

Kathryn DeRiemer, M.P.H., M.S., Ph.D.

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| Clinical Interests | Dr. DeRiemer's research interests include: infectious disease epidemiology, airborne transmission of pathogens, antibiotic resistance, diversity of pathogen strain genotypes and phenotypes. |
| Title | Associate Professor |
| Specialty | Infectious Diseases, Public Health Sciences-Division of Epidemiology |
| Department | Public Health Sciences |
| Division | Epidemiology |
| Languages | Portuguese, Spanish |
| Education | M.P.H., UC Berkeley, Berkeley, California, 1994 Ph.D., UC Berkeley, Berkeley, California, 1999 M.S., University of Washington, Seattle, Washington, 1976 B.S., Stanford University, Stanford, California, 1994 |
| Fellowships | Postdoctoral Fellow, Stanford University Medical Center, Stanford, CA, 2002-05 |
| Professional Memberships | AAAS Stop TB Diagnostics Working Group The Collaborative Group for Meta-Analysis of Individual Patient Data in MDR-TB |
| Select Recent Publications | Migliori GB, Sotgiu G, Gandhi NR, Falzon D, DeRiemer K, Centis R, Hollm-Delgado MG, Palmero D, P?rez-Guzm?n C, Vargas MH, D?Ambrosio L, Spanevello A, Bauer M, Chan ED, Schaaf HS, Keshavjee S, Holtz TH, Menzies D, for the The Collaborative Group for Meta-Analysis of Individual Patient Data in MDR-TB. Drug resistance beyond XDR-TB: results from a large individual patient data meta-analysis. European Respiratory Journal. Epub 11 Oct 2012 Stopka TJ, Lutnick A, Wenger LD, DeRiemer K, Geraghty EM, Kral AH. Demographic, risk and spatial factors associated with over-the-counter syringe purchase among injection drug users. Amer J Epidemiol 2012; 176(1):14-23. The Collaborative Group for Meta-Analysis of Individual Patient Data in MDR-TB: Members (in alphabetic order): Ahuja S, Ashkin D, Avendano M, Banerjee R, Bauer M, Bayona JN, Becerra MC, Benedetti A, Burgos M, Centis R, Chan ED, Chiang C-Y, Cox H, D?Ambrosio L, DeRiemer K, Dung NH, Enarson D, Falzon D, Flanagan K, Flood J, Garcia-Garcia L, Gandhi N, Granich RM, Hollm-Delgado MG, Holtz TH, Iseman MD, Jarlsberg LG, Keshavjee S, Kim HR, Koh WJ, Lancaster J, |

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Abernethy NF, DeRiemer K, Small PM. Methods to Identify Standard Data Elements in Clinical and Public Health Forms. *American Medical Informatics Association (AMIA) Ann Symp Proc* 2011; 19: 27.

de Jong BC, Antonio M, Awine T, Ogungbemi K, de Jong YP, Gagneux S, DeRiemer K, Zozio T, Rastogi N, Borgdorff M, Hill PC, Adegbola RA. Use of spoligotyping and large-sequence polymorphisms to study the population structure of the Mycobacterium tuberculosis complex in a cohort study of consecutive smear-positive tuberculosis cases in the Gambia. *J Clin Microbiol* 2009; 47(4):994-1001.

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