

From: [Scott Allen Jackson](#)
To: [Jones, Jennifer](#); [Abberton, Michael \(IITA\)](#); [Cannon, Steven \[AGRON\]](#); [Joost, Richard](#); [Nelson, Randall](#); [Okamuro, Jack](#); [Peter Wenzl](#); [Suk-Ha Lee](#); "邱丽娟 OIU, Lijuan"; 李英慧; [Masao Ishimoto](#); [Akito Kaga](#); [Bob Stupar](#); [Ricardo Abdelnoor](#); [Marcelo Fernandes de Oliveira - Embrapa Soja - CNPSO](#); [Steven Cannon](#); [Francois Belzile](#)
Cc: [Jenny Leverett](#)
Subject: Re: Follow up to Seattle soybean meeting
Date: Tuesday, November 17, 2015 1:08:02 PM

Seems there is interest to do this again.

Please let me know if Monday from 10-12 does not work for you. There are no scheduled Workshops at that time.

scott

On Nov 17, 2015, at 9:02 AM, Scott Jackson <sjackson@uga.edu> wrote:

Folks, Sorry for the late date. But would there be interest in having a follow up meeting to the SEattle meeting at PAG (San Diego) this year? If so, please let me know this week and I will find a time and room to do this. I think that Steve Cannon was able to get some funding to get the global inventory started. There may be other updates or items to discuss as well.

Thanks Scott

On Apr 16, 2015, at 10:49 AM, Scott Allen Jackson <sjackson@uga.edu> wrote:

Folks, Michael Abberton was not able to make it. His slides are attached.

<soybeanmeetingapril2015.pptx>

See you at 9 in the Kodiak room.

scott

On Apr 13, 2015, at 1:56 PM, Scott Allen Jackson <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)

* 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—Ismimot0/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> 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](#). 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)
- * 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?resLinkId=Germlasm%20and%20Genomics%20Meeting%20Room%20Block%5Esaps%60GGMGMA%60159.00%60USD%60false%604/15/15%604/17/15%603/25/15&app=resvlink&stop_mobi=yes