

To: [REDACTED]@gelderland.nl]  
Cc: [REDACTED]@gmail.com]  
From: [REDACTED]  
Sent: Wed 9/6/2023 10:13:42 AM  
Subject: Re: Hybrid detection - a few details  
Received: Wed 9/6/2023 10:14:23 AM

Dear [REDACTED]  
Thank you for your reply. 13th of September is fine, we'll prepare a quotation.

Best,

[REDACTED]  
[REDACTED]  
Raziskovalec / *Researcher*  
Oddelek za biologijo / *Biology Department*  
Biotehniška fakulteta / *Biotechnical Faculty*  
Jamnikarjeva ulica 101, SI-1000 Ljubljana, Slovenija / *Slovenia*  
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[REDACTED]@gmail.com, [www.bf.uni-lj.si](http://www.bf.uni-lj.si)

On Wed, Sep 6, 2023 at 11:05 AM [REDACTED]@gelderland.nl> wrote:

Dear [REDACTED]

Thank you for your considerations. We gave it some more thoughts and decided to stick with the plan. I realize this also means there is some loss of cost efficiency, but we will take this in consideration when judging your quotation. This also means we would like to stick to the 13<sup>th</sup> of September, please let me know if this is ok with you.

Greeting [REDACTED]

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Van: [REDACTED]@gmail.com>  
Verzonden: zondag 3 september 2023 12:45  
Aan: [REDACTED]@gelderland.nl>; [REDACTED]@gmail.com>  
Onderwerp: Hybrid detection - a few details

Dear [REDACTED]

While preparing a quote we have been asking around for pricing, and the smallest batch size we can get for genome-level SNP genotyping is 48 samples. We are trying to find other people to share the costs of the batch with their samples, but I'm not sure we'll be able to find any. I have a couple of questions:

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- Are you ok to wait for a week or two for us to try and find people to share the SNP genotyping batch? If we manage, we can make the final price 3000 - 4000 € cheaper. If not, I will send you the quote in a day or two - let me know.

- If we can't find other people to share the cost (which is highly likely), I would suggest including other samples in the analysis. This will not increase the price at all, and the results will be better. We can include individuals from various populations that are relevant for your case study, but what would be interesting is if you could include some other samples from your side, blind to us (so that we don't know what they are). A couple of dogs, some jackals if you have them, any additional wolves, maybe some zoo wolves? Any known hybrids would be excellent. These would of course need to be tissue samples.

Best regards,

[REDACTED]

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[REDACTED]

*Raziskovalec / Researcher*

*Oddelek za biologijo / Biology Department*

*Biotehniška fakulteta / Biotechnical Faculty*

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