

**From:** [Scott Allen Jackson](mailto:Scott.Allen.Jackson)  
**To:** [Francois.Belzile@fsaa.ulaval.ca](mailto:Francois.Belzile@fsaa.ulaval.ca)  
**Subject:** Fwd: Seattle Meeting, April 15-17  
**Date:** Monday, February 23, 2015 10:06:23 AM

---

Hi Dr. Belzile,

I'm sorry for the late notice, but I was just given your name as someone that may be interested in this discussion/meeting on the application of genomics to the world's collection of soybean genetic resources. If you are interested, please let me know and feel free to register at the hotel with the information below. There is no money for participant support, everyone is doing it on their own. We have representatives from Embrapa-soja, USDA, US universities, China, Japan, CGIAR and Global Crop Diversity Trust/Divseek.

Let me know if you have any questions and if you will be able to attend.

Thanks, Scott  
Scott Jackson  
Director, Center for Applied Genetic Technologies  
Institute for Plant Breeding, Genetics and Genomics  
University of Georgia  
111 Riverbend Rd.  
Athens, GA 30621

ph 706.542.4021  
[sjackson@uga.edu](mailto:sjackson@uga.edu)

Begin forwarded message:

**From:** Scott Allen Jackson <[sjackson@uga.edu](mailto:sjackson@uga.edu)>  
**Subject:** Seattle Meeting, April 15-17  
**Date:** February 19, 2015 at 12:19:56 AM EST  
**Cc:** Jenny Leverett <[jkrogersw@gmail.com](mailto:jkrogersw@gmail.com)>  
**To:** "Jones, Jennifer" <[jjones@smithbucklin.com](mailto:jjones@smithbucklin.com)>, "Abberton, Michael (IITA)" <[m.abberton@cgiar.org](mailto:m.abberton@cgiar.org)>, "Cannon, Steven [AGRON]" <[scannon@iastate.edu](mailto:scannon@iastate.edu)>, "Joost, Richard" <[RJoost@smithbucklin.com](mailto:RJoost@smithbucklin.com)>, "Nelson, Randall" <[Randall.Nelson@ars.usda.gov](mailto:Randall.Nelson@ars.usda.gov)>, "Okamoto, Jack" <[Jack.Okamoto@ars.usda.gov](mailto:Jack.Okamoto@ars.usda.gov)>, Peter Wenzl <[peter.wenzl@cropprtrust.org](mailto:peter.wenzl@cropprtrust.org)>, Suk-Ha Lee <[sukhalee@snu.ac.kr](mailto:sukhalee@snu.ac.kr)>, "邱丽娟 QIU, Lijuan" <[qiulijuan@caas.cn](mailto:qiulijuan@caas.cn)>, 李英慧 <[liyinghui@caas.cn](mailto:liyinghui@caas.cn)>, "Masao Ishimoto" <[ishimoto@affrc.go.jp](mailto:ishimoto@affrc.go.jp)>, Akito Kaga <[kaga@nias.affrc.go.jp](mailto:kaga@nias.affrc.go.jp)>, "Bob Stupar" <[stup0004@umn.edu](mailto:stup0004@umn.edu)>, Ricardo Abdelnoor <[ricardo.abdelnoor@embrapa.br](mailto:ricardo.abdelnoor@embrapa.br)>, <[marcelofernandes.oliveira@embrapa.br](mailto:marcelofernandes.oliveira@embrapa.br)>, Steven Cannon <[steven.cannon@ARS.USDA.GOV](mailto:steven.cannon@ARS.USDA.GOV)>

A reminder for those that have not registered yet. Please register per the directions below as soon as possible.

Thanks, Scott

—————sent previously Feb 12—————

Hi everyone,

Details below on the meeting on the application of genomics for soybean genomic resources.

1) **Hotel:** I have a room block reserved at 15-17 April at the Seattle Downtown Courtyard Marriott (612 2nd Avenue Seattle WA 98104 ). **You need to immediately register your room** under your name and credit card. You can do so by using the following link: [Book your group rate for Germplasm and Genomics Meeting Room Block](#). The full link is at bottom of email in case the embedded link does not work.

There is a light rail (<http://www.soundtransit.org/Fares-and-Passes/Link-fares>) that will take you from the airport to 0.1 miles from the hotel (get off at Pioneer Square). The cost is \$2.75 USD.

2) **Proposed meeting agenda** (this is a beginning outline. If you have additional ideas/thoughts, please let me know).

Purpose: to explore the possible application of genomics and phenomics to the international soybean genetic resources.

Agenda

\* Introductions

\* Peter Wenzl: introduction to DivSeek initiative ([www.divseek.org](http://www.divseek.org))

\* Short 8 minute reports by country/collection on a) collection, b) application of genomics/high density markers for characterization, c) phenotyping and d) utilization.

Brazil

China

Japan

Korea

United States

CGIAR

\* Randy Nelson: 50K SNPs on US collection (15 min)

\* Lijuan Qiu: sequence analysis of soybean and G. soja (15 Min)

\* Robert Stupar: non-SNP variation (15 min)

Discussion 1: How to collectively characterize collections

Discussion 2: Data sharing

Discussion 3: Germplasm issues: redundancy, sharing, etc.

Thanks and let me know if you have any questions,

Scott Jackson

Link for hotel reservations:

[https://www.marriott.com/meeting-event-hotels/group-corporate-travel/groupCorp.mi?resLinkData=Germplasm%20and%20Genomics%20Meeting%20Room%20Block%5Eseaps%60GGMGGMA%60159.00%60USD%60false%604/15/15%604/17/15%603/25/15&app=resvlink&stop\\_mobi=yes](https://www.marriott.com/meeting-event-hotels/group-corporate-travel/groupCorp.mi?resLinkData=Germplasm%20and%20Genomics%20Meeting%20Room%20Block%5Eseaps%60GGMGGMA%60159.00%60USD%60false%604/15/15%604/17/15%603/25/15&app=resvlink&stop_mobi=yes)