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Genome Sequencing of *Ralstonia solanacearum* Biovar 3, Phylotype I, Strains Rs-09-161 and Rs-10-244, Isolated from Eggplant and Chili in India

Raman Ramesh, Sapna Gaitonde, Gauri Achari, Trupti Asolkar, Narendra Pratap Singh, Sebastien Carrere, Stephane Genin, and Nemo Peeters

Genome Announc. May/June 2014 2:e00323-14; doi:10.1128/genomeA.00323-14

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Draft Whole-Genome Sequence of a New Variant of *Vibrio cholerae* O1 El Tor Strain Isolated from a Cholera Patient in Russia

Nina I. Smirnova, Alexander V. Cherkasov, Yaroslav M. Krasnov, Dmitriy A. Agafonov, and Vladimir V. Kuttyrev

Genome Announc. May/June 2014 2:e00432-14; doi:10.1128/genomeA.00432-14

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Draft Genome Sequence of *bla*_{NDM-1}-Positive *Escherichia coli* O25b-ST131 Clone Isolated from an Environmental Sample

Kirthi K. Kutumbaka, Sukkyun Han, James Mategko, Cesar Nadala, Genevieve L. Buser, Maureen P. Cassidy, Zintars G. Beldavs, Scott J. Weissman, Karim E. Morey, Robert Vega, and Mansour Samadpour

Genome Announc. May/June 2014 2:e00462-14; doi:10.1128/genomeA.00462-14

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Draft Genome Sequence of *Lactobacillus animalis* 381-IL-28

Joseph M. Sturino, Mahitha Rajendran, and Eric Altermann

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Draft Genome Sequence of *Raoultella ornithinolytica* TNT, a Trinitrotoluene-Denitrating and Plant Growth-Promoting Strain Isolated from Explosive-Contaminated Soil

Sofie Thijs, Jonathan Van Hamme, Panagiotis Gkorezis, Francois Rineau, Nele Weyens, and Jaco Vangronsveld

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Genome Sequence of *Bacillus cereus* Strain A1, an Efficient Starch-Utilizing Producer of Hydrogen

Ting Zhang, Meidan Bao, Yu Wang, Haijia Su, and Tianwei Tan

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Draft Genome Sequence of *Mycobacterium triplex* DSM 44626

Mohamed Sassi, Olivier Croce, Catherine Robert, Didier Raoult, and Michel Drancourt

Genome Announc. May/June 2014 2:e00499-14; doi:10.1128/genomeA.00499-14

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Complete Genome Sequence of the Subsurface, Mesophilic Sulfate-Reducing Bacterium *Desulfovibrio aespoensis* Aspo-2

Karsten Pedersen, Andreas Bengtsson, Johanna Edlund, Lisa Rabe, Terry Hazen, Romy Chakraborty, Lynne Goodwin, and Nicole Shapiro

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Complete Genome Sequence of the Attenuated Novobiocin-Resistant *Streptococcus iniae* Vaccine Strain ISNO

Julia W. Pridgeon, Dunhua Zhang, and Lee Zhang

Genome Announc. May/June 2014 2:e00510-14; doi:10.1128/genomeA.00510-14

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Draft Genome Sequence of *Bacillus thuringiensis* Strain LM1212, Isolated from the Cadaver of an *Oryctes gigas* Larva in Madagascar

Guiming Liu, Chao Deng, Lai Song, Qi Peng, Jie Zhang, Didier Lereclus, and Fuping Song

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Draft Genome Sequence of *Frankia* sp. Strain BMG5.23, a Salt-Tolerant Nitrogen-Fixing Actinobacterium Isolated from the Root Nodules of *Casuarina glauca* Grown in Tunisia

Faten Ghodhbane-Gtari, Sheldon G. Hurst IV, Rediet Oshone, Krystalynne Morris, Feseha Abebe-Akele, W. Kelley Thomas, Amir Ktari, Karima Salem, Maher Gtari, and Louis S. Tisa

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Draft Genome Sequence of *Mycobacterium farcinogenes* NCTC 10955

Olivier Croce, Catherine Robert, Didier Raoult, and Michel Drancourt

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Complete Genome Sequences of *Pseudomonas monteilii* SB3078 and SB3101, Two Benzene-, Toluene-, and Ethylbenzene-Degrading Bacteria Used for Bioaugmentation

Morten S. Dueholm, Mads Albertsen, Seth D'Imperio, Vaibhav P. Tale, Derrick Lewis, Per Halkjær Nielsen, and Jeppe Lund Nielsen

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Complete Genome of *Rhodococcus pyridinivorans* SB3094, a Methyl-Ethyl-Ketone-Degrading Bacterium Used for Bioaugmentation

Morten S. Dueholm, Mads Albertsen, Seth D'Imperio, Vaibhav P. Tale, Derrick Lewis, Per Halkjær Nielsen, and Jeppe Lund Nielsen

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Draft Genome Sequence of *Enterobacter cloacae* Strain JD6301

Jessica G. Wilson, William T. French, Anna Lipzen, Joel Martin, Wendy Schackwitz, Tanja Woyke, Nicole Shapiro, James W. Bullard, Franklin R. Champlin, and Janet R. Donaldson

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Genome Sequence of *Sporolactobacillus terrae* DSM 11697, the Type Strain of the Species

Kaiming Huang, Jun Ni, Ke Xu, Hongzhi Tang, Fei Tao, and Ping Xu

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Draft Genome Sequence of *Bacillus cereus* Strain BcFL2013, a Clinical Isolate Similar to G9241

Jay E. Gee, Chung K. Marston, Scott A. Sammons, Mark A. Burroughs, and Alex R. Hoffmaster

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Complete Genome Sequence of *Winogradskyella* sp. Strain PG-2, a Proteorhodopsin-Containing Marine Flavobacterium

Yohei Kumagai, Susumu Yoshizawa, Kenshiro Oshima, Masahira Hattori, Wataru Iwasaki, and Kazuhiro Kogure

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Draft Genome Sequence of the Field Isolate *Brucella melitensis* Strain Bm IND1 from India

Sashi Bhushan Rao, Vivek K. Gupta, Mukesh Kumar, Nagendra R. Hegde, Gary A. Splitter, Pallu Reddanna, and Girish K. Radhakrishnan

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Comparative Genomic Analysis of Two Multidrug-Resistant Clinical Isolates of ST395 Epidemic Strain of *Pseudomonas aeruginosa* Obtained 12 Years Apart

Benoit Valot, Laurence Rohmer, Michael A. Jacobs, Samuel I. Miller, Xavier Bertrand, and Didier Hocquet

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Draft Genome Sequence of Methicillin-Resistant *Staphylococcus epidermidis* Strain ET-024, Isolated from an Endotracheal Tube Biofilm of a Mechanically Ventilated Patient

Ilse Vandecandelaere, Filip Van Nieuwerburgh, Dieter Deforce, Hans J. Nelis, and Tom Coenye

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Metagenomic Approach for Understanding Microbial Population from Petroleum Muck

M. N. Joshi, S. V. Dhebar, S. V. Dhebar, P. Bhargava, A. S. Pandit, R. P. Patel, A. K. Saxena, and S. B. Bagatharia

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Complete Genome Sequences of *Salmonella enterica* Serovar Heidelberg Strains Associated with a Multistate Food-Borne Illness Investigation

Peter S. Evans, Yan Luo, Tim Muruvanda, Sherry Ayers, Brian Hiatt, Maria Hoffman, Shaohua Zhao, Marc W. Allard, and Eric W. Brown

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Draft Genome Sequence of *Xanthomonas arboricola* pv. pruni Strain Xap33, Causal Agent of Bacterial Spot Disease on Almond

J. Garita-Cambronero, M. Sena-Vélez, A. Palacio-Bielsa, and J. Cubero

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Draft Genome Sequence of *Deffluviimonas indica* Strain 20V17^T, Isolated from a Deep-Sea Hydrothermal Vent Environment in the Southwest Indian Ocean

Lijing Jiang姜 丽晶, Minnan Long龙 敏南, and Zongze Shao邵 宗泽

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Genome Sequences of Two *Leuconostoc pseudomesenteroides* Strains Isolated from Danish Dairy Starter Cultures

T. B. Pedersen, W. P. Kot, L. H. Hansen, S. J. Sørensen, J. R. Broadbent, F. K. Vogensen, and Y. Ardo

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Genome Sequence of *Leuconostoc mesenteroides* subsp. *cremoris* Strain T26, Isolated from Mesophilic Undefined Cheese Starter

T. B. Pedersen, W. P. Kot, L. H. Hansen, S. J. Sørensen, J. R. Broadbent, F. K. Vogensen, and Y. Ardo

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Whole-Genome Sequences of Low-Virulence Strain CB3 and Mild Strain CB7 of *Chlamydia psittaci*

Jun Chu, Ruixue Sun, Zongxue Wu, Shanshan Liu, Dongfang Li, Qiang Zhang, Yong Ling, Yanping Gong, Renhua Wu, Honglong Wu, Jizhang Zhou, Cheng He, and Peixiang Ni

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Draft Genome Sequence of *Pseudomonas* sp. Strain Ant30-3, a Psychrotolerant Bacterium with Biodegradative Attribute Isolated from Antarctica

Hyunmin Koo, Malay K. Basu, Michael Crowley, Jackie Aislabie, and Asim K. Bej

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Draft Genome Sequence of the *Mycobacterium tuberculosis* Clinical Isolate C2, Belonging to the Latin American–Mediterranean Family

Yu-Chieh Liao, Yih-Yuan Chen, Hsin-Hung Lin, Jia-Ru Chang, Ih-Jen Su, Tsi-Shu Huang, and Horng-Yunn Dou

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Genome Sequence of a Promising Hydrogen-Producing Facultative Anaerobic Bacterium, *Brevundimonas naejangsensis* Strain B1

Haijia Su, Ting Zhang, Meidan Bao, Yixin Jiang, Yu Wang, and Tianwei Tan

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Complete Genome Sequence of the Highly Transformable *Pseudomonas stutzeri* Strain 28a24

Brian A. Smith, Kevin M. Dougherty, and David A. Baltrus

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Draft Genome Sequence of *Acetobacter aceti* Strain 1023, a Vinegar Factory Isolate

John E. Hung, Christopher P. Mill, Sandra W. Clifton, Vincent Magrini, Ketaki Bhide, Julie A. Francois, Aaron E. Ransome, Lucinda Fulton, Jyothi Thimmapuram, Richard K. Wilson, and T. Joseph Kappock

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Genome Sequences of Seven *Mycoplasma hyosynoviae* Strains Isolated from the Joint Tissue of Infected Swine (*Sus scrofa*)

Eric Bumgardner, Russell F. Bey, Weerayuth Kittichotirat, Roger E. Bumgarner, and Paulraj K. Lawrence

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Complete Genome Sequence of a Virulent Strain, *Streptococcus iniae* ISET0901, Isolated from Diseased Tilapia

Julia W. Pridgeon, Dunhua Zhang, and Lee Zhang

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Complete Genome Sequence of a Moderately Virulent *Aeromonas hydrophila* Strain, pc104A, Isolated from Soil of a Catfish Pond in West Alabama

Julia W. Pridgeon, Dunhua Zhang, and Lee Zhang

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Draft Genome Sequence of *Geobacillus thermopakistanensis* Strain MAS1

Masood Ahmed Siddiqui, Naeem Rashid, Saravanaraj Ayyampalayam, and William B. Whitman

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Draft Genome Sequences of *Geobacillus* sp. Strains CAMR5420 and CAMR12739

Pieter De Maayer, Carolyn E. Williamson, Christopher T. Vennard, Michael J. Danson, and Don A. Cowan

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***Yersinia pseudotuberculosis* ST42 (O:1) Strain Misidentified as *Yersinia pestis* by Mass Spectrometry Analysis**

Patrick Jérôme, Philippe Le Flèche, Yann Blouin, Holger C. Scholz, François M. Thibault, Françoise Raynaud, Gilles Vergnaud, and Christine Pourcel

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Draft Genome Sequences of Two Clinical Isolates of *Streptococcus mutans*

Hui Zheng, Lihong Guo, Ning Du, Jiuxiang Lin, Lai Song, Guiming Liu, and Feng Chen

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Draft Genome Sequence of Insecticidal *Streptomyces* sp. Strain PCS3-D2, Isolated from Mangrove Soil in Philippines

Aileen N. Bayot-Custodio, Edwin P. Alcantara, and Teofila O. Zulaybar

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Genome Sequence of the ϵ -Poly-L-Lysine-Producing Strain *Streptomyces albulus* NK660, Isolated from Soil in Gutian, Fujian Province, China

Yanyan Gu, Chao Yang, Xiaomeng Wang, Weitao Geng, Yang Sun, Jun Feng, Yuanyuan Wang, Yufen Quan, You Che, Chi Zhang, Ting Gong, Wei Zhang, Weixia Gao, Zhenqiang Zuo, Cunjiang Song, and Shufang Wang

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Genome Sequence of *Thermoanaerobaculum aquaticum* MP-01^T, the First Cultivated Member of *Acidobacteria* Subdivision 23, Isolated from a Hot Spring

Blake W. Stamps, Nathaniel A. Losey, Paul A. Lawson, and Bradley S. Stevenson

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Complete Genome Sequence of Amino Acid-Utilizing *Eubacterium acidaminophilum* al-2 (DSM 3953)

Anja Poehlein, Jan R. Andreessen, and Rolf Daniel

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Draft Genome Sequence of *Photobacterium halotolerans* S2753, Producer of Bioactive Secondary Metabolites

Henrique Machado, Maria Månsson, and Lone Gram

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Draft Genome Sequences of Two Genetic Variant Strains of *Edwardsiella piscicida*, JF1305 and RSB1309, Isolated from Olive Flounder (*Paralichthys olivaceus*) and Red Sea Bream (*Pagrus major*) Cultured in Japan, Respectively

Kazuki Oguro, Kazuki Tamura, Jin Yamane, Masato Shimizu, Takeshi Yamamoto, Takuya Ikawa, Kouhei Ohnishi, Syun-ichirou Oshima, and Masayuki Imajoh

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Draft Genome Sequence of *Stenotrophomonas maltophilia* Strain M30, Isolated from a Chronic Pressure Ulcer in an Elderly Patient

Pol Huedo, Óscar Conchillo-Solé, Daniel Yero, Sònia Martínez-Servat, Xavier Daura, and Isidre Gibert

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Genome Sequence of *Aeromonas taiwanensis* LMG 24683^T, a Clinical Wound Isolate from Taiwan

Hsuan-Chen Wang, Wen-Chien Ko, Hung-Yu Shu, Po-Lin Chen, Yu-Chun Wang, and Chi-Jung Wu

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Draft Genome Sequences of Nine Enteropathogenic *Escherichia coli* Strains from Kenya

Tracy H. Hazen, Michael S. Humphrys, John Benjamin Ochieng, Michele Parsons, Cheryl A. Bopp, Ciara E. O'Reilly, Eric Mintz, and David A. Rasko

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Genome Sequences of Strain ATCC 29281 and Pin and Northern Red Oak Isolates of *Lonsdalea quercina* subsp. *quercina*

Jorge Ibarra Caballero, Marcelo M. Zerillo, Jacob Snelling, Whitney Cranshaw, Christina Boucher, and Ned Tisserat

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Genome Sequence of SCB34, a Sequence Type 131 Multidrug-Resistant *Escherichia coli* Isolate Causing Neonatal Early-Onset Sepsis

Susana Chavez-Bueno, Michael W. Day, Inimary T. Toby, Darrin R. Akins, and David W. Dyer

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Draft Genome Sequence of *Aeromonas hydrophila* Strain Ae34, Isolated from a Septicemic and Moribund Koi Carp (*Cyprinus carpio koi*), a Freshwater Aquarium Fish

S. S. De S. Jagoda, Engkong Tan, Appudurai Arulkanthan, Shigeharu Kinoshita, Shugo Watabe, and Shuichi Asakawa

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Draft Genome Sequence of *Bacillus pumilus* 7P, Isolated from the Soil of the Tatarstan Republic, Russia

Elena I. Shagimardanova, Anna A. Toymentseva, Nelly P. Balaban, Ayslu M. Mardanova, Yulia V. Danilova, Oleg A. Gusev, Elena Kostryukova, Irina Karpova, Aleksandr Manolov, Dmitriy Alexeev, and Margarita R. Sharipova

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Genome Sequence of *Arthrobacter siccitolerans* 4J27, a Xeroprotectant-Producing Desiccation-Tolerant Microorganism

M. Manzanera, L. Santa-Cruz-Calvo, J. I. Vilchez, C. García-Fontana, G. A. Silva-Castro, C. Calvo, and J. González-López

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Whole-Genome Sequence of *Burkholderia* sp. Strain RPE67, a Bacterial Gut Symbiont of the Bean Bug *Riptortus pedestris*

Kazutaka Takeshita, Tomoko F. Shibata, Naruo Nikoh, Tomoaki Nishiyama, Mitsuyasu Hasebe, Takema Fukatsu, Shuji Shigenobu, and Yoshitomo Kikuchi

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Genome Sequence of *meso*-2,3-Butanediol-Producing Strain *Serratia marcescens* ATCC 14041

Lixiang Li, Yu Wang, Kun Li, Fei Su, Cuiqing Ma, and Ping Xu

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Complete Genome Sequence of the *p*-Nitrophenol-Degrading Bacterium *Pseudomonas putida* DLL-E4

Xiaojun Hu, Jue Wang, Fei Wang, Qiongzheng Chen, Yan Huang, and Zhongli Cui

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Genome Sequences of Nine *Bordetella holmesii* Strains Isolated in the United States

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Draft Genome Sequence of “*Candidatus Cronobacter colletis*” NCTC 14934^T, a New Species in the Genus *Cronobacter*

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Draft Genome Sequence of a *Legionella pneumophila* Serogroup 4 Strain Causing Legionellosis

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Draft Genome Sequence of *Kitasatospora cheerisanensis* KCTC 2395, Which Produces Plecomacrolide against Phytopathogenic Fungi

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Draft Genome Sequence of *Bacillus subtilis* strain KATMIRA1933

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Draft Genome Sequences of *Geomicrobium* sp. Strains JCM 19037, JCM 19038, JCM 19039, and JCM 19055, Isolated from Aquatic Samples

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Draft Genome Sequences of *Vibrio* sp. Strains Isolated from Tetrodotoxin-Bearing Scavenging Gastropod

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Draft Genome Sequence of the Lignin-Degrading *Burkholderia* sp. Strain LIG30, Isolated from Wet Tropical Forest Soil

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Genome Sequence of *Pseudomonas* sp. Strain P482, a Tomato Rhizosphere Isolate with Broad-Spectrum Antimicrobial Activity

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Genome Sequence of *Streptomyces olindensis* DAUFPE 5622, Producer of the Antitumoral Anthracycline Cosmomycin D

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Draft Genome Sequence of Enterotoxigenic *Escherichia coli* Strain W25K

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Draft Genome Sequence of *Kozakia baliensis* SR-745, the First Sequenced *Kozakia* Strain from the Family *Acetobacteraceae*

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Complete Genome Sequence of a Human Enterovirus 71 Strain Isolated from a Fatal Case in Shanghai, China, in 2012

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Complete Genome Sequences of Three Novel Human Papillomavirus Types, 175, 178, and 180

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Complete Genome Sequences of Nine Mycobacteriophages

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Complete Genome Sequence of K14JB01, a Novel Variant Strain of Porcine Epidemic Diarrhea Virus in South Korea

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Complete Genome Sequence of Canine Papillomavirus Type 11

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Genome Sequence of the Novel Reassortant Mammalian Orthoreovirus Strain MRV00304/13, Isolated from a Calf with Diarrhea from the United States

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Complete Genome Sequence of Foot-and-Mouth Disease Virus Type A Circulating in Bangladesh

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Ran Liu, Guilin Zhang, Yinhui Yang, Rongli Dang, and Tongyan Zhao

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