

CONTENTS • JANUARY/FEBRUARY 2013 • VOLUME 1, NO. 1

EUKARYOTES

- Draft Genome Sequence of *Herpotrichiellaceae* sp. UM 238 Isolated from Human Skin Scraping e00148-12
Kee Peng Ng, Su Mei Yew, Chai Ling Chan, Ruixin Tan, Tuck Soon Soo-Hoo, Shiang Ling Na, Hamimah Hassan, Yun Fong Ngeow, Chee-Choong Hoh, Kok Wei Lee, Wai-Yan Yee
- Draft Genome Sequence of *Fusarium fujikuroi* B14, the Causal Agent of the Bakanae Disease of Rice e00035-13
Haeyoung Jeong, Seunghoon Lee, Gyung Ja Choi, Theresa Lee, Sung-Hwan Yun

PROKARYOTES

- Complete Genome Sequence of a *Propionibacterium acnes* Isolate from a Sarcoidosis Patient e00016-12
Kana Minegishi, Chihiro Aikawa, Asuka Furukawa, Takayasu Watanabe, Tsubasa Nakano, Yoshitoshi Ogura, Yoshiyuki Ohtsubo, Ken Kurokawa, Tetsuya Hayashi, Fumito Maruyama, Ichiro Nakagawa, Yoshinobu Eishi
- Genome Sequence of *Sinorhizobium meliloti* Rm41 e00013-12
Stefan Weidner, Birgit Baumgarth, Michael Göttfert, Sebastian Jaenicke, Alfred Pühler, Susanne Schneiker-Bekel, Javier Serrania, Rafael Szczepanowski, Anke Becker
- Genome Sequences of *Salmonella enterica* Serovar Heidelberg Isolates Isolated in the United States from a Multistate Outbreak of Human *Salmonella* Infections e00004-12
Maria Hoffmann, Yan Luo, Patricia C. Lafon, Ruth Timme, Marc W. Allard, Patrick F. McDermott, Eric W. Brown, Shaohua Zhao
- Genome Sequences of Two *Lactococcus garvieae* Strains Isolated from Meat e00018-12
Giovanni Ricci, Chiara Ferrario, Francesca Borgo, Giovanni Eraclio, Maria Grazia Fortina
- Genome Sequence of *Helicobacter heilmannii* Sensu Stricto ASB1 Isolated from the Gastric Mucosa of a Kitten with Severe Gastritis e00033-12
Annemieke Smet, Filip Van Nieuwerburgh, Jessica Ledesma, Bram Flahou, Dieter Deforce, Richard Ducatelle, Freddy Haesebrouck
- Complete Genome Sequence of *Simiduia agarivorans* SA1^T, a Marine Bacterium Able To Degrade a Variety of Polysaccharides e00039-12
Silk Yu Lin, Wung Yang Shieh, Jwo-Sheng Chen, Sen-Lin Tang
- Genome Sequence of *Sphingomonas xenophaga* QYY, an Anthraquinone-Degrading Strain e00031-12
Yuanyuan Qu, Xuwang Zhang, Hao Yu, Hongzhi Tang, E Shen, Hao Zhou, Qiao Ma, Xiangyu Cao, Jiti Zhou, Ping Xu
- Draft Genome Sequence of the *Paenibacillus* sp. Strain ICGEB2008 (MTCC 5639) Isolated from the Gut of *Helicoverpa armigera* e00026-12
Nidhi Adlakha, Hemant Ritturaj Kushwaha, Raman Rajagopal, Syed Shams Yazdani

Draft Genome Sequence of a Single Cell of SAR86 Clade Subgroup IIIa	e00030-12
<i>D. B. Rusch, M.-J. Lombardo, J. Yee-Greenbaum, M. Novotny, L. M. Brinkac, R. S. Lasken, C. L. Dupont</i>	
Draft Genome of <i>Klebsiella pneumoniae</i> Sequence Type 512, a Multidrug-Resistant Strain Isolated during a Recent KPC Outbreak in Italy	e00035-12
<i>Francesco Comandatore, Paolo Gaibani, Simone Ambretti, Maria Paola Landini, Daniele Daffonchio, Piero Marone, Vittorio Sambri, Claudio Bandi, Davide Sassera</i>	
Genome Sequence of the Acidophilic Bacterium <i>Acidocella</i> sp. Strain MX-AZ02	e00041-12
<i>Luis E. Servín-Garcidueñas, Roger A. Garrett, Ricardo Amils, Esperanza Martínez-Romero</i>	
Whole-Genome Shotgun Sequencing of <i>Mycobacterium abscessus</i> M156, an Emerging Clinical Pathogen in Malaysia	e00063-12
<i>Siew Woh Choo, Yan Ling Wong, Ching Yew Beh, Naline Lokanathan, Mee Lian Leong, Chia Sui Ong, Kee Peng Ng, Yun Fong Ngeow</i>	
Genome Sequence of <i>Rhizobium lupini</i> HPC(L) Isolated from Saline Desert Soil, Kutch (Gujarat)	e00071-12
<i>Leena Agarwal, Hemant J. Purohit</i>	
Genome Sequence of <i>Pseudomonas</i> sp. Strain Chol1, a Model Organism for the Degradation of Bile Salts and Other Steroid Compounds	e00014-12
<i>Johannes Holert, Intikhab Alam, Michael Larsen, André Antunes, Vladimir B. Bajic, Ulrich Stingl, Bodo Philipp</i>	
Genome Sequence of the Attenuated Carbosap Vaccine Strain of <i>Bacillus anthracis</i>	e00067-12
<i>Robin Harrington, Brian D. Ondov, Diana Radune, Mary Beth Friss, Joy Klubnik, Lynn Diviak, Jonathan Hnath, Stephen R. Cendrowski, Thomas E. Blank, David Karaolis, Arthur M. Friedlander, James P. Burans, M. J. Rosovitz, Todd Treangen, Adam M. Phillippe, Nicholas H. Bergman</i>	
Genome Sequence of <i>Chlamydia psittaci</i> Strain 01DC12 Originating from Swine	e00078-12
<i>Helena M. B. Seth-Smith, Michelle Sait, Konrad Sachse, Wolfgang Gaede, David Longbottom, Nicholas R. Thomson</i>	
Draft Genome Sequences of Two Clinical Isolates of <i>Lactobacillus rhamnosus</i> from Initial Stages of Dental Pulp Infection	e00073-12
<i>Zhiliang Chen, Marc R. Wilkins, Neil Hunter, Mangala A. Nadkarni</i>	
Draft Genome Sequence of Fructophilic <i>Lactobacillus florum</i>	e00025-12
<i>Eun Bae Kim, Charlotte A. Tyler, Lauren M. Kopit, Maria L. Marco</i>	
Draft Genome Sequence of the Opportunistic Human Pathogen <i>Morganella morganii</i> SC01	e00051-12
<i>Indu Khatri, Chetna Dureja, Saumya Raychaudhuri, Srikrishna Subramanian</i>	
Draft Genome Sequence of <i>Rhizobium mesoamericanum</i> STM3625, a Nitrogen-Fixing Symbiont of <i>Mimosa pudica</i> Isolated in French Guiana (South America)	e00066-12
<i>Lionel Moulin, Damien Mornico, Rémy Melkonian, Agnieszka Klonowska</i>	
Draft Genome Sequences of <i>Pseudomonas fluorescens</i> BS2 and <i>Pusillimonas noertemannii</i> BS8, Soil Bacteria That Cooperate To Degrade the Poly- γ -D-Glutamic Acid Anthrax Capsule	e00057-12
<i>Richard A. Stabler, David Negus, Arnab Pain, Peter W. Taylor</i>	

Draft Genome Sequence of <i>Clostridium</i> sp. Maddingley, Isolated from Coal-Seam Gas Formation Water	e00081-12
<i>Carly P. Rosewarne, Paul Greenfield, Dongmei Li, Nai Tran-Dinh, Mark I. Bradbury, David J. Midgley, Philip Hendry</i>	
Draft Genome Sequence of <i>Methanobacterium</i> sp. Maddingley, Reconstructed from Metagenomic Sequencing of a Methanogenic Microbial Consortium Enriched from Coal-Seam Gas Formation Water	e00082-12
<i>Carly P. Rosewarne, Paul Greenfield, Dongmei Li, Nai Tran-Dinh, David J. Midgley, Philip Hendry</i>	
Draft Genome Sequences and Annotation of <i>Enterococcus faecium</i> Strain LCT-EF20	e00083-12
<i>De Chang, Yuanfang Zhu, Jiapeng Chen, Xiangqun Fang, Tianzhi Li, Junfeng Wang, Yinghua Guo, Longxiang Su, Guogang Xu, Yajuan Wang, Zhenhong Chen, Changting Liu</i>	
Draft Genome Sequences of Two Multidrug Resistant <i>Klebsiella pneumoniae</i> ST258 Isolates Resistant to Colistin	e00113-12
<i>Francesco Comandatore, Davide Sassera, Simone Ambretti, Maria Paola Landini, Daniele Daffonchio, Piero Marone, Vittorio Sambri, Claudio Bandi, Paolo Gaibani</i>	
Genome Sequence of the Sulfate-Reducing Bacterium <i>Desulfotomaculum hydrothermale</i> Lam5 ^T	e00114-12
<i>Oulfat Amin, Marie-Laure Fardeau, Odile Valette, Agnès Hirschler-Réa, Valérie Barbe, Claudine Médigue, Benoit Vacherie, Bernard Ollivier, Philippe N. Bertin, Alain Dolla</i>	
Whole-Genome Shotgun Sequence of <i>Pseudomonas viridisflava</i> , a Bacterium Species Pathogenic to <i>Arabidopsis thaliana</i>	e00116-12
<i>Francois Lefort, Gautier Calmin, Julien Crovadore, Magne Osteras, Laurent Farinelli</i>	
Whole Genome Sequencing of <i>Thermus oshimai</i> JL-2 and <i>Thermus thermophilus</i> JL-18, Incomplete Denitrifiers from the United States Great Basin	e00106-12
<i>Senthil K. Murugapiran, Marcel Huntemann, Chia-Lin Wei, James Han, John C. Detter, Cliff S. Han, Tracy H. Erkkila, Hazuki Teshima, Amy Chen, Nikos Kyrpides, Konstantinos Mavrommatis, Victor Markowitz, Ernest Szeto, Natalia Ivanova, Ioanna Pagani, Jenny Lam, Austin I. McDonald, Jeremy A. Dodsworth, Amrita Pati, Lynne Goodwin, Lin Peters, Sam Pitluck, Tanja Woyke, Brian P. Hedlund</i>	
Genome Sequence of <i>Klebsiella pneumoniae</i> Ecl8, a Reference Strain for Targeted Genetic Manipulation	e00027-12
<i>Maria Fookes, Jing Yu, Shyamasree De Majumdar, Nicholas Thomson, Thamarai Schneiders</i>	
Draft Genome Sequence of the First Isolate of Extensively Drug-Resistant (XDR) <i>Mycobacterium tuberculosis</i> in Malaysia	e00056-12
<i>Kee Peng Ng, Su Mei Yew, Chai Ling Chan, Jennifer Chong, Soo Nee Tang, Tuck Soon Soo-Hoo, Shiang Ling Na, Hamimah Hassan, Yun Fong Ngeow, Chee Choong Hoh, Kok Wei Lee, Wai Yan Yee</i>	
Draft Genome Sequences of Five Strains in the Genus <i>Thauera</i>	e00052-12
<i>Binbin Liu, Åsa Frostegård, James P. Shapleigh</i>	
Draft Genome Sequences of Two Virulent Serotypes of Avian <i>Pasteurella multocida</i>	e00058-12
<i>Juan E. Abrahante, Timothy J. Johnson, Samuel S. Hunter, Samuel K. Maheswaran, Melissa J. Hauglund, Darrell O. Bayles, Fred M. Tatum, Robert E. Briggs</i>	

Draft Genome Sequence of a Subarctic Humic Substance-Degrading Pseudomonad <i>Ha Ju Park, Seung Chul Shin, Dockyu Kim</i>	e00070-12
Genome Sequence of <i>Klebsiella oxytoca</i> M5al, a Promising Strain for Nitrogen Fixation and Chemical Production <i>Guanhui Bao, Yanping Zhang, Chenyu Du, Zugen Chen, Yin Li, Zhu'an Cao, Yanhe Ma</i>	e00074-12
Complete Chromosome Sequence of <i>Carnobacterium maltaromaticum</i> LMA 28 <i>Catherine Cailliez-Grimal, Stéphane Chaillou, Jamila Anba-Mondoloni, Valentin Loux, Muhammad Inam Afzal, Abdur Rahman, Gilles Kergourlay, Marie-Christine Champomier-Vergès, Monique Zagorec, Paw Dalgaard, Jorgen J. Leisner, Hervé Prévost, Anne-Marie Revol-Junelles, Frédéric Borges</i>	e00115-12
Draft Genome Sequences of Four Nosocomial Methicillin-Resistant <i>Staphylococcus aureus</i> (MRSA) Strains (PPUKM-261-2009, PPUKM-332-2009, PPUKM-377-2009, and PPUKM-775-2009) Representative of Dominant MRSA Pulsotypes Circulating in a Malaysian University Teaching Hospital <i>Hui-Min Neoh, Zeti-Azura Mohamed-Hussein, Xin-Ee Tan, Raja Mohd Fadhil B Raja Abd Rahman, Salasawati Hussin, Noraziah Mohamad Zin, Rahman Jamal</i>	e00103-12
Complete Genome of <i>Lactococcus lactis</i> subsp. <i>cremoris</i> UC509.9, Host for a Model Lactococcal P335 Bacteriophage <i>Stuart Ainsworth, Aldert Zomer, Victor de Jager, Francesca Bottacini, Sacha A. F. T. van Hijum, Jennifer Mahony, Douwe van Sinderen</i>	e00119-12
Draft Genome Sequence of <i>Brucella abortus</i> BCB027, a Strain Isolated from a Domestic Deer <i>Lulu Wang, Yefeng Qiu, Zeliang Chen, Jie Xu, Zhoujia Wang, Yuehua Ke, Tiefeng Li, Dali Wang, Liuyu Huang, Yaqin Yu, Qing Zhen</i>	e00130-12
Genome Sequence of the Human Abscess Isolate <i>Streptococcus intermedius</i> BA1 <i>Paul J. Planet, Ryan Rampersaud, Saul R. Hymes, Susan Whittier, Phyllis A. Della-Latta, Apurva Narechania, Sean C. Daugherty, Ivette Santana-Cruz, Robert DeSalle, Jacques Ravel, Adam J. Ratner</i>	e00117-12
Draft Genome Sequence of the Methicillin-Resistant <i>Staphylococcus aureus</i> Isolate MRSA-M2 <i>Janette M. Harro, Sean Daugherty, Vincent M. Bruno, Mary Ann Jabra-Rizk, David A. Rasko, Mark E. Shirtliff</i>	e00037-12
Draft Genome of the Nitrogen-Fixing Bacterium <i>Pseudomonas stutzeri</i> Strain KOS6 Isolated from Industrial Hydrocarbon Sludge <i>Tatiana V. Grigoryeva, Aleksandr V. Laikov, Rimma P. Naumova, Aleksandr I. Manolov, Andrey K. Larin, Irina Y. Karpova, Tatiana A. Semashko, Dmitry G. Alexeev, Elena S. Kostryukova, Rudolf Müller, Vadim M. Govorun</i>	e00072-12
Draft Genome Sequence of the Type Species of the Genus <i>Citrobacter</i> , <i>Citrobacter freundii</i> MTCC 1658 <i>Shailesh Kumar, Chandandeep Kaur, Kazuyuki Kimura, Masahiro Takeo, Gajendra Pal Singh Raghava, Shanmugam Mayilraj</i>	e00120-12
Complete Genome Sequence of <i>Listeria monocytogenes</i> LL195, a Serotype 4b Strain from the 1983–1987 Listeriosis Epidemic in Switzerland <i>Thomas Weinmaier, Martin Riesing, Thomas Rattei, Jacques Bille, Carolina Arguedas-Villa, Roger Stephan, Taurai Tasara</i>	e00152-12

Complete Genome Sequence of <i>Bacillus thuringiensis</i> Strain 407 Cry-	e00158-12
<i>Anna E. Sheppard, Anja Poehlein, Philip Rosenstiel, Heiko Liesegang, Hinrich Schulenburg</i>	
Draft Genome Sequence of <i>Catellicoccus marimammalium</i> , a Novel Species Commonly Found in Gull Feces	e00019-12
<i>Michael R. Weigand, Hodon Ryu, Laura Bozek, Konstantinos T. Konstantinidis, Jorge W. Santo Domingo</i>	
Complete Genome Sequence of a <i>Francisella tularensis</i> subsp. <i>holarctica</i> Strain from Germany Causing Lethal Infection in Common Marmosets	e00135-12
<i>Markus H. Antwerpen, E. Schacht, P. Kaysser, W. D. Splettstoesser</i>	
Complete Genome Sequence of <i>emm1 Streptococcus pyogenes</i> A20, a Strain with an Intact Two-Component System, CovRS, Isolated from a Patient with Necrotizing Fasciitis	e00149-12
<i>Po-Xing Zheng, Kun-Ta Chung, Chuan Chiang-Ni, Shu-Ying Wang, Pei-Jane Tsai, Woei-Jer Chuang, Yee-Shin Lin, Ching-Chuan Liu, Jiunn-Jong Wu</i>	
Genome Sequence of <i>Bacillus licheniformis</i> CGMCC3963, a Stress-Resistant Strain Isolated in a Chinese Traditional Solid-State Liquor-Making Process	e00060-12
<i>Qun Wu, Suqin Peng, Yao Yu, Yixue Li, Yan Xu</i>	
Draft Genome Sequence of <i>Pseudomonas aeruginosa</i> Strain N002, Isolated from Crude Oil-Contaminated Soil from Geleky, Assam, India	e00104-12
<i>Abhijit Sarma Roy, Reshita Baruah, Dhrubajyoti Gogoi, Maina Borah, Anil Kumar Singh, Hari Prasanna Deka Boruah</i>	
Complete Genome Sequence of <i>Mycoplasma hyorhinis</i> Strain SK76	e00101-12
<i>Steve Goodison, Virginia Urquidi, Dibyendu Kumar, Leticia Reyes, Charles J. Rosser</i>	
Draft Genome Sequence of <i>Vibrio parahaemolyticus</i> SNUVpS-1 Isolated from Korean Seafood	e00132-12
<i>Jin Woo Jun, Ji Hyung Kim, Casiano H. Choresca, Sang Phil Shin, Jee Eun Han, Se Chang Park</i>	
Draft Genome Sequence of a Clinical Isolate, <i>Aeromonas hydrophila</i> SNUFPC-A8, from a Moribund Cherry Salmon (<i>Oncorhynchus masou masou</i>)	e00133-12
<i>Jee Eun Han, Ji Hyung Kim, Casiano Choresca, Sang Phil Shin, Jin Woo Jun, Se Chang Park</i>	
Genome Sequence of <i>Xanthomonas campestris</i> pv. <i>campestris</i> Strain Xca5	e00032-12
<i>Stéphanie Bolot, Endrick Guy, Sébastien Carrere, Valérie Barbe, Matthieu Arlat, Laurent D. Noël</i>	
Draft Genome Sequence of <i>Pediococcus lolii</i> NGRI 0510Q ^T Isolated from Ryegrass Silage	e00156-12
<i>Katsumi Doi, Kazuki Mori, Kosuke Tashiro, Yasuhiro Fujino, Yuko Nagayoshi, Yoshiharu Hayashi, Satoru Kuhara, Toshihisa Ohshima</i>	
Complete Genome Sequence of the Probiotic <i>Enterococcus faecalis</i> Symbioflor 1 Clone DSM 16431	e00165-12
<i>Moritz Fritzenwanker, Carsten Kuenne, Andre Billon, Torsten Hain, Kurt Zimmermann, Alexander Goesmann, Trinad Chakraborty, Eugen Domann</i>	
Genome Sequence of <i>Lactobacillus saerimneri</i> 30a (Formerly <i>Lactobacillus</i> sp. Strain 30a), a Reference Lactic Acid Bacterium Strain Producing Biogenic Amines	e00097-12
<i>Andrea Romano, Hein Trip, Hugo Campbell-Sills, Olivier Bouchez, David Sherman, Juke S. Lolkema, Patrick M. Lucas</i>	

Towards the Description of the Genome Catalogue of <i>Pseudomonas</i> sp. Strain M1 <i>Pedro Soares-Castro, Pedro M. Santos</i>	e00146-12
Whole Genome Sequencing and Comparative Analysis of <i>Bartonella bacilliformis</i> Strain INS, the Causative Agent of Carrion's Disease <i>D. Tarazona, C. Padilla, O. Cáceres, J. D. Montenegro, H. Bailón, G. Ventura, G. Mendoza, E. Anaya, H. Guio</i>	e00053-12
Draft Genome Sequence of a Humic Substance-Degrading <i>Paenibacillus</i> sp. Isolated from the Subarctic Grasslands at Low Temperature <i>Ha Ju Park, Dockyu Kim</i>	e00170-12
Complete Genome Sequence of the Alfalfa Symbiont <i>Sinorhizobium/Ensifer meliloti</i> Strain GR4 <i>Francisco Martínez-Abarca, Laura Martínez-Rodríguez, José Antonio López-Contreras, José Ignacio Jiménez-Zurdo, Nicolás Toro</i>	e00174-12
Genome Sequence of <i>Weissella ceti</i> NC36, an Emerging Pathogen of Farmed Rainbow Trout in the United States <i>Jason T. Ladner, Timothy J. Welch, Chris A. Whitehouse, Gustavo F. Palacios</i>	e00187-12
Genome Sequence of <i>Moraxella macacae</i> 0408225, a Novel Bacterial Species Isolated from a Cynomolgus Macaque with Epistaxis <i>Jason T. Ladner, Chris A. Whitehouse, Galina I. Koroleva, Gustavo F. Palacios</i>	e00188-12
Draft Genome Sequence of the Nitrate- and Phosphate-Accumulating <i>Bacillus</i> sp. Strain MCC0008 <i>Shreya DebRoy, Amrita Bhattacharjee, Ashoke Ranjan Thakur, Shaon Ray Chaudhuri</i>	e00189-12
Complete Genome Sequence of <i>Mycoplasma cynos</i> Strain C142 <i>Caray A. Walker, Sally A. Mannering, Shelly Shields, Damer P. Blake, Joe Brownlie</i>	e00196-12
Genome of a Gut Strain of <i>Bacillus subtilis</i> <i>Ghislain Schyns, Cláudia R. Serra, Thomas Lapointe, José B. Pereira-Leal, Sébastien Potot, Patrick Fickers, John B. Perkins, Markus Wyss, Adriano O. Henriques</i>	e00184-12
Draft Genome Sequence of an Alphaproteobacterium, <i>Caenispirillum salinarum</i> AK4 ^T , Isolated from a Solar Saltern <i>Indu Khatri, Aditya Singh, Suresh Korpole, Anil Kumar Pinnaka, Srikrishna Subramanian</i>	e00199-12
Draft Genome Sequence of a <i>Clostridium botulinum</i> Isolate from Water Used for Cooling at a Plant Producing Low-Acid Canned Foods <i>Uma Basavanna, Narjol Gonzalez-Escalona, Ruth Timme, Shomik Datta, Brianna Schoen, Eric W. Brown, Donald Zink, Shashi K. Sharma</i>	e00200-12
Genome Sequence of <i>Campylobacter showae</i> UNSWCD, Isolated from a Patient with Crohn's Disease <i>Aidan P. Tay, Nadeem O. Kaakoush, Nandan P. Deshpande, Zhiliang Chen, Hazel Mitchell, Marc R. Wilkins</i>	e00193-12
Genome of <i>Cupriavidus</i> sp. HMR-1, a Heavy Metal-Resistant Bacterium <i>Li-Guan Li, Lin Cai, Tong Zhang</i>	e00202-12

Draft Genome of <i>Spiribacter salinus</i> M19-40, an Abundant Gammaproteobacterium in Aquatic Hypersaline Environments	e00179-12
<i>Maria Jose Leon, Rohit Ghai, Ana Beatriz Fernandez, Cristina Sanchez-Porro, Francisco Rodriguez-Valera, Antonio Ventosa</i>	
Draft Genome Sequences of the <i>Enterococcus faecium</i> Strain LCT-EF258	e00147-12
<i>De Chang, Yuanfang Zhu, Xiangqun Fang, Tianzhi Li, Junfeng Wang, Yinghua Guo, Longxiang Su, Yan Liu, Xuege Jiang, Li Wang, Na Guo, Changting Liu</i>	
Draft Genome Sequence of <i>Rhodococcus opacus</i> Strain M213 Shows a Diverse Catabolic Potential	e00144-12
<i>Ashish Pathak, Stefan J. Green, Andrew Ogram, Ashvini Chauhan</i>	
Genome Sequence of Non-O1 <i>Vibrio cholerae</i> PS15	e00227-12
<i>Sanath Kumar, Ingrid E. Lindquist, Anitha Sundararajan, Chythanya Rajanna, Jared T. Floyd, Kenneth P. Smith, Jody L. Andersen, Guixin He, Ryan M. Ayers, Judith A. Johnson, James J. Werdann, Ava A. Sandoval, Nadia M. Mojica, Faye D. Schilkey, Joann Mudge, Manuel F. Varela</i>	
Draft Genome Sequence of <i>Enterococcus faecalis</i> PC1.1, a Candidate Probiotic Strain Isolated from Human Feces	e00160-12
<i>Páraig Ó Cuív, Eline S. Klaassens, Wendy J. Smith, Stanislas Mondot, A. Scott Durkin, Derek M. Harkins, Les Foster, Jamison McCorrison, Manolito Torralba, Karen E. Nelson, Mark Morrison</i>	
Draft Genome Sequence of a Clinical Strain of <i>Yersinia enterocolitica</i> (IP10393) of Bioserotype 4/O:3 from France	e00150-12
<i>Cyril Savin, Lionel Frangeul, Laurence Ma, Christiane Bouchier, Ivan Moszer, Elisabeth Carniel</i>	
Genome Sequence of <i>Klebsiella pneumoniae</i> KpQ3, a DHA-1 β -Lactamase-Producing Nosocomial Isolate	e00167-12
<i>Raquel Tobes, Francisco M. Codoñer, Elena López-Camacho, Iñigo J. Salanueva, Marina Manrique, Marta Brozynska, Rosa Gómez-Gil, Juan F. Martínez-Blanch, Miguel Álvarez-Tejado, Eduardo Pareja, Jesús Mingorance</i>	
Complete Genome Sequence of Prepandemic <i>Vibrio parahaemolyticus</i> BB22OP	e00002-12
<i>Roderick V. Jensen, Saylem M. DePasquale, Elizabeth A. Harbolick, Tian Hong, Alison L. Kornell, David H. Kruchko, Thero Modise, Cimarron E. Smith, Linda L. McCarter, Ann M. Stevens</i>	
First Genome Sequence of a Syntrophic Acetate-Oxidizing Bacterium, <i>Tepidanaerobacter acetatoxydans</i> Strain Re1	e00213-12
<i>Shahid Manzoor, Erik Bongcam-Rudloff, Anna Schnürer, Bettina Müller</i>	
Complete Genome Sequence of the Porcine Strain <i>Brachyspira pilosicoli</i> P43/6/78 ^T	e00215-12
<i>Changyou Lin, Henk C. den Bakker, Haruo Suzuki, Tristan Lefébure, Lalit Ponnala, Qi Sun, Michael J. Stanhope, Martin Wiedmann, Gérald E. Duhamel</i>	
Genome Sequence of <i>Mycoplasma feriruminatoris</i> sp. nov., a Fast-Growing <i>Mycoplasma</i> Species	e00216-12
<i>Anne Fischer, Ivette Santana-Cruz, Michelle Giglio, Suvarna Nadendla, Elliott Drabek, Edy M. Vilei, Joachim Frey, Joerg Jores</i>	
Draft Genome Sequence of the Halophilic Bacterium <i>Halobacillus</i> sp. Strain BAB-2008	e00222-12
<i>M. N. Joshi, A. S. Pandit, A. Sharma, R. V. Pandya, A. K. Saxena, S. B. Bagatharia</i>	

Complete Genome Sequence of the Piezophilic, Mesophilic, Sulfate-Reducing Bacterium <i>Desulfovibrio hydrothermalis</i> AM13 ^T	e00226-12
Boyang Ji, Gregory Gimenez, Valérie Barbe, Benoît Vacherie, Zoé Rouy, Amira Amrani, Marie-Laure Fardeau, Philippe Bertin, Didier Alazard, Sabine Leroy, Emmanuel Talla, Bernard Ollivier, Alain Dolla, Nathalie Pradel	
Genome Sequence of Naphthalene-Degrading Soil Bacterium <i>Pseudomonas putida</i> CSV86	e00234-12
Prashant S. Phale, Vasundhara Paliwal, Sajan C. Raju, Arnab Modak, Hemant J. Purohit	
Genome Sequence of <i>Alcaligenes</i> sp. Strain HPC1271	e00235-12
Atya Kapley, Sneha Sagarkar, Himgouri Tanksale, Nandita Sharma, Asifa Qureshi, Anshuman Khardenavis, Hemant J. Purohit	
Draft Genome Sequence of <i>Escherichia coli</i> Strain LCT-EC59	e00242-12
Tianzhi Li, Jiapeng Chen, De Chang, Xiangqun Fang, Junfeng Wang, Yinghua Guo, Longxiang Su, Guogang Xu, Yajuan Wang, Zhenhong Chen, Changting Liu	
Draft Genome Sequence of Uropathogenic <i>Escherichia coli</i> Strain J96	e00245-12
Eric A. Klein, Zemer Gitai	
Draft Genome Sequence of Chromate-Resistant and Biofilm-Producing Strain <i>Pseudomonas alcaliphila</i> 34	e00125-12
Luisa Santopolo, Emmanuela Marchi, Francesca Decorosi, Marco Galardini, Matteo Brilli, Luciana Giovannetti, Carlo Viti	
Draft Genome Sequences of Three <i>Salmonella enterica</i> Serotype Agona Strains from China	e00203-12
Jianmin Zhang, Guojie Cao, Xuebin Xu, Huimin Jin, Xiaowei Yang, Marc Allard, Eric Brown, Jianghong Meng	
Draft Genome Sequence of Type Strain <i>Clostridium pasteurianum</i> DSM 525 (ATCC 6013), a Promising Producer of Chemicals and Fuels	e00232-12
Sugima Rappert, Lifu Song, Wael Sabra, Wei Wang, An-Ping Zeng	
Complete Genome of <i>Enterobacteriaceae</i> Bacterium Strain FGI 57, a Strain Associated with Leaf-Cutter Ant Fungus Gardens	e00238-12
Frank O. Aylward, Daniel M. Tremmel, David C. Bruce, Patrick Chain, Amy Chen, Karen Walston Davenport, Chris Detter, Cliff S. Han, James Han, Marcel Huntemann, Natalia N. Ivanova, Nikos C. Kyrpides, Victor Markowitz, Kostas Mavromatis, Matt Nolan, Ioanna Pagani, Amrita Pati, Sam Pitluck, Shweta Deshpande, Lynne Goodwin, Tanja Woyke, Cameron R. Currie	
Draft Genome Sequence of a Phosphate-Accumulating <i>Bacillus</i> sp., WBUNB004	e00251-12
Shreya DebRoy, Pallavi Mukherjee, Sujata Roy, Ashoke Ranjan Thakur, Shaon RayChaudhuri	
Draft Genome Sequence of <i>Herbaspirillum huttense</i> subsp. <i>putei</i> IAM 15032, a Strain Isolated from Well Water	e00252-12
Vanely de Souza, Vitor C. Piro, Helisson Faoro, Michelle Z. Tadra-Sfeir, Vanessa K. Chicora, Dieval Guizelini, Vinicius Weiss, Ricardo A. Vialle, Rose A. Monteiro, Maria Berenice R. Steffens, Jeroniza N. Marchaukoski, Fabio O. Pedrosa, Leonardo M. Cruz, Leda S. Chubatsu, Roberto T. Raittz	
Draft Genome Sequence of a Nitrate- and Phosphate-Removing <i>Bacillus</i> sp., WBUNB009	e00254-12
Shreya DebRoy, Pallavi Mukherjee, Sujata Roy, Ashoke Ranjan Thakur, Shaon RayChaudhuri	

Complete Genome Sequence of the Industrial Strain <i>Gluconobacter oxydans</i> H24	e000003-13
<i>Xin Ge, Yan Zhao, Wei Hou, Weicai Zhang, Weiwei Chen, Jianhua Wang, Nan Zhao, Jian Lin, Wenxi Wang, Mengxia Chen, Qingge Wang, Yinghui Jiao, Zhigang Yuan, Xianghua Xiong</i>	
Whole-Genome Sequencing and Comparative Analysis of <i>Yersinia pestis</i> , the Causative Agent of a Plague Outbreak in Northern Peru	e00249-12
<i>O. Cáceres, J. Montenegro, C. Padilla, D. Tarazona, H. Bailón, P. García, M. Céspedes, P. Valencia, H. Guio</i>	
Genome Sequence of <i>Methanobrevibacter</i> sp. Strain JH1, Isolated from Rumen of Korean Native Cattle	e000002-13
<i>Jong-Hwan Lee, Moon-Soo Rhee, Sanjay Kumar, Geun-Hye Lee, Dong-Ho Chang, Dae-Soo Kim, Sang-Haeng Choi, Dong-Woo Lee, Min-Ho Yoon, Byoung-Chan Kim</i>	
Complete Genome Sequence of <i>Anoxybacillus flavithermus</i> TNO-09.006, a Thermophilic Sporeformer Associated with a Dairy-Processing Environment	e00010-13
<i>Martien P. M. Caspers, Jos Boekhorst, Tjakko Abe, Roland J. Siezen, Remco Kort</i>	
Genome Sequence of the “Indian Bison Type” Biotype of <i>Mycobacterium avium</i> subsp. <i>paratuberculosis</i> Strain S5	e00005-13
<i>Shoor Vir Singh, Naveen Kumar, Shree Narayan Singh, Tapas Bhattacharya, Jagdip Singh Sohal, Pravin Kumar Singh, Ajay Vir Singh, Brajesh Singh, Kundan Kumar Chaubey, Saurabh Gupta, Nitu Sharma, Shailesh Kumar, Gajendra Pal Singh Raghava</i>	
Draft Genome Sequence of <i>Brevibacillus</i> sp. Strain BAB-2500, a Strain That Might Play an Important Role in Agriculture	e00021-13
<i>M. N. Joshi, A. Sharma, A. S. Pandit, R. V. Pandya, A. K. Saxena, S. B. Bagatharia</i>	
Genome Sequence of <i>Enterobacter cancerogenus</i> YZ1	e00023-13
<i>Yifeng Wei, Yu Yang, Lisha Zhou, Zhangyi Liu, Xiangyan Wang, Rentao Yang, Qingqing Su, Yuping Zhou, Jiao Zhao, Jun Yang</i>	
Genome Sequence of <i>Lawsonia intracellularis</i> Strain N343, Isolated from a Sow with Hemorrhagic Proliferative Enteropathy	e00027-13
<i>Michelle Sait, Kevin Aitchison, Nick Wheelhouse, Kim Wilson, F. Alex Lainson, David Longbottom, David G. E. Smith</i>	
Draft Genome Sequence of the <i>Wolbachia</i> Endosymbiont of <i>Drosophila suzukii</i>	e00032-13
<i>Stefanos Siozios, Alessandro Cestaro, Rupinder Kaur, Ilaria Pertot, Omar Rota-Stabelli, Gianfranco Anfora</i>	
Whole-Genome Sequence of <i>Mycobacterium intracellulare</i> Clinical Strain MOTT-H4Y, Belonging to INT5 Genotype	e00006-13
<i>Hyungki Lee, Byoung-Jun Kim, Kijeong Kim, Seok-Hyun Hong, Yoon-Hoh Kook, Bum-Joon Kim</i>	
Genome Sequences and Photosynthesis Gene Cluster Composition of a Freshwater Aerobic Anoxygenic Phototroph, <i>Sandarakinorhabdus</i> sp. Strain AAP62, Isolated from the Shahu Lake in Ningxia, China	e00034-13
<i>Yonghui Zeng, Fuying Feng, Yapeng Liu, Yunxu Li, Michal Kobližek</i>	
Draft Genome Sequence of the Phyllosphere Model Bacterium <i>Pantoea agglomerans</i> 299R	e00036-13
<i>Mitja N. P. Remus-Emsermann, Eun Bae Kim, Maria L. Marco, Robin Tecon, Johan H. J. Leveau</i>	

VIRUSES

- Genome Sequence of a Novel Virus of the Species Human Adenovirus D Associated with Acute Gastroenteritis** **e00068-12**
Yuki Matsushima, Hideaki Shimizu, Atsuko Kano, Etsuko Nakajima, Yoko Ishimaru, Shuvra Kanti Dey, Yuki Watanabe, Fuyuka Adachi, Kohnosuke Mitani, Tsuguto Fujimoto, Tung Gia Phan, Hiroshi Ushijima
- Complete Genome Sequence of the *Crocuta crocuta* Papillomavirus Type 1 (CcrPV1) from a Spotted Hyena, the First Papillomavirus Characterized in a Member of the *Hyaenidae*** **e00062-12**
Hans Stevens, Elisabeth Heylen, Karlien De Keyser, Roger Maes, Matti Kiupel, Annabel Wise, Keith Nelson, Kay Holekamp, Anne Engh, Christy McKnight, Marc Van Ranst, Annabel Rector
- Complete Genome Sequence of the *Pseudomonas fluorescens* Bacteriophage UFV-P2** **e00006-12**
Monique R. Eller, Rafael L. Salgado, Pedro M. P. Vidigal, Maura P. Alves, Roberto S. Dias, Leandro L. de Oliveira, Cynthia C. da Silva, Antônio F. de Carvalho, Sérgio O. De Paula
- Complete Genome Sequence of the *Streptococcus suis* Temperate Bacteriophage φNJ2** **e00008-12**
Fang Tang, Alex Bossers, Frank Harders, Chengping Lu, Hilde Smith
- Complete Genome Sequence of a Novel Human Enterovirus 85 (HEV85) Recombinant with an Unknown New Serotype HEV-B Donor Sequence Isolated from a Child with Acute Flaccid Paralysis** **e00015-12**
Qiang Sun, Yong Zhang, Hui Cui, Shuangli Zhu, Zhen Zhu, Guohong Huang, Xiaolei Li, Bo Zhang, Dongmei Yan, Hongqiu An, Wenbo Xu
- Complete Genome Sequences of Two Feline Leukemia Virus Subgroup B Isolates with Novel Recombination Sites** **e00036-12**
H. Stewart, O. Jarrett, M. J. Hosie, B. J. Willett
- Complete Genome Sequence of *Bacillus thuringiensis* Bacteriophage BMBtp2** **e00011-12**
Zhaoxia Dong, Donghai Peng, Yueying Wang, Lei Zhu, Lifang Ruan, Ming Sun
- Genome Sequence of a Novel Archaeal Rudivirus Recovered from a Mexican Hot Spring** **e00040-12**
Luis E. Servín-Garcidueñas, Xu Peng, Roger A. Garrett, Esperanza Martínez-Romero
- Complete Genome Sequence of a Novel Porcine Parvovirus (PPV) Provisionally Designated PPV5** **e00021-12**
Chao-Ting Xiao, Patrick G. Halbur, Tanja Opriessnig
- Complete Genome Sequence of a Novel Natural Recombinant Porcine Reproductive and Respiratory Syndrome Virus Isolated from a Pig Farm in Yunnan Province, Southwest China** **e00003-12**
Yulin Yan, Aiguo Xin, Gaohong Zhu, Hui Huang, Qian Liu, Zhiyong Shao, Yating Zang, Ling Chen, Yongke Sun, Hong Gao
- Complete Genome Sequences of Classical Swine Fever Virus Isolates Belonging to a New Subgenotype, 2.1c, from Hunan Province, China** **e00080-12**
Da-Liang Jiang, Guo-Hua Liu, Wen-Jie Gong, Run-cheng Li, Yun-Fei Hu, Changchun Tu, Xing-Long Yu
- Complete Genome Sequence of a Newcastle Disease Virus Strain Belonging to a Recently Identified Genotype** **e00100-12**
François-Xavier Briand, Aurélie Henry, Paul Brown, Pascale Massin, Véronique Jestin

Complete Genome Sequence of the First Aichi Virus Isolated in Taiwan	e00107-12
<i>Jenn-Tzong Chang, Yao-Shen Chen, Bao-Chen Chen, David Chao, Tsung-Hsien Chang</i>	
Genome Sequences of a Novel HIV-1 Circulating Recombinant Form, CRF55_01B, Identified in China	e00050-12
<i>Xiaoxu Han, Minghui An, Weiqing Zhang, Weiping Cai, Xi Chen, Yutaka Takebe, Hong Shang</i>	
Complete Genome Sequence of a Porcine Reproductive and Respiratory Syndrome Virus Variant with a New Deletion in the 5' Untranslated Region	e00090-12
<i>Kaichuang Shi, Chun'ai Tao, Changhua Lin, Gang Li, Jun Li</i>	
Complete Nucleotide Sequence of Canine Distemper Virus HLJ1-06, Isolated from Foxes in China	e00065-12
<i>Qian Jiang, Xiaoliang Hu, Yanhua Ge, Huan Lin, Yong Jiang, Jiasen Liu, Dongchun Guo, Changde Si, Liandong Qu</i>	
Complete Genome Sequence of an Avian Paramyxovirus Type 4 from North America Reveals a Shorter Genome and New Genotype	e00075-12
<i>Manoharan Parthiban, Manimaran Kaliyaperumal, Sa Xiao, Baibaswata Nayak, Anandan Paldurai, Shin-Hee Kim, Brian S. Ladman, Lauren A. Preskenis, Jack Gelb, Jr., Peter L. Collins, Siba K. Samal</i>	
Complete Genome Sequence of Bacteriophage EC6, Capable of Lysing <i>Escherichia coli</i> O157:H7	e00085-12
<i>Birendra R. Tiwari, Jungmin Kim</i>	
Complete Genome Analysis of Canine Respiratory Coronavirus	e00093-12
<i>Seong-in Lim, Sarah Choi, Ji-Ae Lim, Hye-Young Jeoung, Jae-Young Song, R. C. dela Pena, Dong-Jun An</i>	
Full Genome Sequence of a Novel Human Enterovirus C (EV-C118) Isolated from Two Children with Acute Otitis Media and Community-Acquired Pneumonia in Israel	e00121-12
<i>Cristina Daleno, Antonio Piralla, Alessia Scala, Fausto Baldanti, David Greenberg, Nicola Principi, Susanna Esposito, CAP-PRI Study Group</i>	
Complete Genome Sequences of Six Avian-Like H1N1 Swine Influenza Viruses from Northwestern China	e00098-12
<i>Jing-Yu Wang, Juan-Juan Ren, Yuan-Hao Qiu, Hung-Jen Liu</i>	
Complete Genome Sequence of a Rabies Virus Isolate from Cattle in Guangxi, Southern China	e00137-12
<i>Hai-Bo Tang, Xiao-Xia He, Yi-Zhi Zhong, Su-Huan Liao, Tao-Zhen Zhong, Lin-Juan Xie, Yan Pan, Zhan-Ling Lu, Xian-Kai Wei, Yang Luo, Ting Rong Luo</i>	
Complete Genome Sequence of Sacbrood Virus Strain SBM2, Isolated from the Honeybee <i>Apis cerana</i> in Vietnam	e00076-12
<i>Nga Thi Bich Nguyen, Thanh Hoa Le</i>	
Complete Genome Sequence of a Highly Pathogenic Porcine Reproductive and Respiratory Syndrome Virus Variant Isolated from a Backyard Piglet	e00123-12
<i>Kaichuang Shi, Shenglan Mo, Huaxin Huang, Changhua Lin, Chun'ai Tao, Jun Li, Gang Li</i>	
Complete Genome Sequences of <i>Edwardsiella tarda</i>-Lytic Bacteriophages KF-1 and IW-1	e00089-12
<i>Motoshige Yasuike, Emi Sugaya, Yoji Nakamura, Yuya Shigenobu, Yasuhiko Kawato, Wataru Kai, Atushi Fujiwara, Motohiko Sano, Takanori Kobayashi, Toshihiro Nakai</i>	

Complete Genome Sequence of a Novel Recombinant Human Norovirus Genogroup II Genotype 4 Strain Associated with an Epidemic during Summer of 2012 in Hong Kong <i>Martin C. W. Chan, Paul K. S. Chan</i>	e00140-12
Complete Genome Sequence of a Human Coxsackievirus B3 from a Child with Myocarditis in Beijing, China <i>Jiang Du, Zhiqiang Wu, Ying Xue, Ting Zhang, Fan Yang, Qi Jin</i>	e00163-12
Complete Genome Sequence of the Novel Duck Circovirus Strain GH01 from Southwestern China <i>Zhilong Zhang, Renyong Jia, Mingshu Wang, Yanyan Lu, Dekang Zhu, Shun Chen, Zhongqiong Yin, Yin Wang, Xiaoyue Chen, Anchun Cheng</i>	e00166-12
The Genome of <i>Cronobacter sakazakii</i> Bacteriophage vB_CsaP_GAP227 Suggests a New Genus within the <i>Autographivirinae</i> <i>Reza Abbasifar, Andrew M. Kropinski, Parviz M. Sabour, Hans-Wolfgang Ackermann, Argentina Alanis Villa, Arash Abbasifar, Mansel W. Griffiths</i>	e00122-12
Complete Genome Sequence of a Natural Recombinant H9N2 Influenza Virus from Wild Birds in Republic of Korea <i>Dong-Hun Lee, Jae-Keun Park, Seong-Su Yuk, Tseren-Ochir Erdene-Ochir, Jung-Hoon Kwon, Joong-Bok Lee, Seung-Yong Park, In-Soo Choi, Chang-Seon Song</i>	e00159-12
Genome Sequence of the Hepatitis C Virus Subtype 6n Isolated from Malaysia <i>Kim Tien Ng, Yeat Mei Lee, Haider Abdulrazzaq Abed Al-Darraji, Xueshan Xia, Yutaka Takebe, Kok Gan Chan, Ling Lu, Sanjiv Mahadeva, Adeeba Kamarulzaman, Kok Keng Tee</i>	e00168-12
Complete Genome Sequence of GII.4 Human Norovirus HS191 <i>Stanislav V. Sosnovtsev, Karin Bok, QiuHong Wang, Linda J. Saif, Kim Y. Green</i>	e00169-12
Complete Genome Sequence of a Highly Virulent Newcastle Disease Virus Currently Circulating in Mexico <i>Sa Xiao, Anandan Paldurai, Baibaswata Nayak, Armando Mirande, Peter L. Collins, Siba K. Samal</i>	e00177-12
Complete Genome Sequence of Human Adenovirus Type 7 Associated with Fatal Infant Pneumonia <i>Liuying Tang, Junjing An, Pengbo Yu, Wenbo Xu</i>	e00182-12
Complete Genome Sequences of Two Newcastle Disease Virus Strains of Genotype VIII <i>Yongzhong Cao, Min Gu, Xiaorong Zhang, Wenbo Liu, Xiufan Liu</i>	e00180-12
Complete Genome of a Genotype I Japanese Encephalitis Virus Isolated from a Patient with Encephalitis in Vientiane, Lao PDR <i>Fabien Aubry, Manivanh Vongsouvath, Antoine Nougairède, Rattanaphone Phetsouvanh, Bountoy Sibounheuang, Rémi Charrel, Sayaphet Rattanavong, Koueko Phommasonne, Onanong Sengvilaiprasertth, Xavier de Lamballerie, Paul N. Newton, Audrey Dubot-Pérès</i>	e00157-12
Complete Genome Sequence of a Street Rabies Virus Isolated from a Dog in Nigeria <i>Ming Zhou, Zutao Zhou, Grace S. N. Kia, Clement W. Gnanadurai, Christina M. Leyson, Jarlath U. Umoh, Jacob P. Kwaga, Haruna M. Kazeem, Zhen F. Fu</i>	e00214-12
Full Genome Sequence of Giant Panda Rotavirus Strain CH-1 <i>Ling Guo, Qigui Yan, Shaolin Yang, Chengdong Wang, Shijie Chen, Xiaonong Yang, Rong Hou, Zifang Quan, Zhongxiang Hao</i>	e00241-12

- Complete Genome Sequence of a Variant Porcine Epidemic Diarrhea Virus Strain Isolated in Central China e00243-12**
Xiao-Meng Wang, Bei-Bei Niu, He Yan, Dong-Sheng Gao, Jin-Yao Huo, Lu Chen, Hong-Tao Chang, Chuan-qing Wang, Jun Zhao
- Complete Genome Sequences of Bovine Parainfluenza Virus Type 3 Strain BN-1 and Vaccine Strain BN-CE e00247-12**
Takashi Ohkura, Takehiro Kokuhō, Misako Konishi, Ken-ichiro Kameyama, Kaoru Takeuchi
- Complete Genome Sequences of New Emerging Newcastle Disease Virus Strains Isolated from China e00129-12**
Hualei Liu, Yan Lv, Claudio L. Afonso, Shengqiang Ge, Dongxia Zheng, Yunling Zhao, Zhiliang Wang
- Complete Genome Sequence of *Escherichia coli* O157:H7 Bacteriophage phiJLA23 Isolated in Mexico e00219-12**
Luis Amarillas, Cristobal Chaidez, Yadira Lugo, Josefina León-Félix
- Complete Genome Sequence of a Novel Myovirus Which Infects Atypical Strains of *Edwardsiella tarda* e00248-12**
Motoshige Yasuike, Emi Sugaya, Yoji Nakamura, Yuya Shigenobu, Yasuhiko Kawato, Wataru Kai, Satoshi Nagai, Atushi Fujiwara, Motohiko Sano, Takanori Kobayashi, Toshihiro Nakai