

From: Lau, Matthew K. matthewklau@fas.harvard.edu  
Subject: Re: sequence transfer  
Date: November 7, 2016 at 4:17 PM  
To: Jim Bochicchio jboch@broadinstitute.org  
Cc: Jennifer Wineski jwineski@broadinstitute.org, Sarah Young stowey@broadinstitute.org, Caroline Cusick ccusick@broadinstitute.org

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Hi Jim, that works for me. I just saw & accepted the updated meeting invite too.

Best,  
Matt

On Nov 7, 2016, at 2:18 PM, Jim Bochicchio <[jboch@broadinstitute.org](mailto:jboch@broadinstitute.org)> wrote:

Hi Matt

Can we bump our meeting until the 18th? We had a weekly check in with our analyst today and he doesn't think all of the assemblies will be ready this week.

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The data will be stored here for a month before being deleted, so please move it over at your earliest convenience. If you have any questions or concerns, please email [btlhelp@broadinstitute.org](mailto:btlhelp@broadinstitute.org)

Aspera Username: SN0107253  
Aspera Password:  
B26QN2YENHKT29B  
Aspera FilePasscode: 6miXig7JX19I91f

For detailed instructions, examples and  
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From: Lau, Matthew K. matthewklau@fas.harvard.edu

Subject: Re: sequence transfer

Date: November 2, 2016 at 1:14 PM

To: Jennifer Wineski jwineski@broadinstitute.org

Cc: Sarah Young stowey@broadinstitute.org, Jim Bochicchio jboch@broadinstitute.org, Caroline Cusick ccusick@broadinstitute.org



Hi Jenn, Thu Nov 10 at 11am works for me!

Best,  
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75 Ames St | Cambridge MA 02142 USA

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Caroline Cusick  
Product Coordinator  
Broad Institute of MIT and Harvard  
75 Ames St | Cambridge MA 02142 USA



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rCollab...tor.xlsx



From: Lau, Matthew K. matthewklau@fas.harvard.edu  
Subject: Re: sequence transfer  
Date: October 27, 2016 at 4:01 PM  
To: Caroline Cusick ccusick@broadinstitute.org  
Cc: Jim Bochicchio jboch@broadinstitute.org, Sarah Young stowey@broadinstitute.org

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Hi Caroline, great, thanks for the update and the heads-up!

Cheers,

Matt

On Oct 27, 2016, at 10:36 AM, Caroline Cusick  
<[ccusick@broadinstitute.org](mailto:ccusick@broadinstitute.org)> wrote:

Hi Matt,

The Aspera is still staging, but it should come down either today/tomorrow, and I'll send along the login information. As a heads up, the total size of the raw files is ~0.69 TB.

thanks!  
Caroline

On Thu, Oct 27, 2016 at 10:23 AM, Lau, Matthew K.  
<[matthewklau@fas.harvard.edu](mailto:matthewklau@fas.harvard.edu)> wrote:

Hi Caroline and Sarah, just checking in the status of the sequences. Did your bioinformatician work his magic? And/or, have the files been zipped and hosted on your file transfer system?

Best,

Matt

*Postdoctoral Fellow*  
Harvard Forest  
Harvard University  
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Petersham, MA 01366  
[978-756-6165](tel:978-756-6165)

"Knowledge is knowing a tomato is a fruit.  
Wisdom is not putting it in a  
fruit salad." — Miles Kingston

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