



# EuroFAANG

Accelerating genome to phenome research  
for farmed animals in Europe

EuroFAANG is a coordinated effort to unravel the connection between the genetic make-up of an animal and the observable physical and physiological traits. The EuroFAANG projects aim to address challenges in farmed animal production.

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Bringing together a wide range of genomics, bioinformatics, modelling and open data expertise, the six projects are laying the scientific foundations for a new era of farmed animal production based on:

Disease resistance

Biological efficiency

Precision breeding

Reduced environmental impact

Feeding a growing population

**The EuroFAANG community is supported by  
six key projects**



Document genome function to understand the basis for trait variation and disease resistance in farmed fish.



[www.aqua-faang.eu](http://www.aqua-faang.eu)  
@AQUA\_FAANG



Identify genome features for phenotypic diversity in cattle.



[www.bovreg.eu](http://www.bovreg.eu)  
@BovReg



Develop new breeding strategies to help ruminants adapt to climatic changes.



[www.rumigen.eu](http://www.rumigen.eu)  
@RUMIGENH2020



Understanding microbiomes of the ruminant holobiont.



[www.holoruminant.eu](http://www.holoruminant.eu)  
@holoruminant



Identify genome features whose activity, during development and when facing environmental challenges, determines complex traits in chicken and pigs.



[www.gene-switch.eu](http://www.gene-switch.eu)  
@GeneSwitch



Provide new knowledge and tools for genome and epigenome enabled breeding in monogastrics.



[www.geronimo-h2020.eu](http://www.geronimo-h2020.eu)  
@GeronimoH2020

Visit the project websites to find the latest tools and resources for the breeding and livestock farming communities. Project data will be available on the FAANG Data Portal powered by EMBL's European Bioinformatics Institute (EMBL-EBI).



<https://data.faang.org/projects>

<https://eurofaang.eu>

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These projects have received funding from the European Union's Horizon 2020 Research and Innovation Programme under the Grant agreements no 817923, 8156668, 817998, 101000236 and 101000226.



**EuroFAANG**