

**From:** [Scott Allen Jackson](#)  
**To:** [Masao Ishimoto](#)  
**Cc:** [Akito Kaga](#)  
**Subject:** Re: Seattle Meeting, April 15-17  
**Date:** Tuesday, April 14, 2015 8:27:06 AM

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Great. I will follow up with details.  
scott

On Apr 14, 2015, at 12:26 AM, Masao Ishimoto <[ishimoto@affrc.go.jp](mailto:ishimoto@affrc.go.jp)> wrote:

Dear Scott,

We will also arrive early enough for dinner on Wed (15th).  
We look forward to hearing from you .

Regards,

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Masao ISHIMOTO

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On 2015/04/14 2:56, "Scott Allen Jackson" <[sjackson@uga.edu](mailto:sjackson@uga.edu)> wrote:

Folks, the meeting is this week. Meeting agenda is below. Please let me know if you have any questions. For dinners, if you plan to arrive early enough for dinner on Wed (15th), let me know and I can make a reservation and we can go together. I will make a group reservation for the 16th.

Meeting Agenda (Start at 9:00 AM in the Kodiak room at the Seattle Courtyard Marriott, 612 2nd Avenue)

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)

<<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—Fernandes/Abdelnoor  
China—Qiu  
Japan—Ismimoto/Kaga  
Korea—Lee  
United States—Nelson  
CGIAR—Abberton (will provide a PDF/PPT)  
\* 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.

If there are any emergencies, my cell is 765.409.4973

THanks, Scott

On Feb 19, 2015, at 12:19 AM, Scott Allen Jackson <[sjackson@uga.edu](mailto:sjackson@uga.edu)> wrote:

A reminder for those that have not registered yet (~9 people). 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 <<http://www.marriott.com/meeting-event-hotels/group-corporate-travel/group> Corp.mi?resLinkData=Germplasm%20and%20Genomics%20Meeting%20Room%20Block%

5E

seaps%60GGMGMA%60159.00%60USD%60false%604/15/15%604/17/15%603/25/15&app=r

esvlink&stop\_mobi=yes>. 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).

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\* 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/group](https://www.marriott.com/meeting-event-hotels/group-corporate-travel/groupCorp.mi?)  
Corp.mi?

resLinkData=Germplasm%20and%20Genomics%20Meeting%20Room%20Block%  
5E

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esvlink&stop\_mobi=yes