Genome of an arbuscular mycorrhizal fungus provides i symbiosis

Proceedings of the National Academy of Sciences of the Unite 110, 20117-20122

DOI: 10.1073/pnas.1313452110

Citation Report

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1	The Largest Subunit of RNA Polymerase II as a New Marker Gene to Study Assemblages of Arbuscular Mycorrhizal Fungi in the Field. PLoS ONE, 2014, 9, e107783.	1.1	55
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4	Exploring the Transcriptome of Mycorrhizal Interactions. Advances in Botanical Research, 2014, 70, 53-78.	0.5	8
5	Lipid Droplets of Arbuscular Mycorrhizal Fungi Emerge in Concert with Arbuscule Collapse. Plant and Cell Physiology, 2014, 55, 1945-1953.	1.5	41
6	A dipeptide transporter from the arbuscular mycorrhizal fungus Rhizophagus irregularis is upregulated in the intraradical phase. Frontiers in Plant Science, 2014, 5, 436.	1.7	47
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8	Transcriptional responses of Medicago truncatula upon sulfur deficiency stress and arbuscular mycorrhizal symbiosis. Frontiers in Plant Science, 2014, 5, 680.	1.7	31
9	Genome-wide analysis of copper, iron and zinc transporters in the arbuscular mycorrhizal fungus Rhizophagus irregularis. Frontiers in Plant Science, 2014, 5, 547.	1.7	120
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19	DNA-based detection and identification of Glomeromycota: the virtual taxonomy of environmental sequences. Botany, 2014, 92, 135-147.	0.5	170

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21	Signaling events during initiation of arbuscular mycorrhizal symbiosis. Journal of Integrative Plant Biology, 2014, 56, 250-261.	4.1	102
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