

From: [Scott Allen Jackson](#)
To: [Norman Warthmann](#)
Subject: Re: australia - policy brief
Date: Thursday, April 30, 2015 9:32:24 PM

I did my MS on potato:)

I'll check out the paper. I'm way behind in reading...

scott

On Apr 30, 2015, at 9:28 PM, Norman Warthmann

[<norman.warthmann@anu.edu.au>](mailto:norman.warthmann@anu.edu.au) wrote:

Dear Scott,

Thanks a lot.

I am currently sequencing potato for CIP, only 100 for a start. I am advocating low-pass whole genome combined with reduced representation, and I am trying this out right now. Yes, I guess “jumpstarting” is the right word. We will need metadata standards soon. I am trying to re-analyse data from the short read archive and the state of affairs regarding consistent meta information is abhorrent.

Did you see the most recent promising progress on reference graphs?

<http://www.nature.com/ng/journal/vaop/ncurrent/full/ng.3257.html>

best

Norman

On 1 May 2015, at 11:08 , Scott Allen Jackson [<sjackson@uga.edu>](mailto:sjackson@uga.edu) wrote:

Thanks NOrman,

I voted and left comments. I don't know how I missed your brief. It's great. Thanks for taking the effort!

as for Divseek, I'm somewhat ambivalent. We are moving forward on soybean and common bean to genotype by sequencing 10s of thousands of accessions to jumpstart divseek for legumes.

Hope to see you in Australia.

scott

On Apr 30, 2015, at 8:19 PM, Norman Warthmann

[<norman.warthmann@anu.edu.au>](mailto:norman.warthmann@anu.edu.au) wrote:

Dear Scott,

our regular plant science seminar time slot is Wednesdays at 1pm. I have penciled you into the calendar for Wed, November 11, 2015. I can fix that booking anytime in case that suits. If not, I can schedule a special seminar for you any other date and time.

Actually, I was about to contact you anyway to say that I am sorry that you did not make it into the DivSeek steering committee. It wasn't for the lack of votes for you from Canberra. It is particularly annoying that people got preference, who didn't even attend the earlier meetings.

However, on that note. I made a move on my idea about a copyleft-type license for our crop plant genomes. I teamed up with a biodiversity-governance specialist and we submitted a policy brief to the UN. It was in response to a call for briefs about science/policy issues to be included into the "2015 UN Global Sustainable Development Report". It is by no means clear at this point that it will be included. It will depend on the amount of public support it will receive. Hence you may consider rating it (by clicking on the 5th star) and leaving a comment. It would also be good, if you could help to make it widely known. It should serve as a starting point for discussion. I am attaching the pdf also.

<https://gsdr2015.wordpress.com/2015/02/02/thinking-a-global-open-genome-sequence-data-framework-for-sustainable-development>

We also have a press release though:
<http://www.anu.edu.au/news/all-news/united-nations-urged-to-ensure-open-plant-genomes>

best wishes and see you soon hopefully!

best
Norman

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<5934Thinking a global open genome sequence data
framework for sustainable development.pdf>

On 29 Apr 2015, at 11:17 , Scott Allen Jackson
<sjackson@uga.edu> wrote:

Norman,

I'll be in Brisbane 5-8November for a peanut
meeting (in case you are interested:-)).
WOuld be cool to get together if at all
possible.

scott jackson