

**From:** [Scott Allen Jackson](#)  
**To:** [Jenny Leverett](#)  
**Subject:** Re: Seattle Meeting, April 15-17  
**Date:** Friday, February 13, 2015 12:18:32 PM

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No. Some aren't coming.

Sent from my iPhone

On Feb 13, 2015, at 9:37 AM, Jenny Leverett <[jkrogersw@gmail.com](mailto:jkrogersw@gmail.com)> wrote:

Scott,

I was looking at the email and noticed that there are more than 13 people on this list. I only have 13 rooms included in the block. Maybe I need to add more?

Jenny

On Thu, Feb 12, 2015 at 4:54 PM, Scott Allen Jackson <[sjackson@uga.edu](mailto:sjackson@uga.edu)> wrote:

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%60GGMGMA%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%60GGMGMA%60159.00%60USD%60false%604/15/15%604/17/15%603/25/15&app=resvlink&stop_mobi=yes)