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RESEARCHERS COMPLETE DRAFT GENOME SEQUENCE FOR CASSAVA

\$1.3 Million grant from Bill & Melinda Gates Foundation will fund the next phase of research critical to global food security

St. Louis, MO Nov. 9, 2009 – A team of academic, government and industry researchers has completed a first draft of the cassava (*Manihot esculenta*) genome. The project is an important first step in accelerating the pace of research on this subsistence crop and addressing some of the many limitations that face cassava farmers around the world.

Cassava is a root crop that serves as the primary food source for more than 750 million people each day. Although it has many properties that make it an important food across much of Africa and Asia, it also has many limitations. Cassava has poor nutritional content and is susceptible to many pathogens, particularly in Africa, where one third of the continental harvest is lost each year to viral diseases. One of these, Cassava Brown Streak Disease, or CBSD, is currently the major threat to food security in some parts of Eastern Africa.

The impetus for the genome sequence began in 2003 with the formation of The Global Cassava Partnership (GCP-21), co-chaired by Dr. Claude Fauquet, director of the International Laboratory for Tropical Agriculture Biology (ILTAB) at the Donald Danforth Plant Science Center (DDPSC) in St. Louis, and Dr. Joe Tohme of the International Center for Tropical Agriculture (CIAT) in Cali, Colombia. This, in turn, led to a 2006 proposal by Fauquet, Tohme and 12 other international scientists to DOE JGI's Community Sequencing Program, which was selected for a pilot project.

The full genome project gathered momentum in early 2009 when 454 Life Sciences and DOE JGI each pledged the resources to use 454's Genome Sequencer FLX platform with long-read GS FLX Titanium chemistry to rapidly generate the DNA sequence data needed for the project. More than 61 million sequencing reads were generated and assembled into a draft genome that contains an estimated 95 percent of cassava genes. It is one of the first large genome projects to primarily use 454 Life Sciences' long-read sequencing platform, which enabled both improved quality of the draft, and its rapid generation.

The annotated draft genome sequence is available at DOE JGI's Phytozome Web site, www.phytozome.net/cassava.

“By 2050, 90 percent of humankind will live in developing countries where agriculture is the most important economic activity. Crops grown by small farmers are central to international food security, health, economic growth, energy needs, poverty reduction and social stability,” Fauquet said. “The information contained in the cassava genome will provide tremendous opportunities to improve this important crop, bringing it into the mainstream of plant research thereby reducing the time and cost of delivering improved cultivars to farmers who need it most.”

“Sequencing of the cassava genome is a major and significant achievement for the cassava research community. The data obtained will accelerate the development of new and more efficient molecular markers to characterize the *Manihot* collections and to assist breeding programs to develop varieties adapted to the needs of the poor in developing countries. It will also strengthen ongoing functional genomics efforts,” said Tohme

In response to the urgency of this threat, and building upon the newly available cassava genome sequence, the Bill & Melinda Gates Foundation has awarded a \$1.3 million grant to University of Arizona researchers who will lead an international consortium to develop a genome variation database that will provide breeding tools to aid farmers in improving cassava, with a special focus on increased resistance to the CBSD virus.

This grant is part of the foundation’s Agricultural Development initiative, which is working with a wide range of partners to provide millions of small farmers in the developing world with tools and opportunities to boost their yields, increase their incomes and build better lives for themselves and their families. The foundation is working to strengthen the entire agricultural value chain – from seeds and soil to farm management and market access—so that progress against hunger and poverty is sustainable over the long term.

The availability of the genome sequence enables the newly-funded project to study how cassava varieties differ from each other. “The contributions of 454 Life Sciences and DOE JGI in making the cassava genome a reality have opened a new chapter in cassava research worldwide. We’re excited about the opportunity for cassava breeders to access new tools for improving a staple African crop,” said Katherine Kahn, program officer with the Agricultural Development initiative at the Bill & Melinda Gates Foundation.

Steve Rounsley, PhD, associate professor in the School of Plant Sciences at the UA and a member of the BIO5 Institute, will coordinate the project that includes partners at the Institute for Genome Sciences, University of Maryland, Baltimore, the U.S. Department of Energy Joint Genome Institute (DOE JGI), and 454 Life Sciences, a Roche Company.

“This is a perfect example of how quickly things can happen when everyone is aligned behind an important cause. Most of the data for the genome were generated within 8 weeks of getting DOE JGI and 454 Life Sciences on board,” said the UA’s Rounsley, who led the collaboration.

“We are pleased to contribute our sequencing technology to this important global initiative,” explained Michael Egholm, Chief Technology Officer and Vice President of Research and Development at 454 Life Sciences. “This project, along with other recently completed complex plant genome projects, demonstrates that 454 Sequencing systems are rapidly becoming the standard for *de novo* sequencing and assembly.”

Researchers will use next-generation technologies to sample many varieties of cassava and develop a large database of markers that can be used to identify genes involved in many important traits. The team will collaborate with researchers in Kenya, Uganda and Tanzania in applying these genetic markers toward identifying resistance to Cassava Brown Streak Disease. All of the information and tools the project develops will be freely available worldwide.

Traditional cassava improvement is slow and difficult. The availability of large numbers of markers will help make breeding schemes more efficient. For instance, traits that may only show up in mature plants can be identified in seedlings with a cheap DNA test. Since cassava is used for industrial starch production, and has potential as a biofuel source, there are commercial applications of these breeding tools. However, the most important applications will be those that improve the lives of those who depend upon cassava for their daily calorie intake.

The Global Cassava Partnership (GCP-21)

Founded and chaired by Dr. Claude Fauquet and Dr. Joe Tohme, the GCP-21 is an alliance of the cassava research and development community, that are working to identify the major constraints to unlock the productivity potential of cassava. The sequencing of the cassava genome was one of these major constraints, and its establishment will permit the development of many more tools that will be made available for free to the cassava community. These modern tools will provide new tools to breeders and biotechnologists to speed up the improvement of cassava to benefit of hundreds of millions of people in the world. For more information please visit, <http://www.danforthcenter.org/GCP21/>.

About The Donald Danforth Plant Science Center

Founded in 1998, the Donald Danforth Plant Science Center is a not-for-profit research institute with a mission to improve the human condition through plant science. Research at the Danforth Center will feed the hungry and improve human health, preserve and renew the environment, and enhance the St. Louis region and Missouri as a world center for plant science. Please visit www.danforthcenter.org for additional information.

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About the International Center for Tropical Agriculture (CIAT)

The International Center for Tropical Agriculture (CIAT) is a not-for-profit organization that works to reduce hunger and poverty, and improve human health in the tropics through research aimed at increasing the eco-efficiency of agriculture. CIAT is one of the 15 centers funded mainly by the 64 countries, private foundations, and international organizations that make up the Consultative Group on International Agricultural Research (CGIAR). The plant genetic resources conserved by CIAT are a component of the world "designate collection" of the UN Food and Agriculture Organization (FAO). CIAT holds in trust one of the largest cassava collection in the world. CIAT makes its germplasm available free of charge, upon request, to farmers, farmer associations, breeders, geneticists, agronomists, extension agencies, universities, and biodiversity institutes with a clearly articulated need. Please visit www.ciat.cgiar.org for additional information.

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About the UA's BIO5 Institute

The BIO5 Institute at the University of Arizona is at the leading edge of life sciences research, bringing together faculty and other researchers from five disciplines – agriculture, medicine, pharmacy, basic science and engineering – to tackle complex biology-based problems affecting humanity today. Learn more at <http://bio5.org/>

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About the Institute for Genome Sciences, University of Maryland, Baltimore

The Institute for Genome Sciences (IGS) at the University of Maryland, Baltimore is an international research center dedicated to advancing the use of genomics to improve human life. Led by Dr. Claire Fraser-Liggett, a preeminent genome scientist and microbiologist, IGS is located in a 10-acre BioPark in downtown Baltimore. IGS scientists integrate genomics, bioinformatics and metagenomics into biomedical research. For more information, see <http://www.igs.umaryland.edu>.

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About U.S. Dept. of Energy Joint Genomic Institute (DOE JGI)

The U.S. Department of Energy Joint Genome Institute (<http://www.jgi.doe.gov>), supported by DOE's Office of Science, is committed to advancing genomics in support of DOE missions related to clean energy generation and environmental characterization and cleanup. DOE JGI, headquartered in Walnut Creek, Calif., provides integrated high-throughput sequencing and computational analysis that enable systems-based scientific approaches to these challenges. Follow DOE JGI on Twitter.

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About 454 Life Sciences

454 Life Sciences, a center of excellence of Roche Applied Science, develops and commercializes the innovative 454 Sequencing system for ultra-high-throughput DNA sequencing. Specific applications include de novo sequencing and re-sequencing of genomes, metagenomics, RNA analysis, and targeted sequencing of DNA regions of interest. The hallmarks of the 454 Sequencing system are its simple, unbiased sample preparation and long, highly accurate sequence reads, including paired-end reads. The system has enabled hundreds of peer-reviewed studies in diverse research fields, such as cancer and infectious disease research, drug discovery, marine biology, anthropology, paleontology and many more.

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