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Nota di contenuto	<p>""Front Matter""; ""Reviewers""; ""Acknowledgments""; ""Contents""; ""Tables, Figures, and Boxes""; ""Workshop Overview""; ""Appendix A: Contributed Manuscripts""; ""A1 Effector and memory T cell responses to commensal bacteria--Yasmine Belkaid, Nicolas Bouladoux, and Timothy W. Hand""; ""A2 What are the consequences of the disappearing human microbiota?--Martin J. Blaser and Stanley Falkow""; ""A3 Pathways in microbe-induced obesity--Laura M. Cox and Martin J. Blaser""</p> <p>""A4 Microbial exposure during early life has persistent effects on natural killer T Cell function--Torsten Olszak, Dingding An, Sebastian Zeissig, Miguel Penilla Vera, Julia Richter, Andre Franke, Jonathan N. Glickman, Reiner Siebert, Rebecca M. Barron, Dennis L. Kasper, and Richard S. Blumberg""</p> <p>""A5 The application of ecological theory toward an understanding of the human microbiome--Elizabeth K. Costello, Keaton Stagaman, Les Dethlefsen, Brendan J. M. Bohannan, and David A. Relman""</p> <p>""A11 Community ecology and the vaginal microbiome--Larry J. Forney and Jacques Ravel""</p> <p>""A12 Investigating bacterial-animal symbioses with</p>

light sheet microscopy--Michael J. Taormina, Matthew Jemielita, W. Zac Stephens, Adam R. Burns, Joshua V. Troll, Raghuvier Parthasarathy, and Karen Guillemin"; ""A13 Clinical application of fecal microbiota transplantation in Clostridium difficile infection and beyond--Josbert J. Keller and Els van Nood""

""A14 Consumption of human milk glycoconjugates by infant-associated bifidobacteria: Mechanisms and implications--Daniel Garrido, David C. Dallas, and David A. Mills""""A15 Bacteriophage adhering to mucus provide a non host-derived immunity--Jeremy J. Barr, Rita Auro, Mike Furlan, Katrine L. Whiteson, Marcella L. Erb, Joe Pogliano, Aleksandr Stotland, Roland Wolkowicz, Andrew S. Cutting, Kelly S. Doran, Peter Salamon, Merry Youle, and Forest Rohwer""

""A16 Topographic diversity of fungal and bacterial communities in human skin--Keisha Findley, Julia Oh, Joy Yang, Sean Conlan, Clayton Deming, Jennifer A. Meyer, Deborah Schoenfeld, Effie Nomicos, Morgan Park, NIH Intramural Sequencing Center Comparative Sequencing Program, Heidi H. Kong, and Julia A. Segre""

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