



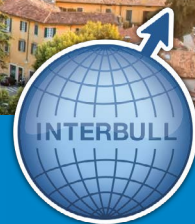
ICAR 2026

VERONA ITALY

FINAL PROGRAMME

Verona, Italy

29 May | 5 June 2026



On behalf of the ICAR board, the Interbull Steering Committee, the local organiser and the Conference Scientific Committee, it is our great pleasure to welcome you to Verona for the ICAR/Interbull 2026 Meeting.

This biannual event of the full ICAR community brings together scientists, technical experts, breeders, the private and public sectors, and other professionals worldwide who share a common commitment to advancing animal recording, genetics, data management, and sustainable livestock production, including animal health and welfare, through collaboration and innovation.

At a time when the livestock sector is evolving rapidly, our community also faces important opportunities and challenges. New ways of breeding livestock populations, along with the increasing use of Precision Livestock Farming (PLF) technologies, such as automated systems and sensor-based data, are transforming how and how much information is collected, analysed, and used on farms. These developments open new possibilities for improving animal health, welfare, efficiency and sustainability, while also raising important questions about data quality, harmonisation, interoperability and the practical delivery of value to farmers and stakeholders.

The ICAR/Interbull Meeting is more than a technical conference. It is a place where ideas become collaborations, where expertise is shared openly, and where collective efforts aim to improve methods to deliver higher-quality services to farmers and stakeholders. In this evolving landscape, ICAR remains committed to supporting the sector, fostering collaboration, and exploring new developments with its members, experts, and partners worldwide.

We would like to warmly thank the scientific committee, all speakers, working group members, sponsors, exhibitors and participants for their valuable contribution and continued engagement. Your expertise and commitment are essential to the success of this meeting and to the future development of our sector.

Special thanks to all the ICAR and Interbull colleagues whose dedication, support, and professionalism contribute to the success of this event.

Finally, we hope that, alongside a productive and stimulating programme, you will also have the opportunity to discover and enjoy beautiful Verona.

Welcome to Verona! We wish you an inspiring and successful ICAR/Interbull 2026 Meeting.

Ana Granados Chapatte, Toine Roozen
ICAR General Manager and Interbull Centre Director

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Sunday, 31st May 2026

AUDITORIUM



08:30>10:30 SESSION I

Experience with Single step evaluation

Chair: Ezequiel Nicolazzi

- 08:30>08:45** Welcome
Interbull Steering Committee Chair, ICAR President
- 08:45>09:00** Variance Component Re-Estimation and Model Simplification in a Single-Step Random Regression Test-Day Model for Nordic Red Dairy Cattle
Timo Pitkanen, *Natural Resources Institute Finland (Luke)*
- 09:00>09:15** Variance component estimation in single-step model using MiX99
Hongding Gao, *Natural Resources Institute Finland (Luke)*
- 09:15>09:30** Modernizing U.S. Calving-Traits Evaluation Under Single-Step
Simone Savoia, *Council On Dairy Cattle Breeding*
- 09:30>09:45** Development of genomic evaluation of breeding value system for milk production traits with on-laboratory quality control of phenotypic data
Ilia Rukin, *JSC "Agroplem"*
- 09:45>10:00** Genetic correlations between juvenile mortality traits and traits in selection in French dairy cattle breeds
Sophie Aguerre, *French Livestock Institute*
- 10:00>10:15** Re-ranking of Imported Angus Sires Following Integration of Foreign Genotypes into a National Genomic Evaluation
Hannah Kruger, *SA Stud Book*
- 10:15>10:30** Advances in single-step for national and multi-country dairy genomic evaluations
Daniela Lourenco, *University of Georgia*

10.30>11.00 ☕ **Coffee Break**

11:00>12:30 **Business Meeting I**

12:30>14:00 🍴 **Lunch**

Sunday, 31st May 2026

AUDITORIUM



14:00>15:30 SESSION II

Implementing new traits in genetic and genomic evaluation systems

Chair: Daniele Vicario, *Associazione Nazionale Allevatori P.R.I*

- 14:00>14:15** Estimation of Genetic Parameters for Lactation Persistency and Productive Maturity in the Italian Brown Swiss Breed
Attilio Rossoni, *ANARB - Italian Brown Cattle Breeders' Association*
- 14:15>14:30** Refinement of U.S. Reproductive Performance Evaluations for the Modern Dairy Herd
Taylor Mcwhorter, *Council on Dairy Cattle Breeding*
- 14:30>14:45** 2025 National Breeding Objective review for Australian dairy cattle
Thuy Nguyen, *DataGene*
- 14:45>15:00** Genetics of increased importance of health traits
Ignacy Misztal, *University of Georgia*
- 15:00>15:30** Q&A session

15:30>16:00 ☕ **Coffee Break**

16:00>17:30 SESSION III

Efficiency of Evaluations and Adaptation/Resilience

Chair: Filippo Miglior, *Lactanet*

- 16:00>16:15** Consequences of ignoring GxE interactions when estimating genetic parameters
Beatriz Castro-Dias-Cuyabano, *INRAE*
- 16:15>16:30** Interaction between adaptation and mitigation traits in Canadian Holsteins
Gabriella Dodd, *University of Guelph*
- 16:30>16:45** Validating the impact of genetic progress for novel and difficult to measure traits in commercial herds
Freddy Fikse, *AbacusBio*
- 16:45>17:00** Walloon strategies targeting new traits for efficiency, mitigation, adaptation and resilience in dairy cattle
Nicolas Gengler, *ULiège-GxABT*
- 17:00>17:15** Genetic residual feed intake and its genetic association to other breeding goal traits
Karoline Bakke, *Geno Breeding and A.I. Association*
- 17:15>17:30** Q&A session

19:30



INTERBULL DINNER AT RISTORANTE MAFFEI

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Monday, 1st June 2026

AUDITORIUM



08:30>10:30 SESSION IV

Big Data for the benefit of genetic and genomic evaluation

Chair: Thuy Nguyen, *DataGene, Australia*

- 08:30>08:45** Using sensor-derived estrus dates to improve the genetic evaluation of fertility in the New Zealand dairy industry
Melissa Stephen, *Massey University*
- 08:45>09:00** Assessing the impact of hormonal interventions on genetic evaluation of fertility in New Zealand
Katarzyna Stachowicz, *AbacusBio Ltd*
- 09:00>09:15** Combining environmental and genomic information in artificial neural networks for phenotypic prediction
Jeremie Vandenplas, *Wageningen University & Research*
- 09:15>09:30** An integrated SaaS ecosystem for standardized phenotypic evaluation and predictive modeling in livestock breeding
Ferdinando Galluzzo, *Breeding & more SRL*
- 09:30>09:45** A Machine Learning Framework for Concurrent Identification of Favorable, Deleterious, and Lethal Variants in Bovine Populations
John Cole, *Council on Dairy Cattle Breeding*
- 09:45>10:00** Guidelines on combining genomic data across populations
Renzo Bonifazi, *Wageningen University & Research*
- 10:00>10:15** European Bovine Evaluation – building joint genetic evaluations across Europe
Johannes Heise, *European Bovine Evaluation FMBA*
- 10:15>10:30** Including Females in MACE evaluation
Julius Mugambe, *Interbull*

10.30>11.00 ☕ Coffee Break

INDUSTRIA ROOM



08:30>12:30 ICAR/IDF ExtraMIR

- 08:30>09:00** Introduction to the ExtraMIR project Reports from the 4 working group leaders
Hélène Soyeurt, *University of Liège*
- 09:00>09:15** ICAR/IDF collaboration
Daniel Nunez, *ICAR* / **Silvia Orlandini**, *ICAR*
- 09:15>09:35** WP1- Update on the reference method comparison
Josee Bordeleau, *MASC*
- 09:35>09:55** Discussion
- 09:55>10:15** WP2- Modelling
Steve Holroyd
- 10:15>10:30** Discussion

10.30>11.00 ☕ Coffee Break

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INDUSTRIA ROOM



- 11:00>11:20** WP3-Quality Assurance
Frederic Dehareng, *Walloon Agricultural Research Center*
- 11:20>11:40** Discussion
- 11:40>12:00** WP4 Communication
Michael Wittacker
- 12:00>12:30** Discussion

12:30>14:00  Lunch

AUDITORIUM



11:00>12:30 **Business Meeting II**

12:30>14:00  Lunch

AUDITORIUM



14:00>15:45 **SESSION V**

Meta-analytical methods

Chair: Sebastian Mucha, *PFH BiPM*

- 14:00>14:15** Visualization of changes in relationships among traits over time - example in type traits
Maciej Misztal, *University of Georgia*
- 14:15>14:30** Estimation of genetic parameters for production traits in Polish Simmental cattle using a random regression test-day model
Dawid Slomian, *National Research Institute of Animal Production*
- 14:30>14:45** Pilot genomic evaluations of French crossbred dairy cattle
Audrey Martin, *INRAE / Eliance*
- 14:45>15:00** Reassessing Wiggans' Method for Estimating Daily Yields in 3x Milking Systems over Four Decades: What Have We Learned?
Nick Wu, *Council on Dairy Cattle Breeding*
- 15:00>15:15** Estimation of across-breed metafounder parameters in all-breed US dairy cattle.
Andres Legarra, *Council on Dairy Cattle Breeding*
- 15:15>15:30** Effects of weighting schemes and base adjustment on US genomic validation
Rodrigo Mota, *Council on Dairy Cattle Breeding*
- 15:30>15:45** Using Interbull validations to improve national genomic predictions
Pete Sullivan, *Senior Geneticist*
- 15:45** Closing of 2026 Interbull Open Meeting
Urs Schnyder, *Qualitas AG*

15:45>16:00  Coffee Break

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COMMERCIO ROOM



14:00>15:30 Milk Analysis SC

15:30>16:00 ☕ Coffee Break

TURISMO ROOM



16:00>17:30 **ICAR Working Group Meeting Artificial Insemination and Related Technologies (Hybrid Meeting)**

1. Welcome and Opening

- Welcome and introduction
- Approval of agenda

2. Commitment Working Group - Review of membership

- Review of membership
- Terms of the reference

3. Bull Semen Proficiency Test

- Overview of results
- Evaluation design
- Future steps

4. Straw ID Alignment

- Steps taken to date
- Questionnaire (please have your phone ready)
- Proposed design
- Future steps

5. Any Other Business

6. Next Meeting

7. Closure

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Tuesday, 2nd June 2026

AUDITORIUM



08:30>13:00 ICAR/IDF WORKSHOP

08:30>08:45

Welcome and Introduction

Christa Egger-Danner, *ZuchtData EDV-Dienstleistungen GmbH*
and **Ilka Klaas**, *DeLaval International AB*
Coordinators of ICAR IDF Sensor Initiative

08:45>09:10

Animal Welfare risk Assessment: Data and the way forward

Yves Van der Stede, *EFSA*

09:10>09:35

Integral monitoring of dairy cattle welfare on farm using precision livestock farming sensors

Pol Llonch Obiols, *University of Barcelona*

09:35>10:00

ICAR IDF Sensor Initiative: Development of a validation protocol

Ilka Klaas, *Delaval and Miel Hostens, University Cornell*

10:00>10:30

Stakeholder perspectives: potential of using sensor data along the dairy value chain with representatives from farmers, veterinarians, consultants and breeding organisations

10:30>11:00

 **Coffee Break**

11:00>12:00

Stakeholder perspectives continued with representatives from dairy processors, welfare assessment users and manufacturers, among others

12:00>12:50

Multi-stakeholder discussion on practical options: Ways forward?
How do we get there? How to make extended use along the dairy value chain?

Moderator: **Ana Granados Chapatte**, *ICAR*
and **Piercristiano Brazzale**, *IDF*

12:50>13:00

Summary and next steps

Christa Egger-Danner, *ZuchtData EDV-Dienstleistungen GmbH*
and **Ilka Klaas**, *DeLaval International AB*

13:00>14:00

 **Lunch**

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Tuesday, 2nd June 2026

TURISMO ROOM



11:00>13:00 ICAR Meeting of the Sheep, Goats and Camelids Working Group

- 11:00>11:05** Opening and welcome
- 11:05>11:15** Organisation and membership
- 11:15>11:25** ICAR survey on milk recording
- 11:25>11:35** What about an ICAR survey on recording on meat sheep
- 11:35>11:50** Survey on genotyping activities: results and questions
- 11:50>12:10** Guidelines on conformation in dairy and meat sheep: update
- 12:10>12:25** EU Reference Centre in small ruminants: update
- 12:25>12:40** BWYPEX Michael Aldrge and perspective on guidelines on feed efficiency and methane emissions
- 12:40>12:50** Toward a codification of sheep and goat breeds within ICAR
- 12:50>12:55** Verona technical session
- 12:55>13:00** Date of next meeting
Any other business and closure

13:00>14:00  *Lunch*

Tuesday, 2nd June 2026

INDUSTRIA ROOM



14:00>15:30 **Open ICAR working group meeting Feed and Gas (Hybrid)**

1. Welcome & Introduction

2. Update Feed&Gas working group

- Membership and Terms of Reference
- Guidelines – ICAR wikipage
- Activities with Global Methane Genetics (Methane Test Centre, Database, webinars & group meetings)

3. Open discussion: Please have your phones ready!

- New topics/traits/name for the group
- Adoption & incentives genetic selection as climate change mitigation tool & policy implementation
- Methane trait definition & harmonization of data pipelines

4. Celebrations

5. AOB & closure

15:30>16:00 ☕ *Coffee Break*

Tuesday, 2nd June 2026

INDUSTRIA ROOM



16:00>18:00 ICAR Animal Data Exchange - working Group Agenda

16:00>16:10

Introductions

Jasper van der Noord, *UNIFORM-agri, Netherlands*

16:10>16:40

ADE technical specification update

Andrew Cooke, *Map of Ag (Rezare Systems)*

- Current Efforts - **Andrew Cooke**, *Map of Ag (Rezare Systems)*
- New data elements to be considered
 - Dairy related data
 - Milk Spectra data
 - Genetic data
 - Other??
 - Beef cattle - **Mike Lynch**
 - Non-Ruminants - **Jean Michel Astruc**, *Livestock Institute*

16:40>17:20

ADE in action

Robert Fourdraine, *Dairy Records Management Systems*

- Practical Implementation of ICAR ADE in Real - World Data Ecosystems - IDDEN Presentation, **Alexey Shelepaev**, *Mtech* and **Valerija Volcic**, *iDDEN GmbH*
- Uniform Agri Presentation - **Jasper van der Noord**, *UNIFORM-agri*
- Smaxtec Presentation - **Katharina Six**
- VAS Presentation - **Jennifer Cline**, *VAS*

17:20>17:50

Roundtable regarding lessons learned implementing ADE

Robert Fourdraine, *Dairy Records Management Systems*

- Our company is interested in using ADE standards - where do I start?
- What are the steps for a successful implementation ?
- Supporting infra-structure
- Developer skillset
- Sharing experiences
 - Systems with poor animal ID
 - Addressing reusable ID's

17:50>18:00

Wrap up / Next Steps

Jasper van der Noord, *UNIFORM-agri*

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08:30>10:00 ICAR General Assembly

10:00>10:30 ☕ Coffee Break

10:30>13:00 ICAR Plenary Session

10:30>10:45

75 years of ICAR

Neil Petreny, *Lactanet* and **Enrico Santus**, *Synergy*

10:45>11:05

The FAO Sustainable livestock Transformation

Thanawat Tiensin, *Director of the Animal Production and Health Division at FAO*

11:05>11:25

Definitions of sustainability of livestock systems, the current landscape, and its importance for global food security and environmental health

Michael Lee, *Deputy Vice Chancellor Harper Adams University, FAO LEAP Scientific Advisor, WFO Scientific Council*

11:25>11:45

Sustainability in dairy: the industry perspective

Pier Cristiano Brazzale, *Former President of IDF*

11:45>12:05

How technology meets sustainability

Claudia Kamphuis, *Researcher Big Data at Wageningen Livestock Research*

12:05>13:00

The 2025-2026 ICAR Brian Wickham Young Person Exchange Program, *BWYPEX*

Moderated by **Jo Newton**, *Agriculture Victoria* and **Katharina Schodl**, *ZuchData*

Main Panel discussion

13:00>14:00 🍴 Lunch

Wednesday, 3rd June 2026

AUDITORIUM



14:00>16:00 Technical Session 1

ICAR / Interbull joint session on “Advances in Genetics and Phenotyping”

Chair: **Urs Schnyder**, *Qualitas AG* and **Daniel Lefebvre**, *Lactanet*

- 14:00>14:15** Limits to selection in U.S. dairy cattle after several generations of genomic evaluation
John B. Cole, *Council on Dairy Cattle Breeding*
- 14:15>14:30** Genetic associations between methane emission and other economically important traits
Björg Heringstad, *Norwegian University of Life Sciences (NMBU)*
- 14:30>14:45** BWYPEX: Discovery, Approval and Management of Genetic Abnormalities in Cattle
Colin Lynch, *Lactanet*
- 14:45>15:00** Digital Innovation at the Service of Breeders: The Synergy Portal and Mobile App for Real-Time Genetic Decision Support
Enrico Carlo Santus, *Synergy*
- 15:00>15:15** Novel approach to develop robust artificial neural network for FT-MIR prediction of enteric methane emissions in dairy cows
Hélène Soyeurt, *University of Liège*
- 15:15>15:30** Using milk mid-infrared spectra from dairy cows to predict methane measured by sniffers and GreenFeed
Anouk van Breukelen, *Wageningen University & Research*
- 15:30>15:45** Prediction of methane emissions based on milk fatty acid profiles shows similar performance to milk MIR spectra
Astrid Köck, *ZuchtData*
- 15:45>16:00** Comparing Methodologies for Deriving Methane Phenotypes from Laser Methane Detector Data in Small Ruminants
Sarah Joe Burn, *BOKU University*

16:00>16:30 ☕ **Coffee Break**

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Wednesday, 3rd June 2026

COMMERCIO ROOM



14:00>16:00 **Technical Session 3**

Animal Identification and National Database

Chair: Ken Evers, ICAR

14:00>14:30 Australia's National Livestock Identification Systems (NLIS)

Elizabeth Bradley, *Integrity Systems Company*

14:30>15:00 Italy's Banca Dati Nazionale (BDN)

Luigi Possenti, *ICT*

15:00>15:30 Advancing national livestock traceability: development and rollout of Rwanda's first cattle identification and registration

Felicien Shumbusho, *RAB*

15:30>16:00 Mexico's Sistema Nacional de Identificación Individual e Ganado (SINIIGA) (National System of Individual Livestock Identification)

Othón Reynoso, *SINIIGA - SINIDA / CNOG*

16:00>16:30 ☕ **Coffee Break**

Wednesday, 3rd June 2026

INDUSTRIA ROOM



14:00>16:00 Technical Session 7

New Developments in Sheep, Goats and Camelids within ICAR

Chair: Jean-Michel Astruc, *Livestock Institute (IDELE)*

- 14:00>14:15** Assessing the adoption of genomic tools in sheep and goat breeding: patterns, challenges, and perspectives
Sotero Salaris, *AGRIS Sardegna*
- 14:15>14:30** Towards ICAR Guidelines on conformation traits in dairy sheep recording
Marija Spehar, *Croatian Agency for Agriculture and Food*
- 14:30>14:45** How ICAR can help in fostering cooperation across country in breeding programs: the case of the Latxa and Manech dairy sheep breeds
Jean-Michel Astruc, *Livestock Institute (IDELE)*
- 14:45>15:00** International collaboration for multinational evaluations in sheep
Michael Aldridge, *AGBU, University of New England*
- 15:00>15:15** Accelerating Digitalisation in the Dairy Goat and Sheep Sector through an Integrated ICAR ADE-Based Management Platform
Tineke Ammerlaan-Sieperda, *UNIFORM-agri*
- 15:15>15:30** Mid-infrared spectroscopy adapted to sheep milk for flock management, performance recording and milk processing
Marine Gelé, *Institut de l'Elevage*
- 15:30>15:45** Standardization of goat milk MIR spectra and fatty acid prediction equations
Marion Calmels, *Seenovia*
- 15:45>16:00** Artificial neural networks outperform partial least squares models for fatty acids estimation using FT-MIR spectrometry on bovine milk: a first step towards transfer learning on small ruminants
Elisabeth Cabaraux, *University of Liège*

16:00>16:30 ☕ **Coffee Break**

- 16:30>16:45** Portable NIRS spectrometer for field diagnosis of anemia in sheep infected with *Haemonchus contortus*
Ana Carolina Chagas, *Brazilian Agricultural Research Corporation*
- 16:45>17:00** Goat colostrum immunological quality: on-farm evaluation
Maddalena Zucali, *University of Milan*

Wednesday, 3rd June 2026

AUDITORIUM



16:30>18:00 **Manufacturers Technical Session**

- 16:30>16:40** From Sample to Value: Building future-ready, smart laboratories to extract the highest value from every milk sample
Pierre Broutin, *Bentley Instruments*
- 16:40>16:50** Trusted solutions for raw milk quality testing
Daniel Schwarz, *FOSS*
- 16:50>17:00** International collaboration added value for creating and spreading predictions at a large scale
Julie Leblois, *European Milk Recording*
- 17:00>17:10** GenoCells: individual somatic cell counts of dairy cows by sequencing tank milk
Jean-Bernard Daviere, *Seenovia*
- 17:10>17:20** DairyWise: turning dairy farm data into profitable herd decisions
Sanna Keski-Nisula, *Mtech Digital Solutions Oy*
- 17:20>17:30** Real-Time PCR in Mastitis Diagnostics
Heidi Hiitio, *Pinehill Goblin Oy*
- 17:30>17:40** Advanced In Line Milk Analyzer and Its Applications
Donatas Majus, *Brolis sensor technology, UAB*
- 17:40>17:50** Potential of on-line SCC in udder health management and herd improvement
Ilka Christine Klaas, *DeLaval International AB*
- 17:50>18:00** Field Implementation of an Inline Milk Composition analyzer and Cloud Analytics System in Dairy Herd
Laima Andruliene, *BouMatic HerdLine*

18:00>19:00



WELCOME RECEPTION

Thursday, 4th June 2026

AUDITORIUM



08:30>10:30 Technical Session 2

Validation related to the Use and Usability of Data

Chair: Steven Sievert, *DHIA* and **Jonas Persson**, *SEGES*

- 08:30>08:45** A new ICAR test facility to validate sniffer devices that measure methane output of dairy cows
Yvette de Haas, *Wageningen University & Research*
- 08:45>09:00** Comparison of Enteric Methane Repeatability Estimates Using GreenFeed and Agscent Measurement Devices
Andrew Lakamp, *University of New England*
- 09:00>09:15** A scalable, automated data pipeline and interactive dashboard for high-throughput methane phenotyping
Lucas Magalhaes Alcantara, *University of Guelph*
- 09:15>09:30** A comparison of data-processing pipelines for methane concentration measurements from sniffers
Coralia Manzanilla-Pech, *Wageningen University & Research*
- 09:30>09:45** The unified framework for processing and storing large-scale sniffers-based emission data
Viktor Milkevych, *Aarhus University*
- 09:45>10:00** Automating Data Unification, Governance, and Analytics in Animal Science: A Cloud-Native, AI-Assisted Framework
Puchun Niu, *Cornell University*
- 10:00>10:15** Whole-genome ICAR parentage verification certification
Ezequiel Luis Nicolazzi, *Council on Dairy Cattle Breeding (CDCB)*
- 10:15>10:30** Monitoring milk meters: the French experience with this practice, between innovation and user feedback
David Saunier, *ELIANCE*

Thursday, 4th June 2026

COMMERCIO ROOM



08:30>10:30 Technical Session 5

From Milk Analysis to Decision Support: Unlocking Insights for Sustainable dairy management

Chair: Josée Bordeleau, *Lactanet* and **Maria Frizzarin**, *Agroscope*

08:30>08:45

Intra- and cross-laboratory monitoring framework for MIR-integrated milk laboratory data

André Mensching, *Vereinigte Informationssysteme Tierhaltung w.V. (vit)*

08:45>09:00

Extramir spectral proficiency test: comparison of spectral standardisation methods

Clement Grelet, *Walloon Agricultural Research Center*

09:00>09:15

Kinetic assessment of the impact of a linseed-rich diet on a milk FT-MIR-based herbage intake indicator

Dichou Killian, *Université de Liège*

09:15>09:30

Modular Open-Source Framework for Spectral Phenotyping Using Neighbourhood-Adaptive PLSR to Predict Nitrogen Use Efficiency in a Dairy Herd

Ross Muers, *SRUC*

09:30>09:45

Does MIR-based information contribute to practical decision-making on the farm?

Dale Laura Monica, *LKV Baden Württemberg*

09:45>10:00

Decoding lipolysis through individual free fatty FT-MIR model: Enhancing predictive accuracy through real-sample enrichment and advanced algorithms

Octave Christophe, *CRA-W*

10:00>10:15

Considerations for the Establishment of Practical Accuracy Limits for On-Farm Milk Analysers for Official Milk Recording

Robert (Rob) Orchard, *MSD Animal Health*

10:15>10:30

Closing the Utilisation Gap: Converting Routine Milk Analysis Data into Decision Support for More Sustainable Dairy Farming

Daniel Schwarz, *FOSS*

Thursday, 4th June 2026

INDUSTRIA ROOM



08:30>10:30 Technical Session 4

Heat Stress, Health and Welfare: Genetics and Management

Chair: **Christa Egger-Danner**, *ZuchtData* and **Marie Haskell**, *SRUC*

08:30>08:45

Novel approaches to breeding for climate-smart dairy cattlea

Gabriella Dodd, *University of Guelph*

08:45>09:00

Genomic prediction of heat tolerance from rumen sensor data in Australian Holstein cows

Laura Jensen, *Agriculture Victoria*

09:00>09:15

Detecting heat stress in dairy cattle using milk composition data

Marine Gelé, *Institut de l'Elevage*

09:15>09:30

Prediction of dairy cows' heat stress response from milk mid-infrared spectra for detection and genetic selection

Pauline Lemal, *University of Liège*

09:30>09:45

Actionable output developed from milk fatty acids to assess rumen health, negative energy balance and production loss in dairy cows

Ines Adriaens

09:45>10:00

Does the automation level of a dairy farm affect animal welfare?

Susanne Demba, *CRV BV*

10:00>10:15

Q Index: A nationwide welfare index based on DHI and cattle register data

Joachim Braunleder, *Vereinigte Informationssysteme Tierhaltung w.V.*

10:15>10:30

Leveraging herd management, genetics, and DHI by integrating all three data types into a new transition cow decision tool

Robert Fourdraine, *Dairy Records Management Systems*

10:30>11:00 ☕ **Coffee Break**

Thursday, 4th June 2026

AUDITORIUM



11:00>13:00 Technical Session 6

Milk Recording in a Technology World: Providing Value Added Services and Data Exchange Solutions for Farms with Robots and Sensors

Chair: Robert Fourdraine, DRMS

11:00>11:15

ICAR ADE as an Enabler for Inclusive Data Integration:
A Cloud-Based Herd Management Case Study from Catalonia

Tineke Ammerlaan - Sieperda, UNIFORM-agri

11:15>11:30

{farm-twin}: An Open-Source ICAR ADE-Aligned Farm-Scale Digital Twin

Matthew Broadbent, Scotland's Rural College

11:30>11:45

LiveStack: An open-source framework for creating livestock data pipelines

Braeden Fieguth, University of Guelph

11:45>12:00

Stakeholder Role Differences in the Use of Precision Livestock Farming Technologies in Europe

Silvia Orlandini, ICAR

12:00>12:15

Generative AI for Dairy Life Trajectory Modeling: Transforming Milk Recording and Sensor Data into Actionable Farm Intelligence

Asha Miles, DRMS

12:15>12:30

An accurate machine learning approach to predict fat, protein and milk yield before and during lactation from test-day records

Ines Adriaens, CRV BV

12:30>12:45

Machine Learning Approach for Early Prediction of 305-Day Standard Lactation

Jakob Ganitzer, ZuchtData EDV-Dienstleistungen GmbH

12:45>13:00

Reconstructing representative daily Fourier transform mid-infrared milk spectra from partial milking robot samples using machine learning

Franceschini Sébastien, University of Liège

13:00>14:00  Lunch

FINAL SCIENTIFIC PROGRAMME



ICAR 2026
VERONA ITALY

Thursday, 4th June 2026

COMMERCIO ROOM



11:00>13:00 **Technical Session 9**
Milk Analyses Round Table (11:00-12:00)

13:00>14:00  Lunch

INDUSTRIA ROOM



11:00>13:00 **Technical Session 8**
From Measurement to Genetics: Methane Emission and Feed Efficiency
Chair: Raffaella Finocchiaro, ANAFIBJ and **Rasmus Bak Stephansen**, Aarhus University

11:00>11:15 Cattle Feed InTake – A system to identify cows, measure feed intake and bod weight in dairy cattle under commercial settings using 3D cameras
Jan Lassen, Vikinggenetics

11:15>11:30 Recording methane and feed intake internationally
Michael Aldridge, AGBU, University of New England

11:30>11:45 Heritability of feeding behaviour traits and the genetic relationships with dry matter intake in growing dairy bulls
Rhiannon Handcock, Livestock Improvement Corporation (LIC)

11:45>12:00 International perspectives on feed intake phenotyping and feed efficienc in cattle
Sarah Jo Burns, BOKU University

12:00>12:15 Tech comparison of methane measurements devices and alternative predictors in dairy cows
Fazel Almasi, Agriculture Victoria Research

12:15>12:30 breed4green: First results from a dataset on methane emissions in commercial farms for genetic studies
Kristina Linke, ZuchtData EDV Dienstleistungen GmbH

12:30>12:45 Building a Global Integrated Methane Data Ecosystem: Harmonizing Measurement, MIR, and Genomic Data to Accelerate Livestock Methane Mitigation
Puchun Niu, Cornell University

13:00>14:00  Lunch

Thursday, 4th June 2026

AUDITORIUM



14:00>16:00 Technical Session 10

Adoption and Incentives to Breed for Methane Mitigation

Chair: Birgit Gedler-Grandl, *WUR-ABG* and **Roel Veerkamp**, *WUR-ABG*

14:00>14:15

Challenges and options around incentivisation of genetic approaches to mitigation of methane emissions from ruminants

Robert Banks, *c/- AGBU*

14:15>14:30

Towards guidelines to validate genetic evaluations for enteric methane emissions

Jeremie Vandenplas, *Wageningen University & Research*

14:30>14:45

Approaching a crossroads in selecting for methane production: Trait definition and its consequences for breeding programs

Marc Rutten, *Wageningen University & Research*

14:45>15:00

The effect of including methane traits in selection indexes for ruminants

Julius Van der Werf, *University of New England*

15:00>15:15

Enabling adoption of methane-reducing breeding programs through GHG accounting and MRV frameworks

Merel Moleman, *Wageningen University & Research*

15:15>15:30

Integrating Genetic Breeding Values for Enteric Methane Emissions into Carbon Accounting Frameworks

Katherine Grant, *AbacusBio*

15:30>15:45

Economics of breeding schemes for improved methane efficiency in dairy cattle

Fernanda Ferreira, *Clean Air Task Force*

15:45>16:00

Assessing Breeding Strategies to Mitigate Methane Emissions in Danish Dairy Cows

Hanna-Marie Nielsen, *Center for Quantitative Genetics and Genomics*

16:00>16:30  **Coffee Break**

Thursday, 4th June 2026

COMMERCIO ROOM



14:00>16:00 Technical Session 13

Exploring Current and Future Needs for the Livestock Science and Farming Community

Chair: **Ana Granados**, ICAR and **Raffaella Finocchiaro**, ANAFIBJ

14:00>14:15

How does rate of genetic gain impact economic optimum productive life?

Joanna Newton, *Agriculture Victoria*

14:15>14:30

European Network on Livestock Phenomics (EU-LI-PHE): an international initiative aimed at facilitating the application of phenomics in animal breeding

Luca Fontanesi, *University of Bologna*

14:30>14:45

Developing a Genetic Submodule in the Ruminant Farm System (RuFaS) Model for Enhanced Whole-Farm Representation

Victor Cabrera, *University of Wisconsin-Madison*

14:45>15:00

ICAR/IDF Sensor Initiative: Use of sensor data to monitor and improve health and welfare of dairy cattle – achievements, challenges and future perspectives Ilka

Christine Klaas, *DeLaval International AB* and **Miel Hostens**, *Cornell University*

15:00>15:15

BROMEDIR – Development and evaluation of a miniaturized FT-MIR instrument for in-farm milk analysis

Mazen Erfan, *Si-Ware Systems* and **Clement Grelet**, *CRA-W*

15:15>15:30

Transforming routine MIR milk data into actionable tools for Dairy Herd Improvement and cheese production

Marine Gelé, *Institut de l'Élevage*

15:30>15:45

HerdPlanner: model-based tactical decision support for reproduction and culling

Elmer Harms, *CRV BV*

15:45>16:00

Reimagining Animal Identification: Ireland's Progress Towards a Fully Genotyped National Herd

Kevin Mc Dermott, *ICBF*

16:00>16:30 ☕ **Coffee Break**

Thursday, 4th June 2026

INDUSTRIA ROOM



14:00>16:00 Technical Session 12

From Guidelines to the Barn: Implementing Section 2 of ICAR's Milk Recording Guidelines - Updates and Lessons from the Field Milk Recording Strategies in Buffalo, an Urgent Update

Chair: Miel Hostens, Cornell University and Angela Salzano, University of Naples Federico II

- 14:00>14:15** A flexible non-parametric approach to assessing dairy cow performance
Ivo Lagarde, CRV
- 14:15>14:30** BWYPEX: A unified Python implementation of standardized 305 day yield calculation methods
Meike Beatrijs Van Leerdam, Cornell University
- 14:30>14:45** Opportunities and challenges in accurate milk recording on farm in the United States
Asha Miles, DRMS
- 14:45>15:00** The Genetic Transformation of the Italian Mediterranean Buffalo: Five Years of Progress and New Frontiers
Mayra Gomez, Italian National Association of Buffalo Breeders
- 15:00>15:15** Bayesian Modeling of Lactation Curves in Italian Mediterranean Buffalo Using the MilkBot Equation
Lucia Trapanese, University of Naples Federico II
- 15:15>15:30** Analyzing milk recordings to support better decision-making through AI: a case study in Italian Mediterranean Buffalo
Valentina Russo, Logogramma SRL
- 15:30>15:45** Evaluating the Robustness of Cumulative Milk Yield Estimation Methods under Limited Test-Day Availability in Italian Mediterranean Buffalo
Lucia Trapanese, University of Naples Federico II
- 15:45>16:00** Phenotypic variation of milk antioxidant activity in Italian Mediterranean buffaloes
Matteo Santinello, University of Naples Federico II

16:00>16:30 ☕ **Coffee Break**

Thursday, 4th June 2026

AUDITORIUM



16:30>17:30 Adoption and Incentives to Breed for Methane Mitigation Roundtable

COMMERCIO ROOM



16:30>18:00 Technical Session 11

Milk Recording Supporting Sustainability Initiatives

Chair: Tone Roalkvam, TINE

- 16:30>16:45** Farm sustainability index using dairy cattle recording data: the case of Slovenia
Betka Logar, Agricultural Institute of Slovenia
- 16:45>17:00** Combining milk recording and other on-farm data to streamline and improve greenhouse gas emissionsg
Andrew Cooke, Map of Ag (Rezare Systems)
- 17:00>17:15** Variation of enteric methane intensity predicted from bulk tank milk in commercial dairy herds
Débora Santschi, Lactanet
- 17:15>17:30** Using Bulk tank Milk Spectra to Predict Enteric Methane Emissions from Lactating Cows in Commercial Dairy Herds
Liliana Fadul, Lactanet
- 17:30>17:45** BWYPEX Program Daily methane emission predicted from milk MIR spectra: the time of model comparison
Maria Frizzarin, Agroscope and Amelie Vanlierde, CRA-W
- 17:45>18:00** Beyond milk collection: Can Indonesian dairy cooperatives lead the sustainable record-keeping transition?
Yuni Resti, BOKU University, IPB University

Thursday, 4th June 2026

INDUSTRIA ROOM



16:30>18:00 Technical Session 14

Beef Cattle – Genetic Evaluation and Recording

Chair: Kim Matthews, *EBLEX*

16:30>16:45

Implementation of genetic evaluation for intramuscular fat in Norwegian beef cattle using ultrasound measurements

Siri Furre, *Tyr*

16:45>17:00

Genetic parameters and breeding values for feed intake and feed efficiency traits from the Staur performance test in Norwegian beef cattle

Siri Furre, *Tyr*

17:00>17:15

Towards standardised LR protocols for international beef cattle genetic evaluations

Gabriel Soares Campo, *Interbull*

17:15>17:30

Genetic parameters for calving traits in Norwegian Beef-on-Dairy crosses

Ida Wøyen Hamfjord, *TYR*

17:30>17:45

PHENO3D: 3D phenotyping of beef calves – scaling image processing into edge-AI and extending predictions across breeds

Bruyas Maxence, *ELIANCE*

17:45>18:00

Discussion

19:30>24:00



GALA DINNER

Friday, 5th June 2026

TECHNICAL TOURS

Tour 1

Nordera farm (half-day)

Departure at 8:45 by bus from Piazzale XXV Aprile (in front of Verona Porta Nuova Train Station)

Return at 14:00 at the Piazzale XXV Aprile (in front of Verona Porta Nuova Train Station)

Tour 2

ST Genetics Laboratory and Valserena Farm (Full day)

Departure at 7:45 by bus from Piazzale XXV Aprile (in front of Verona Porta Nuova Train Station)

Return at 18:30 at the Piazzale XXV Aprile (in front of Verona Porta Nuova Train Station)

Tour 3

ARAV (Associazione Regionale Allevatori Veneto) Laboratory (half-day)

Departure at 8:45 by bus from Piazzale XXV Aprile (in front of Verona Porta Nuova Train Station)

Return at 14:00 at the Piazzale XXV Aprile (in front of Verona Porta Nuova Train Station)

Title	Title	Presenter
Technical Session 2	Quality of data collection and its implications for the reliability of large-scale animal science analyses	Victor Cabrera
Technical Session 2	Preliminary Evaluation of Phytogenic Dry-Off Tablet on Dairy Cow Transition Health and Performance with pH Plus Bolus from smaXtec.	Ackaert Geoffrey
Technical Session 3	Animal identification in the Bosnia and Herzegovina: current situation and possibilities	Miljan Erbez
Technical Session 3	From Sectoral Traceability to a National Identification Policy: The Role of SISBOV in the Development of Brazil's National Cattle Identification Plan (PNIB)	Jorge Gomes
Technical Session 4	A Mobile Infrared Thermography System for Precision Phenotyping and Health Monitoring of Dairy Cows under Heat Stress Conditions	Jeng-Fang Huang
Technical Session 4	Effects of Cold Drinking Water on Panting Behavior and Milk Yield in Lactating Dairy Cows During Summer	Jeng-Fang Huang
Technical Session 4	Assessment and prediction of heat tolerance in dairy cows using blood biochemical indicators	Jeng-Fang Huang
Technical Session 4	Automated Claw Locomotion Scoring System for Dairy Cows	Chia Xin Lee
Technical Session 4	Impact of Automated Milking Systems on Productivity, Milk Composition, and Udder Health: A Case Study in Subtropical Taiwan	Jeng-Fang Huang
Technical Session 5	Optimization of Differential Somatic Cell Count (DSCC) Threshold for Enhanced Early Screening of Escherichia coli mastitis in Dairy Cattle	Ya Hsin Kuo
Technical Session 5	Raw milk composition associated with texture profile attributes of low-moisture mozzarella cheese in Taiwan	SzuHan Wang
Technical Session 5	A Farm Management Monitor for the Dairy on Bulk Milk Laboratory Testing, a longitudinal case study on a high-level producing Dutch Farm	Wim Schielen
Technical Session 6	Automated Claw Locomotion Scoring System for Dairy Cows	Zhen-Fang Huang
Technical Session 7	Genomic reconstruction of pedigree records: a key step in the revitalization of the critically endangered Istrian goat	Ante Kasap
Technical Session 7	Assessing income-based economic efficiency in PDO Manchega sheep breed dairy farms: preliminary results	Manuel Luque
Technical Session 7	Animal Genetic Banking in Georgia: Conservation and DNA/RNA Sequencing of Capra caucasica and Capra cylindrica	Kakha Nadiradze
Technical Session 8	Assessing Breeding Strategies to Mitigate Methane Emissions in Danish Dairy Cows	Hanne Marie Nielsen
Technical Session 11	Automated Bottom-Reading Identification of Sample Vials to Ensure 100% Traceability in Robotic Milking Systems (Patent Pending)	Rodrigo Carnero
Technical Session 12	A comparative evaluation of conventional and automatic milking systems on performance and behaviour of dairy buffaloes	Chiara Evangelista
Technical Session 12	Modeling Lactation Curves Using Daily Yield and Milking Interval to Identify Efficient Cows in Automatic Milking Systems: A Case Study in Taiwan	Jeng-Fang Huang
Technical Session 12	Difference of lactation curve in 305-day milk yield between automatic and conventional milking system	Yueh-Tung Chen
Technical Session 12	Impact of Automated Milking Systems on Productivity, Milk Composition, and Udder Health: A Case Study in Subtropical Taiwan	Hsiao Han Liao
Technical Session 13	The phytogenic active ingredient AHV9 drives dose-dependent methane mitigation in vitro	Ackaert Geoffrey
Technical Session 13	ICAR/IDF initiative on sensor data for functional traits: reference standards and guidelines to quantify rumination for health and animal welfare recording	Miel Hostens
Technical Session 14	Drivers of Mobile App Adoption in Extensive Beef Cattle Farms: preliminary results	Manuel Luque
Technical Session 14	Animal Welfare Assessment in Extensive Avileña-Negra Ibérica and Asturiana de los Valles Beef Cattle Farms	Manuel Luque



INNOVATIVE ANALYTICAL AND AUTOMATION SOLUTIONS FOR THE CENTRAL MILK TESTING LABORATORIES

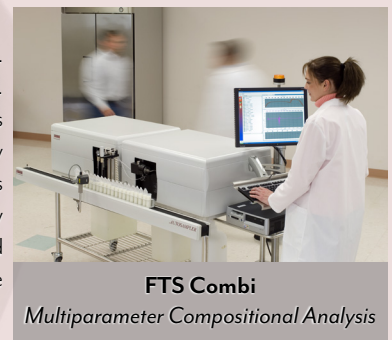
HIGH-THROUGHPUT, STANDARDIZED AND FUTURE-READY SOLUTIONS FOR THE CENTRAL MILK TESTING LABORATORIES

As dairy testing laboratories face growing demands for quality, traceability, throughput, and reliability, the challenge is no longer simply to operate high-performance instruments, but to build a coherent and scalable analytical organization. Bentley Instruments supports this transformation with analytical and automation solutions designed for both current and future industry needs.

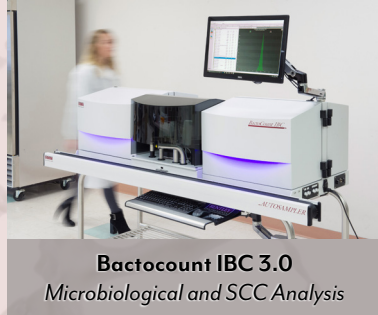
Bentley Instruments' approach is based on a simple principle: a modern laboratory reaches its full potential when the entire workflow operates as one continuous, standardized, and intelligent process. From sample reception to final result reporting, every step can be secured, harmonized, and optimized. Automating sample sorting, aliquoting, preparation, and transfer to analyzers saves time, reduces manual handling, minimizes errors, and ensures consistent analytical performance.

Comprehensive Physicochemical and SCC Analysis

The laboratory of tomorrow relies not only on automation, but also on the quality and scalability of its analytical instruments. For somatic cell counting and physicochemical analysis, the FTS Combi stands out as a comprehensive, future-oriented solution. Meeting the requirements of ISO 9622 | IDF 141 and ISO 13366-2 | IDF 148-2 standards, it can measure up to 64 parameters simultaneously with throughput of up to 600 analyses per hour. It is distinguished by high-value proprietary technologies, notably a patented interferometric technology for real-time control of IR cuvette spacing for optimal spectrum standardization, as well as patented approaches to calibration development and deployment. These advanced capabilities support the creation of highly specific and precise calibration models, including applications such as ketosis prediction, where physicochemical blood-related parameters can be predicted through milk analysis. The FTS Combi therefore positions itself not only as a high-performance analyzer, but also as a true precision platform designed to support the evolving analytical needs of dairy laboratories.



FTS Combi
Multiparameter Compositional Analysis



Bactocount IBC 3.0
Microbiological and SCC Analysis

Scalable Microbiological Testing Built for the Future

Alongside physicochemical and somatic cell analysis, microbiological testing remains a key part of laboratory activities. For this purpose, the Bactocount IBC 3.0 delivers an advanced solution for bacterial counting based on flow cytometry. Built around two lasers and multiple fluorescence detectors, it goes far beyond routine analysis by enabling multiplex capabilities and supporting future applications. This advanced architecture makes the Bactocount IBC 3.0 a scalable platform designed to evolve with laboratory needs over time. With throughput of up to 200 analyses per hour, the Bactocount IBC 3.0 is validated according to ISO 16140 and MicroVal standards for the analysis of raw cow, goat, and sheep milk. It uses a highly stable reagent solution, contributing to both improved operational cost control and a high level of day-to-day laboratory reliability.

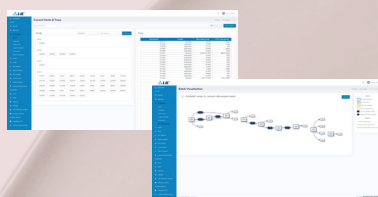
Automation as a Driver of Lab Performance

To fully unlock the value of advanced analytical instruments, laboratories also need automation capable of connecting, standardizing, and securing the entire workflow. At the core of this transformation, laboratory automation becomes a strategic asset. With the ILAS 3200, Bentley Instruments offers a modular, flexible, and fully integrable solution designed to automate and standardize sample preparation and analyzer interfacing. With RFID or barcode traceability, rule-based workflow automation, and seamless LIMS integration, laboratories improve productivity, quality control, and traceability at the same time. Automation is not only about working faster, but about working better — with greater consistency, security, and profitability. The ILAS 3200 supports high-throughput operations, handling up to 1,200 analyses per hour with two FTS Combi analyzers and up to 400 analyses per hour with two Bactocount IBC 3.0 analyzers, making it a true capacity multiplier for central milk testing laboratories.



MOTION DESIGN
AUTOMATION AND SOFTWARE SOLUTIONS

ILAS 3200
Laboratory automation



One Integrated Ecosystem to Streamline Laboratory Operations

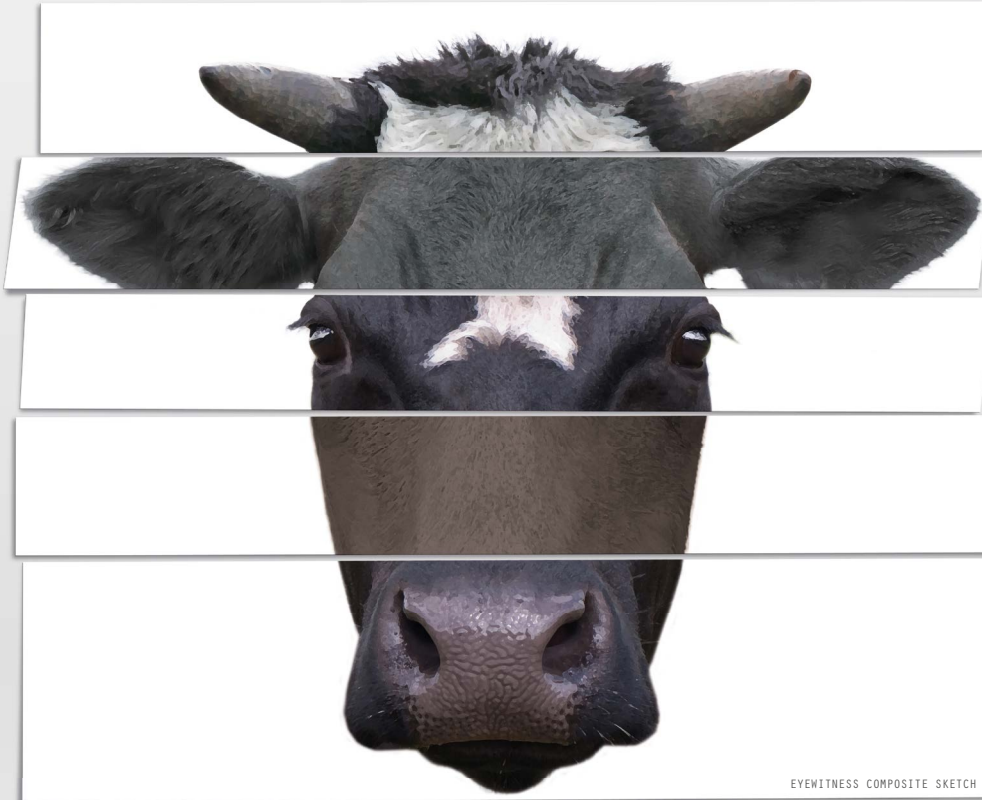
A key strength of Bentley Instruments' approach lies in its ability to combine analytics and automation in one coherent environment, fully integrable with the laboratory's information system. When automation platforms, analyzers, and LIMS communicate seamlessly, laboratories benefit from smoother information flow, stronger traceability, tighter quality control, more secure analytical data, fewer retests, and better overall service valorization. Optional KPI integration can further enhance performance monitoring and operational oversight. The result is a laboratory that is more robust, transparent, and easier to manage strategically.

Bentley Instruments: Enabling the Dairy Testing Laboratory of the Future

Ultimately, the dairy laboratory of the future will be the one that successfully combines state-of-the-art analytical instruments with full automation to transform routine testing into a highly efficient, standardized, and scalable service. With solutions such as the ILAS 3200, the Bactocount IBC 3.0, and the FTS Combi, Bentley Instruments provides laboratories with the tools they need to optimize capacity, streamline workflows, secure result quality, improve profitability, and prepare confidently for tomorrow's analytical requirements. More than an equipment supplier, Bentley Instruments positions itself as a transformation partner for laboratories seeking to turn analytical performance into a true strategic advantage.

“ I THINK HE HAD
A BIGGER NOSE ”

„ ICH DENKE , DIE NASE
WAR ETWAS GRÖSSER “



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EINFACH – SICHER – IDENTIFIZIEREN

Mistakes are easily made, that's why we developed the tissue-sampling ear tag **FLEXOPLUSGENO**. Extraction of a tissue sample is far and away the easiest method of acquiring a “genetic fingerprint” of an animal because it can be carried out at the same time as the ear tag for the animal identification is attached.

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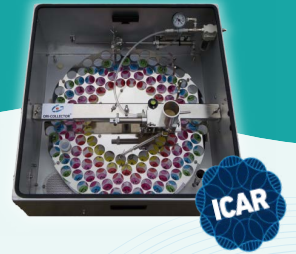


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ICAR 2026
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- Better visibility across cattle data and processes
- Improved efficiency and profitability in operations
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- Advantages:
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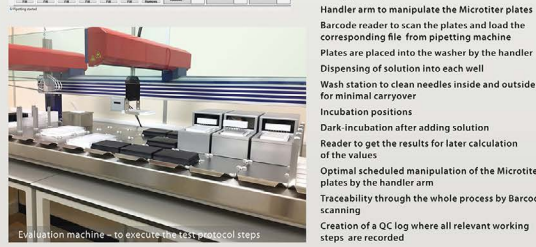
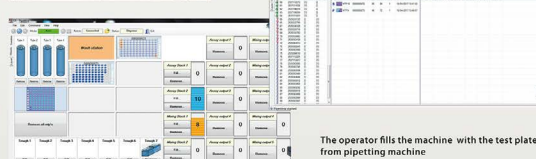
ARS Lxxx Pipetter

Inline or standalone Pipetter to pipette milk into Mtp or Micronicblock



- Pipetting of samples according to pre-programmed analysis code
- Manipulation of Microtiter plates by a handling arm
- Barcode reading of the plates
- Creation of a log-file where the sample information is added after dispensing the milk
- Creation of a QC log where all relevant working steps are recorded
- Adding of control samples in pre-defined wells of the Microtiter plate
- Wash station to clean needles inside and outside for minimal carryover
- Dispensing of solution into each well
- Placing the plate onto a shaker for a pre-definable time
- Transfer from mixing plate into test plate
- Output stacker for finished plates
- Pre-stored information will be amended with appropriate coordinates to the later results

- Additional components:
- Barcode reading
 - Disposable sampling tips
 - Plate shaker
 - Plate washer
 - Plate incubator
 - Plate reader



- The operator fills the machine with the test plates from pipetting machine
- Handler arm to manipulate the Microtiter plates
 - Barcode reader to scan the plates and load the corresponding file from pipetting machine
 - Plates are placed into the washer by the handler
 - Dispensing of solution into each well
 - Wash station to clean needles inside and outside for minimal carryover
 - Incubation positions
 - Dark-incubation after adding solution
 - Reader to get the results for later calculation of the values
 - Optimal scheduled manipulation of the Microtiter plates by the handler arm
 - Traceability through the whole process by Barcode scanning
 - Creation of a QC log where all relevant working steps are recorded

with system to success

WMB

An Innovative Milk Flow Meter for Addressing Urgent Questions in Official Milk Recording



*Since 2018 officially approved for cows, goats and sheep – all with one device. Buffalo approval in progress.



GENERAL INFORMATION



ICAR2026
VERONA ITALY

SOCIAL EVENT

Sunday, 31st May 2026 Interbull Gala Dinner

The Social Event will take place at **Ristorante Maffei** on Sunday, 31st May, starting at **19:30**.

Access is strictly reserved for participants who have purchased a valid dinner ticket. Entry will be granted only upon presentation of the event voucher, which will be provided on-site exclusively to those registered for the social event.

Please note that access without a valid voucher will not be permitted.

Wednesday, 3rd June 2026 ICAR Welcome Cocktail

The ICAR welcome cocktail will follow the close of the programme on 3rd June, starting at **18:00**, and will take place in the exhibition area of the congress center as a light networking reception.

Access is open to all registered congress participants; attendees are kindly requested to wear their badge at all times during the event.

Thursday 4th June 2026 ICAR Gala Dinner

The ICAR Gala Dinner will take place at **Giardino Giusti**
Via Giardino Giusti 2 - Verona, on Thursday, 4 June 2026 at **20:00**.

Access is strictly reserved for participants who have purchased a valid dinner ticket.

Entry will be granted only upon presentation of the event voucher, which will be provided on-site exclusively to those registered for the social event.

Please note that access without a valid voucher will not be permitted.

Kindly note that the venue features gravel pathways; we therefore recommend appropriate footwear and advise against high heels, flip-flops, or open shoes.

CONGRESS VENUE

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ORGANIZING SECRETARIAT

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REGISTRATION DESK OPENING HOURS

Starting from 30 May 16:00 > 18:00

SECRETARIAT OPENING HOURS

29 May	07:00 > 18:00	1 June	08:00 > 18:00
30 May	08:00 > 18:00	2 June	08:00 > 18:00
31 May	07:00 > 18:00	3 June	07:00 > 18:00
		4 June	08:00 > 18:00



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