



RESEARCH  
PROGRAM ON  
Livestock

*More meat, milk and eggs by and for the poor*

# Introduction to bioinformatics and population genomics applied to livestock

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*Patron: Professor Peter C Doherty AC, FAA, FRS*

*Animal scientist, Nobel Prize Laureate for Physiology or Medicine—1996*

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*better lives through livestock*

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Name of the training event	<b>Introduction to bioinformatics and population genomics applied to livestock</b>																																																																											
Aims of the course	The introduction to Bioinformatics and population genomics applied to livestock free online course aims to provide early-career scientists with the theoretical knowledge and practical skills to utilize genetic data to address research questions in livestock genomics and breed characterisation improvement.																																																																											
Dates	May 24 to June 4 2021																																																																											
Venue	The live course sessions took place via the Zoom meeting app. The SLU canvas was the learning platform where course materials, videos and assignments were accessible to participants																																																																											
Instructors and their affiliation	Erik Bongcam Rudloff, Swedish University of Agricultural Sciences (SLU) Anna Maria Johansson, Swedish University of Agricultural Sciences (SLU) Renaud Van Damme, Swedish University of Agricultural Sciences (SLU) Yuri Utsunomiya, Sao Paulo State University (UNESP - Brazil)																																																																											
Participant information, including a number by gender	Forty-two participants, comprising 16 females and 26 males, were admitted to the course. Thirty-three participants (33% female, 67% male) from 20 countries in sub-Saharan Africa (Algeria, Benin, Burkina Faso, Congo, Côte d'Ivoire, Egypt, Ethiopia, Ghana, Kenya, Malawi, Morocco, Nigeria, Rwanda, Senegal, Somalia, South Africa, Sudan, Tanzania, Tunisia and Zimbabwe) completed the courses.																																																																											
Funder	The Livestock CGIAR research program (Livestock CRP) and Global Impact Accelerator Award From the University of Edinburgh																																																																											
Course summary or agenda	<table border="1"> <thead> <tr> <th>Date</th> <th>Time (UTC+2)</th> <th>Topic</th> </tr> </thead> <tbody> <tr> <td>24 May</td> <td>13h-13h30</td> <td>Welcome</td> </tr> <tr> <td>May 24</td> <td>13h30-13h45</td> <td>Practical Information on asynchronous Basic genomic and Rsessions</td> </tr> <tr> <td>May 24</td> <td>13h45-15h</td> <td>What is bioinformatics and genomics Lecture</td> </tr> <tr> <td>May 25</td> <td>13h-14h</td> <td>Bioinformatics resources and biological databases Lecture</td> </tr> <tr> <td>May 25</td> <td>14h-15h</td> <td>Bioinformatics resources and biological databases Tutorial and List</td> </tr> <tr> <td>May 26</td> <td>13h-14h</td> <td>Basics of Bioinformatics using Emboss software suite Lecture</td> </tr> <tr> <td></td> <td>download</td> <td></td> </tr> <tr> <td>May 26</td> <td>14h-15h</td> <td>Basics of Bioinformatics using Emboss software suite Tutorial</td> </tr> <tr> <td>May 27</td> <td>13h-14h</td> <td>Introduction to Linux Lecture download</td> </tr> <tr> <td>May 27</td> <td>14h-15h</td> <td>Linux tutorials and exercises</td> </tr> <tr> <td>24-27 May</td> <td></td> <td>Introduction to R Lecture</td> </tr> <tr> <td>24-27May</td> <td></td> <td>R tutorial and exercises</td> </tr> <tr> <td>24-27 May</td> <td></td> <td>Introduction to population genetics Lecture</td> </tr> <tr> <td>May 28</td> <td>13h-14h</td> <td>Question Time about R</td> </tr> <tr> <td>May 28</td> <td>14h-15h</td> <td>Basic Manipulation of High-density SNP data Lecture and tutorial</td> </tr> <tr> <td>May 31</td> <td>13h-13h30</td> <td>Introduction to NGS technologies Lecture download</td> </tr> <tr> <td>May 31</td> <td>13h30-15h</td> <td>Working on NGS data and File Formats with Ugene Tutorial</td> </tr> <tr> <td>June 1</td> <td>13h-14h</td> <td>Working on NGS data and File Formats with Ugene Tutorial</td> </tr> <tr> <td>June 1</td> <td>14h-15h</td> <td>Theory on signatures of selection Lecture</td> </tr> <tr> <td>June 2</td> <td>13h-14h</td> <td>Theory on ancestry analyses Lecture</td> </tr> <tr> <td>June 2</td> <td>14h-15h</td> <td>Practice demonstrating a selection signatures analysis with cattle data</td> </tr> <tr> <td>June 3</td> <td>13h-15h</td> <td>Practice demonstrating a breed composition analysis with cattle data</td> </tr> <tr> <td>June 4</td> <td>13h-13h20</td> <td>Access and Benefit Sharing of genetic resources</td> </tr> <tr> <td>June 4</td> <td>13h20-14h30</td> <td>Q&amp;A all material</td> </tr> </tbody> </table>	Date	Time (UTC+2)	Topic	24 May	13h-13h30	Welcome	May 24	13h30-13h45	Practical Information on asynchronous Basic genomic and Rsessions	May 24	13h45-15h	What is bioinformatics and genomics Lecture	May 25	13h-14h	Bioinformatics resources and biological databases Lecture	May 25	14h-15h	Bioinformatics resources and biological databases Tutorial and List	May 26	13h-14h	Basics of Bioinformatics using Emboss software suite Lecture		download		May 26	14h-15h	Basics of Bioinformatics using Emboss software suite Tutorial	May 27	13h-14h	Introduction to Linux Lecture download	May 27	14h-15h	Linux tutorials and exercises	24-27 May		Introduction to R Lecture	24-27May		R tutorial and exercises	24-27 May		Introduction to population genetics Lecture	May 28	13h-14h	Question Time about R	May 28	14h-15h	Basic Manipulation of High-density SNP data Lecture and tutorial	May 31	13h-13h30	Introduction to NGS technologies Lecture download	May 31	13h30-15h	Working on NGS data and File Formats with Ugene Tutorial	June 1	13h-14h	Working on NGS data and File Formats with Ugene Tutorial	June 1	14h-15h	Theory on signatures of selection Lecture	June 2	13h-14h	Theory on ancestry analyses Lecture	June 2	14h-15h	Practice demonstrating a selection signatures analysis with cattle data	June 3	13h-15h	Practice demonstrating a breed composition analysis with cattle data	June 4	13h-13h20	Access and Benefit Sharing of genetic resources	June 4	13h20-14h30	Q&A all material
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Links to training material, if available	The course materials, videos and assignments were exchanged via the SLU canvas learning platform. <a href="https://slu-se.instructure.com/courses/4344">https://slu-se.instructure.com/courses/4344</a> (Requires login)
Any other remarks	The free online course sparked a phenomenal response when it was first announced, with over 1,000 early-career scientists from 27 African countries applying for the 40 places available. We received positive feedback from thirty-three participants who completed the course
Course Organizers	Abdulfatai Tijjani, International Livestock Research Institute, ILRI-Ethiopia Emelie Zonabend König, Swedish University of Agricultural Sciences (SLU) Karen Marshall, International Livestock Research Institute, ILRI-Kenya
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