Ok. I think I knew that. Scott

Sent from my iPhone

On Feb 19, 2015, at 4:29 PM, Joost, Richard < RJoost@smithbucklin.com> wrote:

I will not be attending as Jennifer Jones will be representing USB.

Rich

Dr. Richard Joost Director of Supply Programs United Soybean Board USB/SmithBucklin 16305 Swingley Ridge Road Chesterfield MO 63017 PH: (314) 579-1590 FAX: [314] 579=1599

Cell: (314) 308-7444 rjoost@smithbucklin.com

<image001.jpg>

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From: Scott Allen Jackson [mailto:sjackson@uga.edu]
Sent: Wednesday, February 18, 2015 11:20 PM
To: Jones, Jennifer: Albedrein, Michael (III1a): Cannon, Steven [AGRON]; Joost, Richard; Nelson, Randall; Okamuro, Jack; Peter Wenzi; Suk-Ha Lee; 邱丽姆 GIU, Lijuan; 李英慧: Masao Ishimoto; Akito Kaga: Bob Stupar; Ricardo Abdelnoor; marcelofernandes.oliveira@embrapa.br; Steven Cannon

Cc: Jenny Leverett Subject: Seattle Meeting, April 15-17

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-

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

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).

-/. • Oposed meeting agenua (uns is a beginning outline. It 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 (<a href="https://www.divseek.org">www.divseek.org</a>)

\* 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

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:

 $https://www.marriott.com/meeting-event-hotels/group-corporate-travel/groupCorp.mi2 \\ resLinkData=Germplasm%20and%20Genomics%20Meeting%20Room%20Block%5Es$ 

:%60GGMGGMA%60159.00%60USD%60false%604/15/15%604/17/15%603/25/15&app=resvlink&stop\_mobi=yes