## Zimbra

# Re: Data sharing and collaboration

**De:** Florian Rambow <florian.rambow@kuleuven.be>

lun., 15 août 2022 22:30

5 pièces jointes

Objet: Re: Data sharing and collaboration

À: Romain Larive < romain.larive@umontpellier.fr>

Les images externes ne seront pas affichées. Montre les images ci-dessous

Hi Romain,

sorry for my late reply. Things are still a bit chaotic here.

Attached you can find a ppt with explanations for the different data sets and related .xls files, also the supp. table of our paper (mmc2.xls). I hope these files are helpful.

In summary, we never reanalyzed DNA-seq (WES...) from scratch for our paper. We rather relied on the mutational calling of the different cohorts/papers.

If you are interested in the gene expression data of the different cohorts, I can share them via belnet.filesender.

With regards to the PDX cohort from the Wistar Institute, please find the original paper attached. They basically continued with their collection and deepened the molecular profiling.

I was asked to collaborate on the molecular profiling of this massive melanoma PDX cohort. To date, I haven't heard back from them.

If you are eager to contact Helen Rizos again, just forward me your request and I will try to weigh-in.

Amicalement

Florian

p.s. from now on you could also use: <a href="mailto:florian.rambow@uk-essen.de">florian.rambow@uk-essen.de</a> as I will check this Email more often.

On 11/08/2022 08:00, Romain Larive wrote:

Hello Florian,

I hope that the installation in your new laboratory could be completed correctly.

Following our discussion by Zoom at the beginning of July concerning our collaboration for the sharing of your clinical and biological data, have you been able to take stock of the data in your possession? And did you eventually contact the laboratories that generated them to complete them?

I am also interested in the genomic and transcriptomic data generated from PDX in the USA. What are the conditions under which these data would be available and how long would it take?

Concerning the modalities of collaboration, I propose to start from the principle that our publications mentioning the data that you will provide us will give rise to your presence among the authors. The question I have concerns the laboratories from which the data originally came. What do you think about it?

Best regards,

Romain

**De:** "Florian Rambow" <a href="mailto:slope-"><a href="mailto:slope-">slope-"><a href="mailto:slope-"><a href="mailto:slope-">slope-"

**Envoyé:** Mercredi 6 Juillet 2022 11:43:10 **Objet:** Re: Data sharing and collaboration

Cool. Talk to you then.

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On 06/07/2022 10:15, Romain Larive wrote:

Hello Florian,

No problem.

Ok for the 13th (14th is free in France).

10h via Zoom?

https://umontpellier-fr.zoom.us/j/3708678545?pwd=cmhKY3NOenR0ano2OGdGSVA1T0xpZz09

ID de réunion : 370 867 8545

Code secret: 300962

Amicalement,

Romain

**De:** "Florian Rambow" <a href="mailto:slope-be/"><a href="mailto:slope-be/">mailto:slope-be/<a href="mailto:slope-be/"><a href="mailto:slope-be/">mailto:slope-be/<a href="mailto:slope-be/">mai

**Envoyé:** Mardi 5 Juillet 2022 21:53:23 **Objet:** Re: Data sharing and collaboration

Hi Romain,

I am still keen to discuss our collaboration.

However, could we reschedule to next week either 13th or 14th?

Sorry for these last minute changes.

Amicalement

Florian

On 05/07/2022 14:27, Romain Larive wrote:

Hello Florian,

Still OK to discuss via Zoom our collaboration for sharing clinical and biological data?

Sarah, the PhD student working on this project, will be able to present the database she developed to manage the data we have already retrieved.

We will be available all day tomorrow, Thursday morning or Friday morning.

Best regards,

#### Romain

De: "Romain Larive"

<romain.larive@umontpellier.fr>

À: "Florian Rambow"

<florian.rambow@kuleuven.be>

**Envoyé:** Lundi 27 Juin 2022 11:13:05 **Objet:** Re: Data sharing and collaboration

Hello Florian,

Great!

That's right, I would like to reanalyze all the RNAseq and clinical data from the different centers, but also the DNAseq.

A zoom call between the 6th and 8th at noon is fine with me. Just let me know what time would be convenient for you.

I don't know if you are French speaking. If you prefer, we could switch to French.

Best,

Romain

De: "Florian Rambow"

<florian.rambow@kuleuven.be>

À: "Romain Larive"

<romain.larive@umontpellier.fr>

**Envoyé:** Vendredi 24 Juin 2022 20:38:01 **Objet:** Re: Data sharing and collaboration

Hi Romain,

good to hear from you. Your DL approach sounds tempting.

I am happy to share the data of our Cancer Cell paper. I suppose, you are referring to the bulk RNAseq and clinical data of the different centers (Boston, Sydney...)?

How about we set up a zoom call in the week 4th-8th of July to discuss a bit further?

Cheers

Florian

On 24/06/2022 17:21, Romain Larive wrote:

Dear Dr Rambow,

I am Romain Larive, associate professor at the Montpellier Cancer Institute in France.

Our group, in the <u>research team</u> led by Peter Coopman, is interested in the mechanisms of resistance of melanoma cells to kinase inhibitors.

I would like to develop a deep learning approach to analyze clinical and biological data from patients treated with kinase inhibitors. To do this, I would like to use the data described in your paper Marin-Bejar et al, Cancer Cell 2021. I would like to exploit the clinical data (at least the type of treatment, PFS and OS) and the biological data (at least pretreatment WES, if possible post-treatment WES).

It would be an honor to collaborate with you on this data sharing. I am at your disposal for more precise information about our project and to discuss the modalities of collaboration, including authorship.

I am contacting you on the advice of Chris Marine, who told me that you are very busy with the installation of your new lab in Essen. Do not hesitate to tell me to contact you later.

Sincerely,

Romain Larive

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#### **ROMAIN LARIVE**

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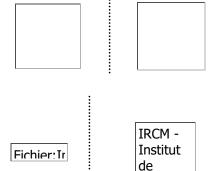
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