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In a series of posts tagged with [ideas-for-genomics-project](http://scalaakka.blogspot.com/search/label/ideas-for-genomics-project) [*] I am describing ideas for business that can be build in area of [genomics](https://en.wikipedia.org/wiki/Genomics) . If you like those ideas and you would like to join as an investor, co-founder or consultant please contact me at [LinkedIn](https://pl.linkedin.com/pub/artur-stanek/a5/643/691) or just drop me an email, stanek.artur@gmail.com .

Now you may be interested to check the [big picture](http://scalaakka.blogspot.com/2015/10/ideas-for-genomics-project-diagram-of.html) of the whole idea.

* - please read the list on the left from the first post (which is at the bottom)

Artur Stanek - Scala, Akka dev with passion.

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[\[posts: written by me\]](#)
[\[posts: web finds\]](#)
[\[posts: all\]](#)

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Ideas for genomics project: genome meta information...

Ideas for genomics project: option on your own geno...

Ideas for genomics project: check if your genome is L...

Ideas for genomics project: grand permissions to you...

Ideas for genomics project: overcome legal limitation...


Ideas for genomics project: anonymise me behind m...

Ideas for genomics project: sharing genome informat...


Ideas for genomics project: signing and identifying ge...

Ideas for genomics project: filter out what is not need...

Ideas for genomics project: store and manage genom...

 Ideas for genomics project: SafeGenome - introductio...

Ideas for genomics project: introduction to the series.




Ideas for genomics project: SafeGenome - introduction and diagram.

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At October 2015 I attended Roche's Code4Life conference. Roche is planning to provide analysis services over genome file but - as they admitted - they are not interested to host that files. I understand that hosting business is more difficult that 'just storing files in the cloud' as we need to deal with - for example - legal limitations.

Year before, when I was working for bioinformatics company in beautiful Los Angeles, I initiated few discussions about moving their business to the cloud. What I discovered that genome safety and privacy is so crucial for them and their clients that going to cloud is almost impossible.

I am not sure if there are so big stoppers that we can not go into the Web. As I mentioned in "Genome - the new intimacy", new tools needs to be provided to allow both sides (genome owners and companies that want to provide services over them) to be satisfied.



Sequencing cost is rapidly going down (above photo was taken at Roche's Code4Life conference). In 2015 it was near \$1000 per genome and going down. Once it will go below critical point people will scan themselves on a mass scale. Where will they keep their scans? Will it be safe for them? Will companies that provides analysis solve this situation?

[<https://1.bp.blogspot.com/-rBL1gW1JUQg/VzG0cRDBful/AAAAAAB-Vg/QmQ-IULgkm8IKzUBKJPYbKJ-t5UvSpxVQCLcB/s1600/2016-05-10%2BArtur%2BStanek%2Bblog%2B-%2Bideas-for-genomics-project1.png>]

