

From: [Scott Allen Jackson](#)
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Subject: Re: Genomic characterization of world soybean germplasm collections
Date: Wednesday, January 21, 2015 3:38:38 PM

Folks, Please reserve Thursday, April 16 for a meeting. Everyone was available except for Dr. Suk Ha Lee. I hope that he will be able to participate by phone or video.

I am planning on Seattle but will follow up with more details. If you require a visa to visit the US, please let me know now and I will get an invitation letter to you. Also, please confirm by response to this email that you plan to attend so we can make a room block reservation at a hotel.

Again, I'll follow up with more planning details but am planning Seattle probably have a dinner Wednesday (for those that arrive early and are awake), an all day meeting on Thursday and departure Friday.

Thanks, Scott

On Jan 15, 2015, at 9:29 PM, Scott Allen Jackson <sjackson@uga.edu> wrote:

Folks,

I have created a poll (<http://doodle.com/cauzf4hc2mdkybi8>) to find a time to have a meeting to discuss the potential of genomics for soybean germplasm collections.

At this time I am thinking to hold it in Seattle as it is easily accessible with many international flights and we may be able to use the Gates FOundation for meeting space. If anyone has any objections to this, or better suggestions, please let me know ASAP.

I can provide invitation letters, if needed, for visas.

If I have missed anyone on this email. Please let me know and please forward to them.

Thanks, Scott Jackson

On Jan 4, 2015, at 8:21 PM, Scott Allen Jackson <sjackson@uga.edu> wrote:

Dear Colleagues,

I'm Scott Jackson from the University of Georgia in the USA. I'm contacting you as persons that have access to or are in charge of germplasm collections in various countries. You may be away of an international initiative to use

genomics to characterize crop germplasm collections in order to better curate them and to exploit them for crop improvement (www.divseek.org).

I am wondering if there is interest to organize the soybean community to take advantage of this initiative. We would have to find funding source(s) for data generation, but could work hand-in-hand with Divseek to take advantage of infrastructure that will be put together to make use of the genomic data.

In the US, most of the germplasm collection has been genotyped with a 50K SNP chip. However, this has limitations and sequencing would complement the utility of the data to uncover haplotypes and non-SNP variation. In addition, it would be useful to understand redundancy within and between collections and to better understand the variation among the various collections.

If there is interest, I propose that we organize a workshop/meeting. I don't have money to pay for travel, but we could organize it as inexpensively as possible and possibly raise some money to help coordinate it. Topics might include: interest to characterize collections at the sequence level; sequencing approaches; utility in curation; funding; etc...

If I have contacted the wrong person, or if someone is missing, please forward this email to the.

Sincerely,

Scott Jackson

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