

From: [Seth C. Murray](#)
To: [Bill Rooney](#)
Cc: [Schuerman, Peter L.](#); [McCutchen, Bill](#); [Avant, Bob](#); [Simpson, Shay](#); [John E" "Mullet](#); [Patricia Klein](#)
Subject: Re: TAMU sweet sorghum study
Date: Tuesday, November 10, 2009 2:32:13 PM

I have no problem sharing the raw data of what we published summaries of (brix, height). There is additional unpublished data on composition (NIRs) and I plan to summarize that for a book chapter, but I do not plan to publish the raw data.

I will default to what others think should be done with it, it obviously has some value.

Thanks,

Seth

----- Original Message -----

From: "Bill Rooney" <wlr@tamu.edu>
To: sethmurray@neo.tamu.edu, "John E" 'Mullet' <jmullet@neo.tamu.edu>, "Patricia Klein" <pklein@tamu.edu>
Cc: "Peter L." 'Schuerman' <PSchuerman@tamu.edu>, "Bill" 'McCutchen' <bmccutchen@tamu.edu>, "Bob Avant" <bavant@tamu.edu>, "Shay" 'Simpson' <shay-simpson@tamu.edu>
Sent: Tuesday, November 10, 2009 2:08:27 PM GMT -06:00 US/Canada Central
Subject: FW: TAMU sweet sorghum study

Greetings:

Please forgive me if we discussed this previously, but I need input from the group per the request from I don't remember if we had a discussion pertaining to Cere's request for phenotype information on sweet sorghum (see below). This would affect some the data that Seth collected as well as some of our current data.

I want to be a good collaborator; at the same time we can just turn everything over for the sake of collaboration. I would welcome your input on what level we should participate and what agreements we make any transfers under. Seth, with regard to your information, I'd like to know if you are even interested in sharing that data.

Regards,

Bill

Dr. William L. Rooney

Professor, Sorghum Breeding and Genetics

Chair, Plant Release Committee

Texas A&M University

College Station, Texas 77843-2474

979 845 2151

From: Timothy Swaller [<mailto:tswaller@ceres.net>]

Sent: Tuesday, November 10, 2009 12:17 PM

To: Bill Rooney

Cc: Jeff Gwyn; Walter Nelson; John Mullet

Subject: TAMU sweet sorghum study

Hi Bill

I am following up on a request that was made a few months back in regards to a population that was phenotyped (NIR, Brix, and height) from 125 diverse accessions and some preliminary marker associations were made (Seth?). Is it possible to get this raw phenotypic information for these 125 accessions (I believe you had mentioned it was going to be available soon)?

We would like to start looking at these types of datasets to begin developing a better comprehensive understanding of these types of studies and the utility they may have for our internal and/or joint programs. Also, this will help us to better understand the benefits and weaknesses of these approaches.

Thanks

Tim

Timothy Swaller

Director, IT and Genomics

Office: 805.376.6545

tswaller@ceres.net

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Seth C. Murray

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From: [Schuerman, Peter L.](#)
To: [Bill Rooney](#); sethmurray@neo.tamu.edu; "Mullet, John E"; "Patricia Klein"
Cc: [McCutchen, Bill](#); [Avant, Bob](#); [Simpson, Shay](#)
Subject: RE: TAMU sweet sorghum study
Date: Tuesday, November 10, 2009 2:30:07 PM

Does turning the information over mean that they need us less? Does retaining it create an opportunity for collaboration?

From: Bill Rooney [<mailto:wlr@tamu.edu>]
Sent: Tuesday, November 10, 2009 2:08 PM
To: sethmurray@neo.tamu.edu; 'Mullet, John E'; 'Patricia Klein'
Cc: Schuerman, Peter L.; McCutchen, Bill; Avant, Bob; Simpson, Shay
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Cc: [Schuerman, Peter L.](#); [McCutchen, Bill](#); [Avant, Bob](#); [Simpson, Shay](#)
Subject: FW: TAMU sweet sorghum study
Date: Tuesday, November 10, 2009 2:08:54 PM

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