry, Microbiology and Laboratory Medicine, Vilnius, Lithuania / Centre of Laboratory Medicine, Laboratory of Infectious Diseases and Tuberculosis, Vilnius University Hospital Santaros Klinikos, Vilnius, Lithuania*

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Viola Dreyer, *1. Research Center Borstel - Leibniz Lung Center, Molecular and Experimental Mycobacteriology*

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Sofie M. Braet, *Institute of Tropical Medicine Antwerp, Belgium*

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Arnau Seguí Moll, *Departament de Genètica i Microbiologia, Facultat de Biociències, Universitat Autònoma de Barcelona, 08193, Bellaterra, Barcelona, Spain*

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Federica Sorella, *University of Bologna*

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Miguel de Diego Fuertes, *Tuberculosis Omics Research Consortium, Department of Family Medicine and Population Health, Faculty of Medicine and Health Sciences, University of Antwerp, Antwerp, Belgium*

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Jerome Ndayisenga, *National Reference Laboratory, Biomedical Services Department, Rwanda Biomedical Centre, Kigali, Rwanda*

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petra de Haas, *KNCV TB foundation, The Hague, The Netherlands*

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HP Liang, *Singapore General Hospital*

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KVB Lima, *Bacteriology and Mycology Section, Instituto Evandro Chagas, Ananindeua, Brazil / Postgraduate Program in Parasitic Biology in the Amazon, Universidade do Estado do Pará, Belém, Brazil /Postgraduate Program in Epidemiology and Health Surveillance, Instituto Evandro Chagas, Ananindeua, Brazil / Brazilian Tuberculosis Research Network, Rede Brasileira de Pesquisa em Tuberculose, Rio de Janeiro, Brazil.*

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Zoë Vandenberghe, *Sciensano / University of Antwerp*

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CC Frota, *Pos-graduate Program in Pathology, Faculty of Medicine, Federal University of Ceará, Fortaleza, Brazil*

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Borowiecka Angelika, *Institut de Systématique, Évolution, Biodiversité (ISYEB),EPHE/PSL University/Muséum national d'Histoire naturelle/CNRS/Sorbonne Université/Université des Antilles, Paris*

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Gloria Guerrero Manriquez, *Unidad Académica de Ciencias Biológicas de la Universidad Autónoma de Zacatecas*

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Magdalena Kuzioła, *Institute of Medical Biology, Polish Academy of Sciences, Mycobacterium Genetics and Physiology Lab, Lodz / The Bio-Med-Chem Doctoral School of the University of Lodz and Lodz Institutes of the Polish Academy of Sciences, Lodz*

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Miriam Blay-Benach, *Institut de Recerca i Tecnologia Agroalimentàries (IRTA), Animal Health, Centre de Recerca en Sanitat Animal (CReSA), Campus de la Universitat Autònoma de Barcelona (UAB), Bellaterra, Catalonia, Spain / Unitat Mixta d'Investigació Institut de Recerca i Tecnologia Agroalimentàries (IRTA)-UAB en Sanitat Animal, CReSA, Campus de la Universitat Autònoma de Barcelona (UAB), Bellaterra, Catalonia, Spain*

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Emma Roycroft, *Irish Mycobacteria Reference Laboratory / Trinity College Dublin*

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ANNA BRZOSTEK, *Intitute for Medical Biology Poilish Academy of Sciences*

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Max Kevane-Campbell, *Goethe University Frankfurt, University Hospital, Department of Internal Medicine, Infectious Diseases, Frankfurt am Main, Germany. / Goethe University Frankfurt, Mycobacterial Infection Research Unit (MIRU), Frankfurt am Main, Germany. / Goethe University Frankfurt, University Hospital, Institute of Medical Microbiology and Infection Control, Frankfurt am Main, Germany.*

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Víctor Lorente-Leal, *Departamento de Fisiología, Genética y Microbiología, Facultad de Ciencias Biológicas, Universidad Complutense de Madrid, Madrid, Spain / Centro de Vigilancia Sanitaria Veterinaria (VISAVET), Universidad Complutense de Madrid, Madrid, Spain*

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Noud Hermans, *Diagnostics Team, Division of TB Elimination & Health Systems Innovations, KNCV Tuberculosis Foundation, The Hague, The Netherlands/ Department Bacterial Surveillance and Response, Center for Infectious Disease Control, Diagnostics and Laboratory Surveillance, National Institute for Public Health and the Environment (RIVM), Bilthoven, The Netherlands*

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Jorge Alfredo Morcillo Muñoz, *Hospital Universitario Departamental de Nariño, Pasto, Colombia/ Universidad de Nariño, Pasto, Colombia*

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Esther Julián, *Departament de Genètica i Microbiologia, Facultat de Biociències, Universitat Autònoma de Barcelona, Bellaterra, Barcelona, Spain*

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Elena Graglia, *Infectious Disease department, Nino Baglieri Hospital, Modica, Italy*

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petra de Haas, *KNCV Tuberculosefonds*

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Arnau Seguí Moll, *Departament de Genètica i Microbiologia, Facultat de Biociències, Universitat Autònoma de Barcelona, 08193, Bellaterra, Barcelona, Spain*

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Patricia Cuenca Lara, *Unitat Mixta d'Investigació Institut de Recerca i Tecnologia Agroalimentàries (IRTA)-UAB en Sanitat Animal, CReSA, Campus de la Universitat Autònoma de Barcelona (UAB), Bellaterra, Catalonia, Spain / Institut de Recerca i Tecnologia Agroalimentàries (IRTA), Animal Health, Centre de Recerca en Sanitat Animal (CReSA), Campus de la Universitat Autònoma de Barcelona (UAB), Bellaterra, Catalonia, Spain*

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Mafalda Sarraguça, *LAQ-REQUIMTE, Faculty of Pharmacy, University of Porto, Portugal*

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Paolo Caiazza, *University of Bologna*

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Hee-Jong (Jason) Hwang, *A&J Science, Republic of Korea*

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Syn Yu Chua, *Singapore General Hospital*

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Erik Michael Rasmussen, *International Reference Laboratory of Mycobacteriology, Statens Serum Institut, Copenhagen, Denmark*

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Sofia Santos, *Instituto Nacional de Saúde Dr. Ricardo Jorge - Porto*

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Wilma Quispe Rojas, *Tuberculosis Omics Research Consortium, Department of Family Medicine and Population Health, Faculty of Medicine and Health Sciences, University of Antwerp, Antwerp, Belgium*