

From: [Greg Porreca](#)
To: [Young, Dan](#)
Cc: [Waller, Patrick](#)
Subject: Re: new app
Date: Friday, May 15, 2009 8:29:12 AM
Attachments: [G0814.70002US00 - PROVISIONAL PT APP-1 GP-EDITS.DOC](#)

Hi Dan-

Attached is my edited version. I used track changes.

Note I added a short paragraph in attempt to cover us if we find our initial barcode pool is not totally uniform random. If this happens, we may want to modify the claims down the road by including a calibration step to correct for it. (as written I think we're limited to using ****only**** unique combinations of target:source in the genotyping, but if there was barcode bias present, and we understood it well enough, we could tolerate more than 1 of ****some**** combinations of target:source depending on abundance in the initial pool).

-Greg

On Thu, May 14, 2009 at 10:29 PM, Daniel Young <Daniel.Young@wolfgreenfield.com> wrote:

Thanks, Greg.

Attached is a draft of Uri's application. We are still thinking about the term "source sequence", and may change it. But please review this draft and let us know your comments. I have included some specific questions in bold.

Thanks
Dan

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

From: Greg Porreca [mailto:gporreca@gsgenetics.com]
Sent: Thursday, May 14, 2009 5:13 PM
To: Young, Daniel W.
Cc: Waller, Patrick
Subject: Re: new app

Bob's address:

Robert Wassman
90 West Shore Drive
Marblehead, MA 01945

On Thu, May 14, 2009 at 4:27 PM, Daniel Young
<Daniel.Young@wolfgreenfield.com> wrote:

Hi Greg,

We will call you in ten minutes to discuss this new application.

Thanks
Dan

From: Greg Porreca [mailto:gporreca@gsogenetics.com]
Sent: Thursday, May 14, 2009 1:30 PM
To: Waller, Patrick; Young, Daniel W.
Subject: new app

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