

## Ideas for genomics project: introduction to the series.

As I am very active in inventing 101 ideas for business each day I should start to write them down long time ago. But anyway, it is never too late to start ... so let me start :).

In a series of posts tagged with [ideas-for-genomics-project](http://scalaakka.blogspot.com/search/label/ideas-for-genomics-project) [http://scalaakka.blogspot.com/search/label/ideas-for-genomics-project] I will describe ideas for business that can be build in area of 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) [https://pl.linkedin.com/pub/artur-stanek/a5/643/691] or just drop me an email, [stanek.artur@gmail.com](mailto:stanek.artur@gmail.com) [mailto:stanek.artur@gmail.com] .

This post is just an introduction to the series. I wonder where I can use my passion and - what's the most important - where the power of [Scala](http://www.scala-lang.org/) [http://www.scala-lang.org/] programming language (and its [ec](http://akka.io/) [http://akka.io/] [osy](https://github.com/gearpump/gearpump) [https://github.com/gearpump/gearpump] [ste](http://spark.apache.org/) [http://spark.apache.org/] [m](#)) can be used in genomics field.

Just to list some of Scala's strengths and features:

- ([Scala](http://www.scala-lang.org/) [http://www.scala-lang.org/]) power of the compiler, strong and rich type system, macros, compiler plugins,
- ([Scala](http://www.scala-lang.org/) [http://www.scala-lang.org/]) internal and external DSLs, powerful and easy to use parser combinators,
- ([Akka](http://akka.io/) [http://akka.io/]) reactive streams (balanced and safe way for parallel processing in a from of graphs with automatic demand/back pressure management that can be - potentially - materialized not only on classical CPUs),
- ([Akka](http://akka.io/) [http://akka.io/]) reactive paradigm, actors model, state machines (FSM), dynamic clusters, parallel processing, conflict free replicated data types, localization transparency, self healing, Akka Typed,
- ([Apache Spark](http://spark.apache.org/) [http://spark.apache.org/], [GearPump](https://github.com/gearpump/gearpump) [https://github.com/gearpump/gearpump], [Apache Flink](https://flink.apache.org/) [https://flink.apache.org/]) massive, parallel processing,
- ([Scala](http://www.scala-lang.org/) [http://www.scala-lang.org/]) value classes / universal traits, lazy streams.

Next post in this series: "SafeGenome - introduction and diagram [http://scalaakka.blogspot.com/2015/10/ideas-for-genomics-project-diagram-of.html]" .

Posted 30th October 2015 by [Artur Stanek \(kermidas\)](#)

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