

Zimbra

romain.larive@umontpellier.fr

Re: Collaboration proposal

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

ven., 11 mars 2022 16:10

Objet : Re: Collaboration proposal 3 pièces jointes**À :** Yibing Yan <yan.yibing@gene.com>**Cc :** aribas <ARibas@mednet.ucla.edu>, Matthew Wong
<wongchenko.matthew@gene.com>

Dear Yibing,

Thank you very much for the RNA-Seq table, we can now relate the expression results to the patients.

On the other hand, the ExomeSamples table does not allow us to know the mutations detected by WES.

We would therefore need the analyzed WES data (the table of detected mutations) and the correspondence with the patients. In your 2019 article, you show in figure 1 the genetic changes on a small panel of genes. We would need to know all detected mutations (SNV and CNV). We would also like to retrieve the raw data from WES (in a second step).

Kind regards,

Romain

De: "Yibing Yan" <yan.yibing@gene.com>**À:** "Romain Larive" <romain.larive@umontpellier.fr>**Cc:** "Yibing Yan" <yan.yibing@gene.com>, "aribas" <ARibas@mednet.ucla.edu>, "Matthew Wong" <wongchenko.matthew@gene.com>**Envoyé:** Jeudi 10 Mars 2022 19:27:03**Objet:** Re: Collaboration proposal

Hi Romain,

Here are the lists that link the patients with either exome or RNAseq data.

Best,
yibing

On Thu, Mar 10, 2022 at 7:02 AM Romain Larive <romain.larive@umontpellier.fr> wrote:

Dear Yibing,

Thank you for your answer, I understand.

We are waiting for your list of detected mutations (SNV and CNV) to advance our study. Would you be able to send it to us directly?

Kind regards,

Romain

De: "Yibing Yan" <yan.yibing@gene.com>**À:** "Romain Larive" <romain.larive@umontpellier.fr>

Cc: "Yibing Yan" <yan.yibing@gene.com>, "aribas" <ARibas@mednet.ucla.edu>, "Matthew Wong" <wongchenko.matthew@gene.com>

Envoyé: Jeudi 10 Mars 2022 15:56:54

Objet: Re: Collaboration proposal

Hi Romain,

I was out for a period of time; and we are also having some issues on exporting raw data in Roche since I do not have direct access to them.

I would send the ID's linking the patient with existing data first, and will try to post the raw data to the journal.

Apologize for the delay.

yibing

On Thu, Mar 10, 2022 at 5:57 AM Romain Larive <romain.larive@umontpellier.fr> wrote:

Dear Professor Ribas, dear Yibing Yan,

I would like to reiterate my proposal for collaboration regarding the sequencing data that you have already published in your article (Yan et al., Clinical Cancer Research 2019), but which are not available on the journal's website.

Following the positive response from Prof. Ribas on 01/28/2022, I clarified to Yibing Yan what we would need.

I am aware of the limited time you can devote to my request. Nevertheless, as this is already published data, would it be possible for you to share it? I am at your disposal to discuss the modalities of collaboration.

This help from your part would allow us to advance in the realization of our project.

Yours sincerely,

Romain Larive

De: "Romain Larive" <romain.larive@umontpellier.fr>

À: "Yibing Yan" <yan.yibing@gene.com>

Cc: "Matthew Wong" <wongchenko.matthew@gene.com>, "aribas" <ARibas@mednet.ucla.edu>

Envoyé: Mercredi 2 Février 2022 17:24:01

Objet: Re: Collaboration proposal

Hi Yibing,

WES RAW data:
FASTQ, SAM or BAM file format would be good.

WES analyzed data:
I attached a TSV file as an example.

RNA-Seq:
The correspondence between Table S1 (patients and clinical data) to Table S2 (Raw sequencing data) is missing.

Best,

Romain

De: "Yibing Yan" <yan.yibing@gene.com>
À: "Romain Larive" <romain.larive@umontpellier.fr>
Cc: "Yibing Yan" <yan.yibing@gene.com>, "Matthew Wong" <wongchenko.matthew@gene.com>, "aribas" <ARibas@mednet.ucla.edu>

Envoyé: Mercredi 2 Février 2022 16:25:10

Objet: Re: Collaboration proposal

Hi Romain,
We'll try to get the data to you soon.

Could you let us know what form of "raw" data you are looking for?

Best,
yibing

On Wed, Feb 2, 2022 at 4:53 AM Romain Larive

<romain.larive@umontpellier.fr> wrote:

Dear Professor Ribas, Yibing Yan and Matt Wongchenko,

Thank you for your positive response.

We would need the whole exome and RNA sequencing data that you have already published in your paper (Yan et al., Clinical Cancer Research 2019), but which are not available on the journal website.

Specifically, we would need:

- RNA-Seq: the mapping of data from Table S1 (patients and clinical data) to Table S2 (Raw sequencing data);
- WES: the raw and analyzed data (the table of detected mutations) and the patient correspondence.

This help from your part would allow us to advance in the realization of our project.

Yours sincerely,

Romain Larive

De: "aribas" <ARibas@mednet.ucla.edu>
À: "Romain Larive" <romain.larive@umontpellier.fr>, "Yibing Yan" <yan.yibing@gene.com>, "Matthew Wong" <wongchenko.matthew@gene.com>

Envoyé: Samedi 29 Janvier 2022 02:00:53

Objet: RE: Collaboration proposal

Hi Romain, I am sorry for misunderstanding. I am responding with cc to my colleagues from Genentech Yibing Yan and Matt Wongchenko, who generated this data. Best, Toni

From: Romain Larive <romain.larive@umontpellier.fr>
Sent: Friday, January 28, 2022 9:28 AM
To: Ribas, Antoni <ARibas@mednet.ucla.edu>
Subject: Re: Collaboration proposal

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Dear Professor Ribas,

Thank you for your answer, even if it was negative.

It is deeply unfortunate. We would only need the sequencing data that you have already published in your paper (Yan et al., Clinical Cancer Research 2019), but which are not available on the journal's website.

I am well aware that you will not have the time to devote to me, nevertheless I would like to suggest that this could be done by someone from your research team.

This help from you would allow us to move forward in the realization of our project.

Yours sincerely,

Romain Larive

De: "aribas" <ARibas@mednet.ucla.edu>
À: "Romain Larive" <romain.larive@umontpellier.fr>
Envoyé: Jeudi 27 Janvier 2022 17:29:03
Objet: RE: Collaboration proposal

Thanks but I am not interested. Best, Toni

From: Romain Larive <romain.larive@umontpellier.fr>
Sent: Thursday, January 27, 2022 8:05 AM
To: Ribas, Antoni <ARibas@mednet.ucla.edu>
Subject: Collaboration proposal

CAUTION - EXTERNAL EMAIL: Do not click links or open attachments unless you recognize the sender.

Dear Professor Ribas,

I am Romain Larive, associate professor at the University of Montpellier in France. [Our research team](#), led by Peter Coopman in the INSERM Institute of Cancer Research of Montpellier, 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 of patients treated with kinase inhibitor(s). For this I would like to use the data described in these article you are corresponding author: Yan et al., Clinical Cancer Research 2019.

It would be an honor to collaborate with you with this data sharing. I am at your disposal to discuss the modalities of collaboration.

Sincerely,

Romain Larive

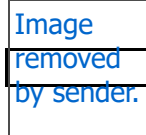
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ROMAIN LARIVE
MAÎTRE DE CONFÉRENCES EN TOXICOLOGIE
DOCTEUR / HDR EN BIOLOGIE - SANTÉ

Laboratoire de Toxicologie du Médicament - Faculté des Sciences Pharmaceutiques et Biologiques de Montpellier

Équipe Signalisation de l'Invasion Tumorale - Institut de Recherche en Cancérologie de Montpellier

romain.larive@umontpellier.fr



FACULTÉ DES SCIENCES PHARMACEUTIQUES ET BIOLOGIQUES
Laboratoire de Toxicologie du Médicament - Bâtiment K - 1er étage

15 avenue Charles Flahault - CS 14491
F-34093 Montpellier Cedex 5, FRANCE

Tél : +33 (0)4 11 75 97 50 | Fax : +33 (0)4 11 75 97 59

WWW.PHARMACIE.EDU.UMONTPELLIER.FR

INSTITUT DE RECHERCHE EN CANCÉROLOGIE DE MONTPELLIER

Campus Val d'Aurelle (U1194)

208 avenue des Apothicaires
F-34298 Montpellier cedex 5, FRANCE

Tél : +33 (0)4 11 28 31 04 | Fax : +33 (0)4 67 61 37 87

WWW.IRCM.FR

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ROMAIN LARIVE
MAÎTRE DE CONFÉRENCES EN TOXICOLOGIE
DOCTEUR / HDR EN BIOLOGIE - SANTÉ

Laboratoire de Toxicologie du Médicament - Faculté des Sciences Pharmaceutiques et Biologiques de Montp

Équipe Signalisation de l'Invasion Tumorale - Institut de Recherche en Cancérologie de Montpellier

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	:	FACULTÉ DES SCIENCES PHARMACEUTIQUES ET BIOLOGIQUES
	:	Laboratoire de Toxicologie du Médicament - Bâtiment K - 1er étage
	:	15 avenue Charles Flahault - CS 14491
	:	F-34093 Montpellier Cedex 5, FRANCE
	:	
	:	
	:	
	:	

Tél : +33 (0)4 11 75 97 50 | Fax : +33 (0)4 11 75 97 59
WWW.PHARMACIE.EDU.UMONTELLIER.FR

	:	INSTITUT DE RECHERCHE EN CANCÉROLOGIE DE MONTPELLIER
	:	
	:	Campus Val d'Aurelle (U1194)
	:	
	:	208 avenue des Apothicaires
	:	F-34298 Montpellier cedex 5, FRANCE
	:	
	:	
	:	
	:	

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WWW.PHARMACIE.EDU.UMONTPELLIER.FR



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