

Rui Tian

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| 45 | A Genomic Encyclopedia of the Root Nodule Bacteria: assessing genetic diversity through a systematic biogeographic survey. <i>Standards in Genomic Sciences</i> , 2015 , 10, 14 | | 46 |
| 44 | Genome sequence of the Lotus spp. microsymbiont Mesorhizobium loti strain R7A. <i>Standards in Genomic Sciences</i> , 2014 , 9, 6 | | 16 |
| 43 | Molecular and serological diversity in Apple chlorotic leaf spot virus from sand pear (<i>Pyrus pyrifolia</i>) in China. <i>European Journal of Plant Pathology</i> , 2011 , 130, 183-196 | 2.1 | 16 |
| 42 | Genome sequence of Ensifer sp. TW10; a Tephrosia wallichii (Biyani) microsymbiont native to the Indian Thar Desert. <i>Standards in Genomic Sciences</i> , 2013 , 9, 304-14 | | 11 |
| 41 | Genome sequence of Ensifer meliloti strain WSM1022; a highly effective microsymbiont of the model legume Medicago truncatula A17. <i>Standards in Genomic Sciences</i> , 2013 , 9, 315-24 | | 9 |
| 40 | Genome sequence of the clover-nodulating Rhizobium leguminosarum bv. trifolii strain TA1. <i>Standards in Genomic Sciences</i> , 2013 , 9, 243-53 | | 8 |
| 39 | Genome sequence of the Lotus corniculatus microsymbiont Mesorhizobium loti strain R88B. <i>Standards in Genomic Sciences</i> , 2014 , 9, 3 | | 7 |
| 38 | High-quality permanent draft genome sequence of Rhizobium sullae strain WSM1592; a Hedysarum coronarium microsymbiont from Sassari, Italy. <i>Standards in Genomic Sciences</i> , 2015 , 10, 44 | | 7 |
| 37 | High-quality permanent draft genome sequence of Ensifer sp. PC2, isolated from a nitrogen-fixing root nodule of the legume tree (Khejri) native to the Thar Desert of India. <i>Standards in Genomic Sciences</i> , 2016 , 11, 43 | | 6 |
| 36 | Genome sequence of Microvirga lupini strain LUT6(T), a novel Lupinus alphasproteobacterial microsymbiont from Texas. <i>Standards in Genomic Sciences</i> , 2014 , 9, 1159-67 | | 5 |
| 35 | High-quality permanent draft genome sequence of the Lebeckia ambigua-nodulating Burkholderia sp. strain WSM4176. <i>Standards in Genomic Sciences</i> , 2015 , 10, 79 | | 5 |
| 34 | Genome sequence of the Lotus spp. microsymbiont Mesorhizobium loti strain NZP2037. <i>Standards in Genomic Sciences</i> , 2014 , 9, 7 | | 4 |
| 33 | High-quality permanent draft genome sequence of the type strain USDA 76, isolated from (<i>L.</i>) Merr. <i>Standards in Genomic Sciences</i> , 2017 , 12, 26 | | 4 |
| 32 | The genome sequences of three isolates of Apple chlorotic leaf spot virus from pear (<i>Pyrus</i> sp.) in China. <i>Canadian Journal of Plant Pathology</i> , 2014 , 36, 396-402 | 1.6 | 4 |
| 31 | Genome sequence of the Listia angolensis microsymbiont Microvirga lotononidis strain WSM3557(T.). <i>Standards in Genomic Sciences</i> , 2014 , 9, 540-50 | | 4 |
| 30 | Genome sequence of Ensifer arboris strain LMG 14919(T); a microsymbiont of the legume Prosopis chilensis growing in Kosti, Sudan. <i>Standards in Genomic Sciences</i> , 2014 , 9, 473-83 | | 4 |
| 29 | High-quality permanent draft genome sequence of Bradyrhizobium sp. Tv2a.2, a microsymbiont of Tachigali versicolor discovered in Barro Colorado Island of Panama. <i>Standards in Genomic Sciences</i> , 2015 , 10, 27 | | 3 |

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| 28 | Genome sequence of the dark pink pigmented <i>Listia bainesii</i> microsymbiont <i>Methylobacterium</i> sp. WSM2598. <i>Standards in Genomic Sciences</i> , 2014 , 9, 5 | 3 |
| 27 | Genome sequence of the <i>Medicago-nodulating</i> <i>Ensifer meliloti</i> commercial inoculant strain RRI128. <i>Standards in Genomic Sciences</i> , 2014 , 9, 602-13 | 3 |
| 26 | High-quality permanent draft genome sequence of <i>Rhizobium leguminosarum</i> bv. <i>viciae</i> strain GB30; an effective microsymbiont of <i>Pisum sativum</i> growing in Poland. <i>Standards in Genomic Sciences</i> , 2015 , 10, 36 | 3 |
| 25 | High-quality permanent draft genome sequence of <i>Ensifer meliloti</i> strain 4H41, an effective salt- and drought-tolerant microsymbiont of <i>Phaseolus vulgaris</i> . <i>Standards in Genomic Sciences</i> , 2015 , 10, 34 | 3 |
| 24 | Genome sequence of <i>Bradyrhizobium</i> sp. WSM1253; a microsymbiont of <i>Ornithopus compressus</i> from the Greek Island of Sifnos. <i>Standards in Genomic Sciences</i> , 2015 , 10, 113 | 3 |
| 23 | Genome sequence of the clover-nodulating <i>Rhizobium leguminosarum</i> bv. <i>trifolii</i> strain SRDI943. <i>Standards in Genomic Sciences</i> , 2013 , 9, 232-42 | 3 |
| 22 | Genome sequence of the <i>Trifolium rueppellianum</i> -nodulating <i>Rhizobium leguminosarum</i> bv. <i>trifolii</i> strain WSM2012. <i>Standards in Genomic Sciences</i> , 2013 , 9, 283-93 | 3 |
| 21 | Genome sequence of the clover-nodulating <i>Rhizobium leguminosarum</i> bv. <i>trifolii</i> strain SRDI565. <i>Standards in Genomic Sciences</i> , 2013 , 9, 220-31 | 3 |
| 20 | Genome sequence of the South American clover-nodulating <i>Rhizobium leguminosarum</i> bv. <i>trifolii</i> strain WSM597. <i>Standards in Genomic Sciences</i> , 2013 , 9, 264-72 | 3 |
| 19 | Genome sequence of the acid-tolerant <i>Burkholderia</i> sp. strain WSM2230 from Karijini National Park, Australia. <i>Standards in Genomic Sciences</i> , 2014 , 9, 551-61 | 3 |
| 18 | High-quality permanent draft genome sequence of the <i>Parapiptadenia rigida</i> -nodulating <i>Burkholderia</i> sp. strain UYPR1.413. <i>Standards in Genomic Sciences</i> , 2015 , 10, 31 | 2 |
| 17 | High-quality permanent draft genome sequence of the <i>Parapiptadenia rigida</i> -nodulating <i>Cupriavidus</i> sp. strain UYPR2.512. <i>Standards in Genomic Sciences</i> , 2015 , 10, 13 | 2 |
| 16 | Genome sequence of <i>Burkholderia mimosarum</i> strain LMG 23256(T), a <i>Mimosa pigra</i> microsymbiont from Anso, Taiwan. <i>Standards in Genomic Sciences</i> , 2014 , 9, 484-94 | 2 |
| 15 | Genome sequence of the lupin-nodulating <i>Bradyrhizobium</i> sp. strain WSM1417. <i>Standards in Genomic Sciences</i> , 2013 , 9, 273-82 | 2 |
| 14 | Genome sequence of the acid-tolerant <i>Burkholderia</i> sp. strain WSM2232 from Karijini National Park, Australia. <i>Standards in Genomic Sciences</i> , 2014 , 9, 1168-80 | 2 |
| 13 | Evolution of a multi-step phosphorelay signal transduction system in <i>Ensifer</i> : recruitment of the sigma factor RpoN and a novel enhancer-binding protein triggers acid-activated gene expression. <i>Molecular Microbiology</i> , 2017 , 103, 829-844 | 4.1 1 |
| 12 | High-quality draft genome sequence of strain STM6155, a microsymbiont from New Caledonia. <i>Standards in Genomic Sciences</i> , 2017 , 12, 7 | 1 |
| 11 | High-quality permanent draft genome sequence of the <i>Lebeckia</i> - nodulating <i>Burkholderia dilworthii</i> strain WSM3556(T). <i>Standards in Genomic Sciences</i> , 2015 , 10, 64 | 1 |

- 10 High-quality permanent draft genome sequence of Bradyrhizobium sp. strain WSM1743 - an effective microsymbiont of an Indigofera sp. growing in Australia. *Standards in Genomic Sciences*, **2015**, 10, 87 1
- 9 High-quality permanent draft genome sequence of Bradyrhizobium sp. Ai1a-2; a microsymbiont of Andira inermis discovered in Costa Rica. *Standards in Genomic Sciences*, **2015**, 10, 33 1
- 8 High-quality permanent draft genome sequence of the Mimosa asperata - nodulating Cupriavidus sp. strain AMP6. *Standards in Genomic Sciences*, **2015**, 10, 80 1
- 7 High-quality permanent draft genome sequence of Bradyrhizobium sp. Th.b2, a microsymbiont of Amphicarpaea bracteata collected in Johnson City, New York. *Standards in Genomic Sciences*, **2015**, 10, 24 1
- 6 Genome sequence of Ensifer medicae strain WSM1369; an effective microsymbiont of the annual legume Medicago sphaerocarpos. *Standards in Genomic Sciences*, **2013**, 9, 420-30 1
- 5 Genome sequence of Ensifer medicae strain WSM1115; an acid-tolerant Medicago-nodulating microsymbiont from Samothraki, Greece. *Standards in Genomic Sciences*, **2014**, 9, 514-26 1
- 4 High-quality permanent draft genome sequence of Ensifer medicae strain WSM244, a microsymbiont isolated from Medicago polymorpha growing in alkaline soil. *Standards in Genomic Sciences*, **2015**, 10, 126 0
- 3 High-quality draft genome sequence of Mlalz-1, a microsymbiont of (L.) miller collected in Lanzarote, Canary Islands, Spain. *Standards in Genomic Sciences*, **2017**, 12, 58
- 2 Genome sequence of Ensifer medicae Di28; an effective N₂-fixing microsymbiont of Medicago murex and M. polymorpha. *Standards in Genomic Sciences*, **2014**, 9, 4
- 1 Genome sequence of the Ornithopus/Lupinus-nodulating Bradyrhizobium sp. strain WSM471. *Standards in Genomic Sciences*, **2013**, 9, 254-63