



RuminOmics

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Connecting the animal genome, the intestinal microbiome and nutrition to enhance the efficiency of ruminant digestion and to mitigate the environmental impacts of ruminant livestock production



Professor John Wallace

Collaborative project Jan. 2012 – Dec. 2015 www.ruminomics.eu





RuminOmics - Response to technology

THE CALL:

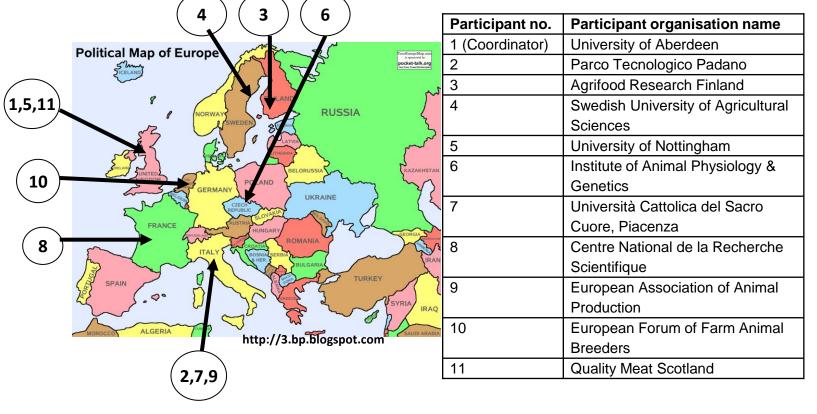
KBBE.2011.1.1-03: Efficiency of ruminant digestive systems and reduction of the ecological footprint through a combination of systems biology, 'omics'

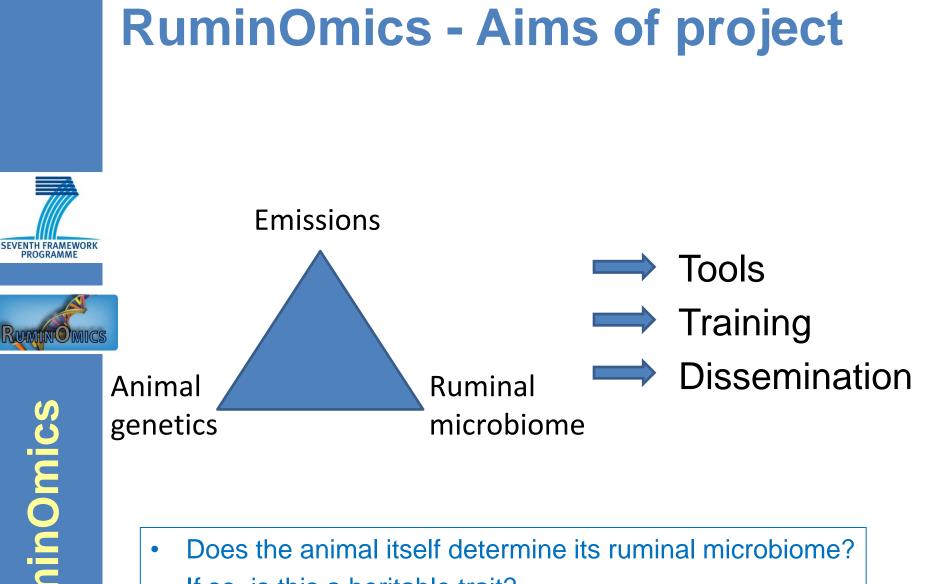


RuminOmics - Partners







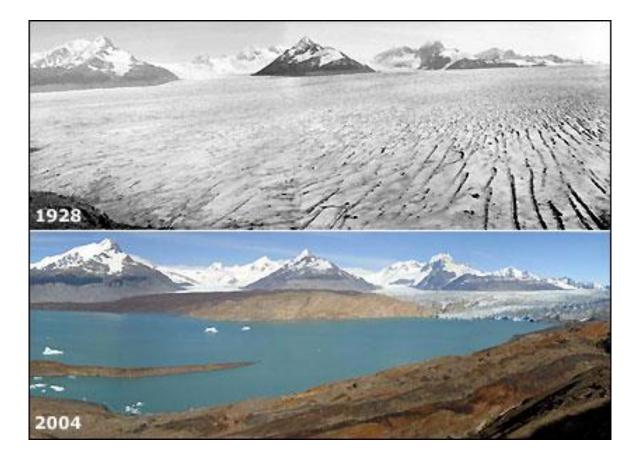


- If so, is this a heritable trait? ۲
- How does nutrition affect this relationship?

Methane, ruminants and the environment

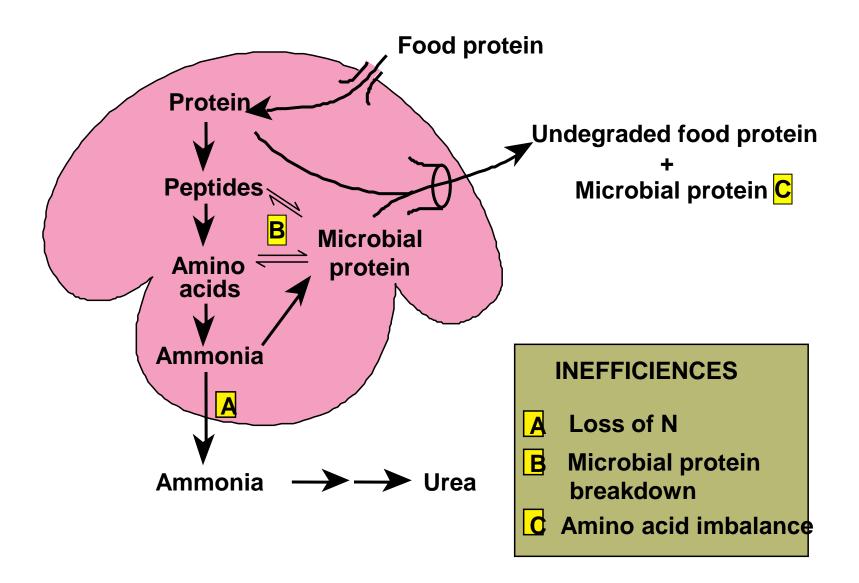




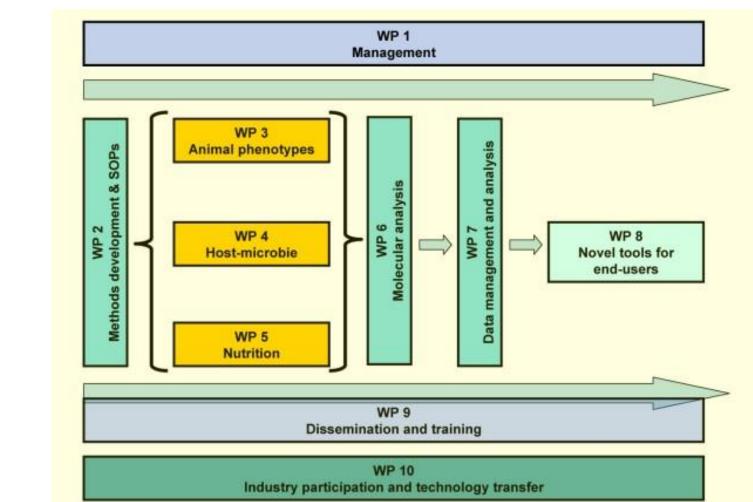




Protein metabolism



Work Package Structure







RuminOmics - Experiments





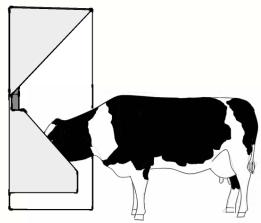
1000 cows in UK, Italy, Sweden and Finland

Methane N emissions FCE Milk quality

Ruminal microbiome

Animal genotype

- **40 cows in Sweden and Finland** Impact of N, CHO, lipid nutrition
- 50 cows in UK, Italy, Sweden and Finland Full metagenome analysis



RuminOmics - Experiments II

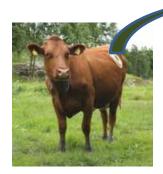
Bovine single-egg twins





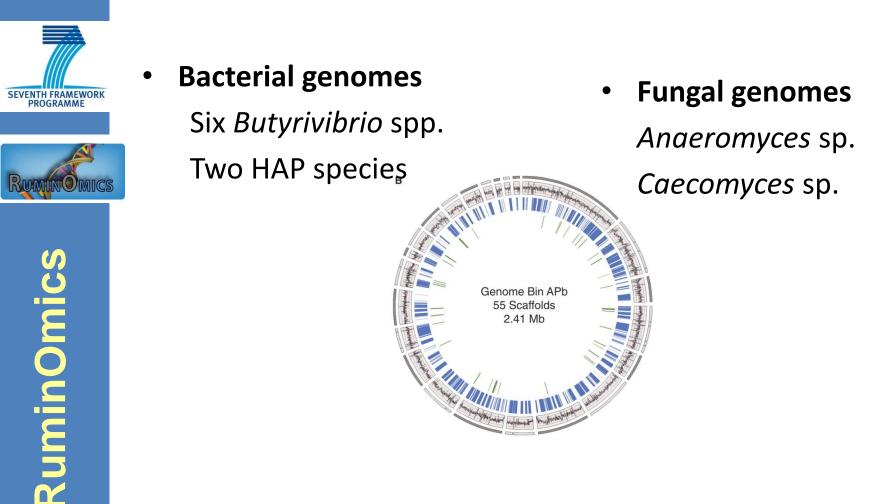


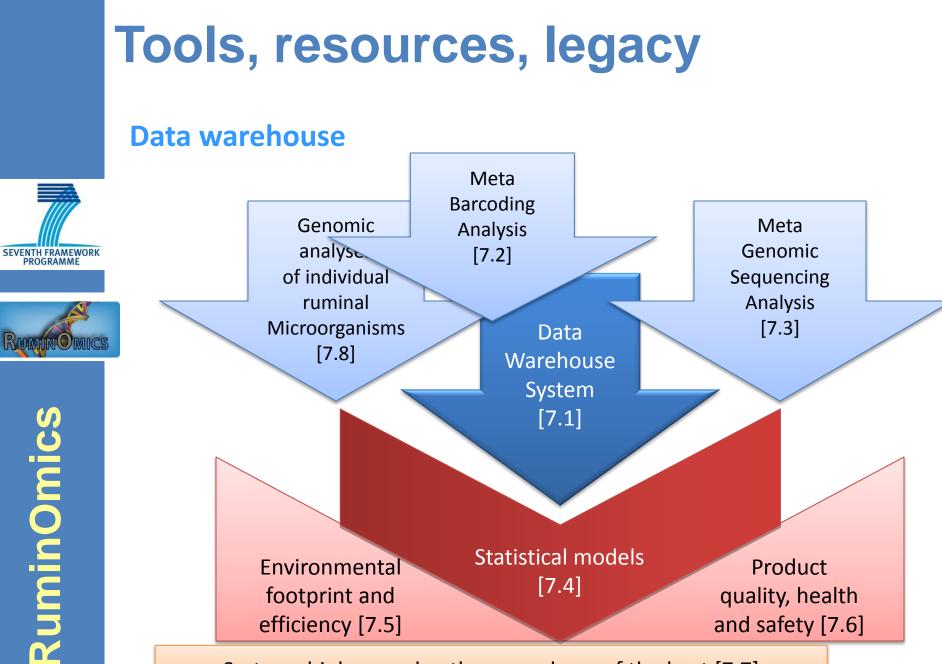
Interspecies digesta transfer





RuminOmics - Experiments III



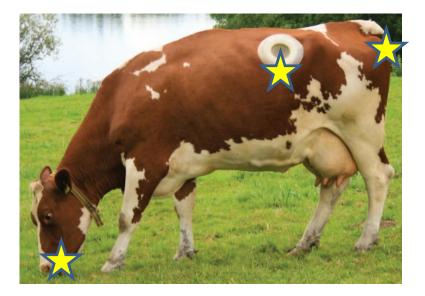


Systems biology and pathway analyses of the host [7.7]

Tools, resources, legacy

Proxies

Buccal-ruminal-faecal microbiomes



RuminOmics

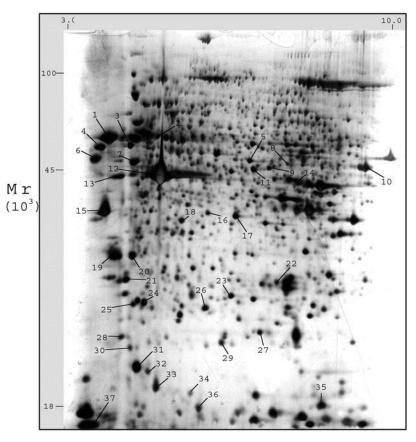
SEVENTH FRAMEWORK PROGRAMME

Tools, resources, legacy

• Tools

Metaproteomics

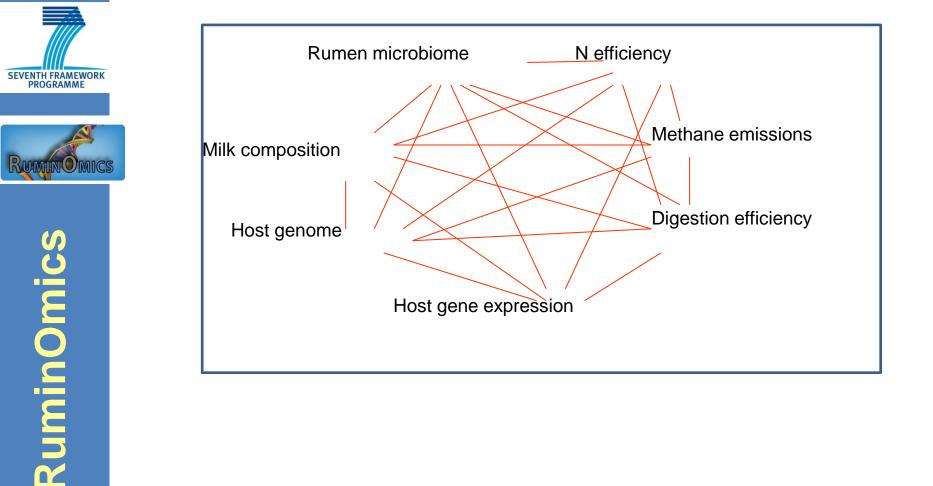
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The Bioinformatics challenge



RuminOmics - Aspirations

- The answer to the animal-microbe conundrum
- Bioinformatics legacy
- Trained & more efficient industry
- Environment legacy