



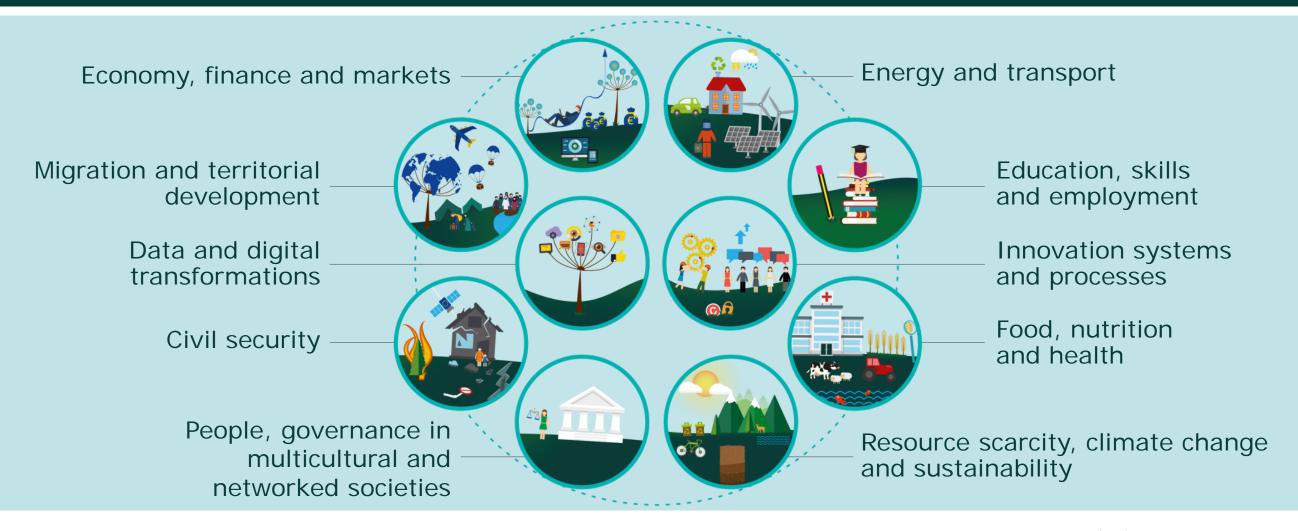
JRC Mission

The Joint Research Centre (JRC) is the European Commission's science and knowledge service.

II Our mission is to support
EU policies with independent evidence
throughout the whole policy cycle



JRC 10 Priority Nexus





JRC sites

Headquarters in Brussels and research facilities located in 6 locations **5 Member States**:

Belgium (Geel)

Germany (Karlsruhe)

Italy (Ispra)

The Netherlands (Petten)

Spain (Seville)





JRC Role: facts & figures

€ 386 million Budget annually, plus € 62 million earned

income

Independent of private, commercial or national interests

30% of activities in policy preparation, 70% in implementation125 instances of support to the EU policy-maker annually

More than **100** economic, bio-physical and nuclear models

Staff: **1500** core research staff (**83%** with PhD's), **3000** total staff



has no policy agenda of its own

42 large scale research facilities, more than 110 online databases

83% of core research staff with PhD's





Over 1,400 scientific publications per year



Directorate for Health, Consumers and Reference Materials



Director: E. Anklam

~ 300 Staff Members

~ 50 % female

Located in Geel, Belgium & Ispra, Italy

F.1 Health in Society



C. Nicholl Ispra Site

F.2 Consumer Products Safety



A.Hoeveler
Ispra Site

F.3 Chemical Safety and Alternative Methods



M. Whelan
Ispra Site

F.4 Food Fraud Detection



F. Ulberth
Geel Site
Ispra Site

F.5 Food and Feed Compliance



H. Emons
Geel Site
Ispra Site

F.6 Reference Materials



D. FlorianGeel Site





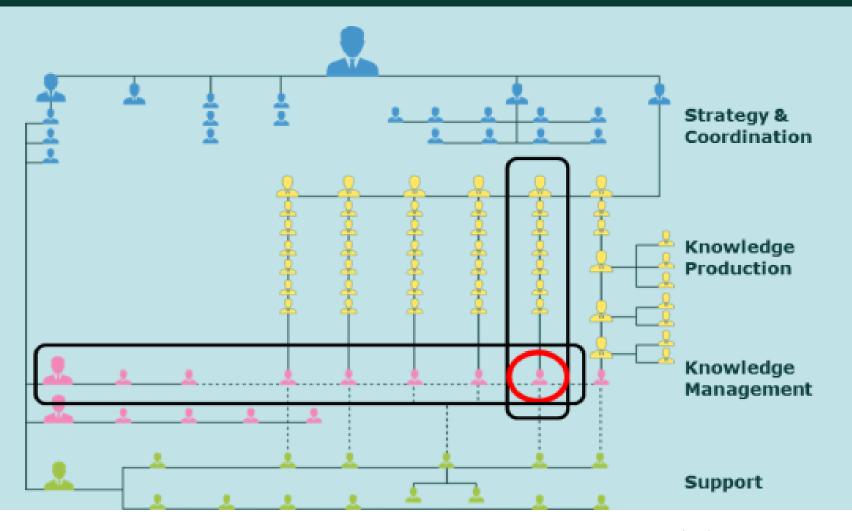
G. Van den Eede Geel Site Ispra Site





JRC.F.7 - Knowledge for Health & Consumer Safety Unit

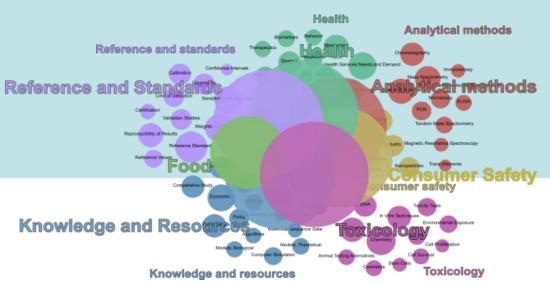
At the interface between Knowledge management and Knowledge production dimensions

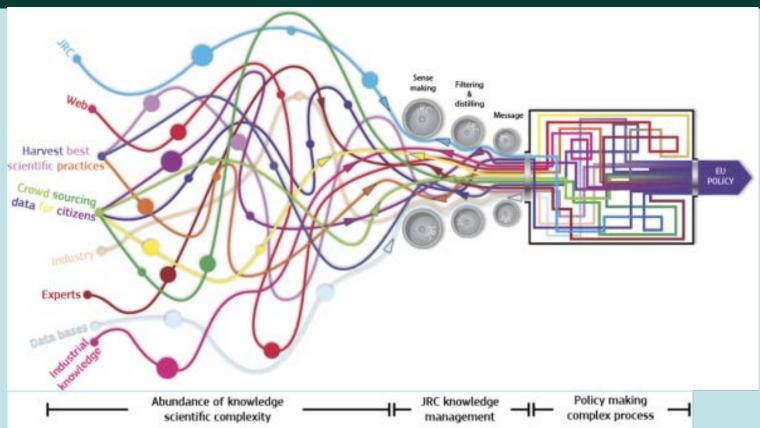




JRC.F.7 Unit role

- **✓** Harvest knowledge and competence
 - **✓** Exploit respective synergies
- ✓ Mapping policy areas
 - ✓ Anticipating policy-relevant knowledge needs
- ✓ Identifying gaps



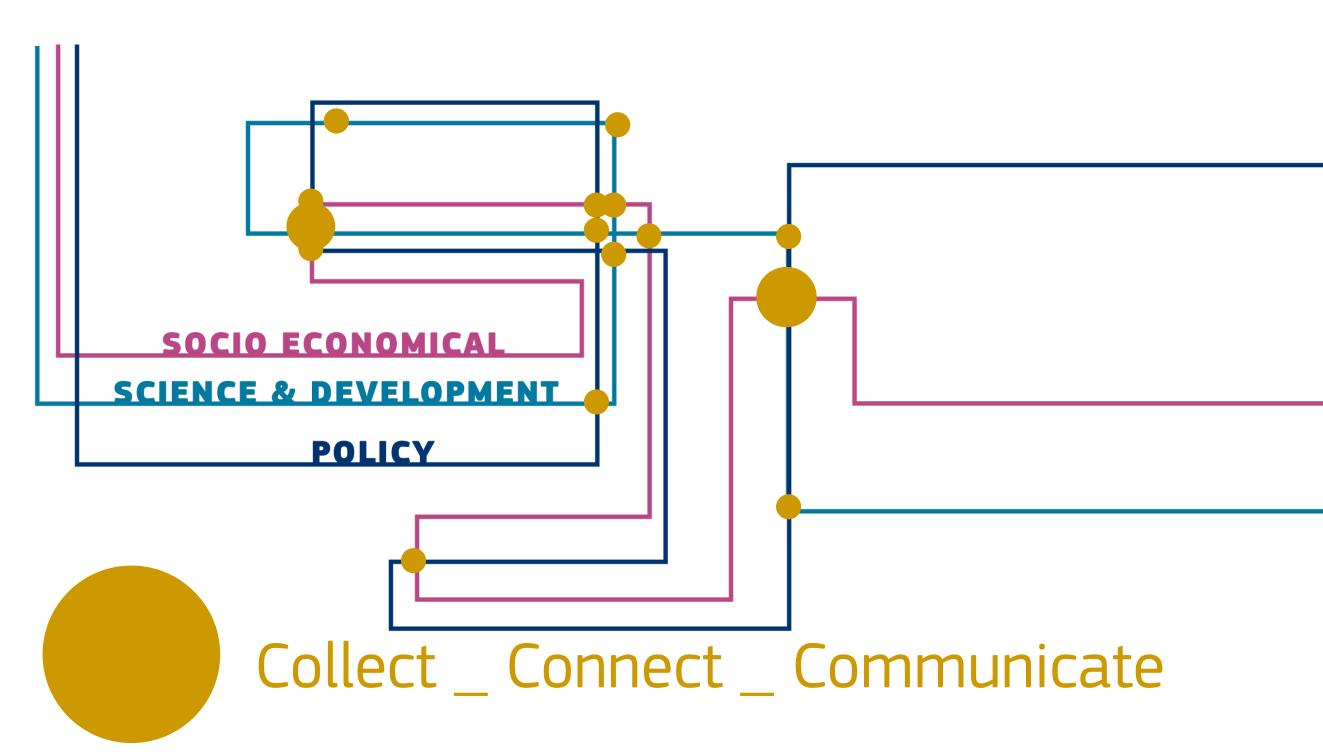


✓ Proposing, networking & contributing

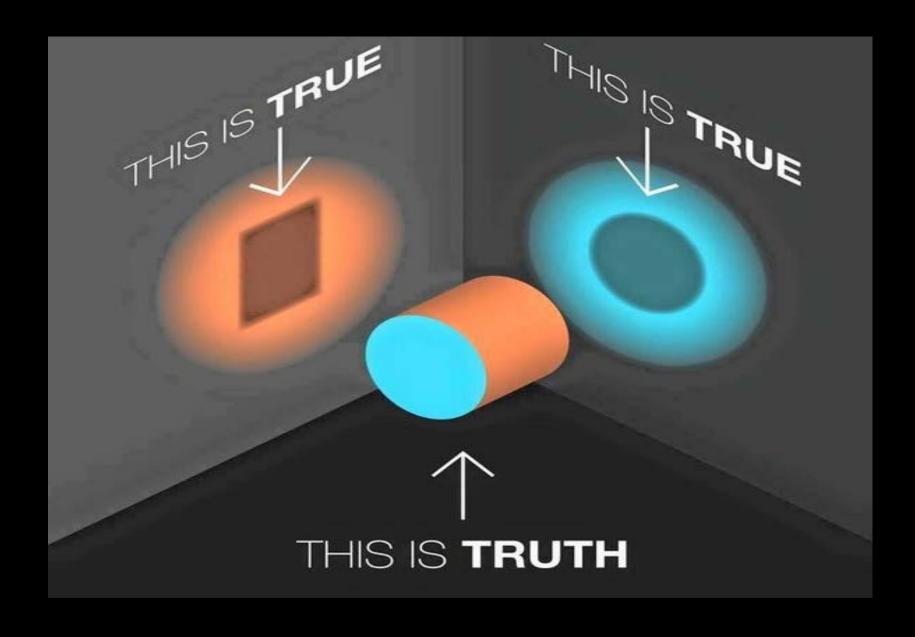
European

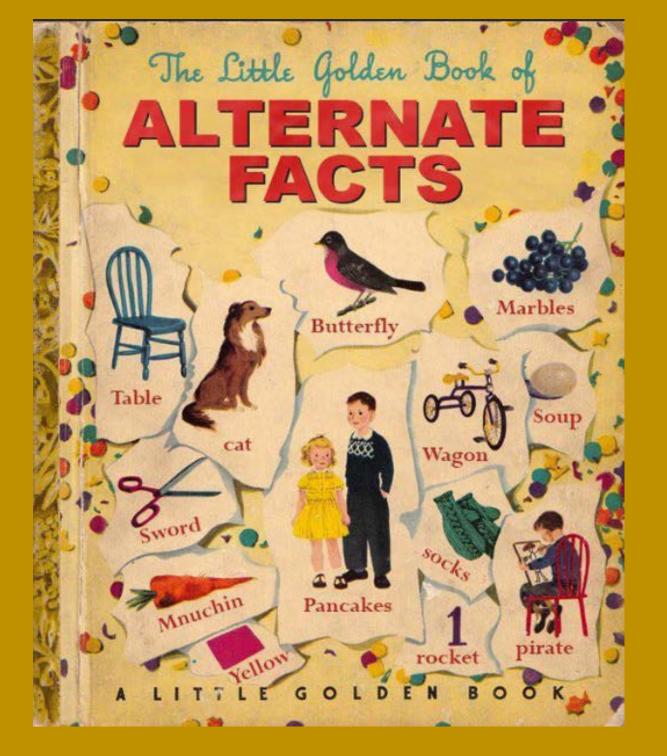
Commission

✓ Exploiting KM tools and approaches



Knowledge Management is about systems thinking

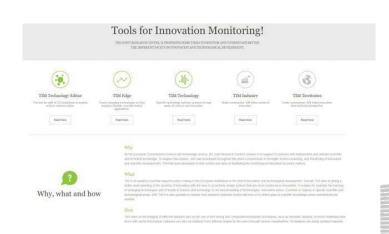






Selected examples – Data Sources

- Traditional scientific data sources:
 Scientific databases, patent databases, DG Research projects, but also e.g. bioRxiv (the preprint server for biology, Cornell Univ) and Europe PMC repository of life sciences articles and resources;
- European Media Monitor (EMM) enhanced for horizon scanning & foresight;
- Tools for Innovation Monitoring (TIM);
- Venture capital databases;
- Speech to text.







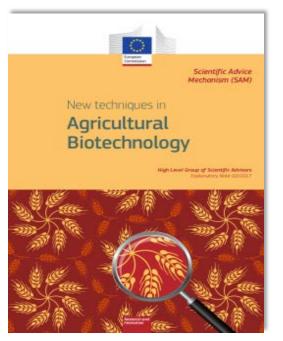
Selected examples

Support to the EC's Scientific Advice Mechanism (SAM)



The SAM's High Level Group requested help in managing the results of literature searches in the context of the development of the group's explanatory note "New Techniques in Agricultural Biotechnology"

F.7 staff developed strategies designed to process large sets of scientific references (such as those obtained by broad literature searches, in different databases) and assist in the identification of documents relevant for specific aspects of the final document.





Regulatory Bioinformatics

Data Policy

Data for Policy

"OMICS in Society" - JRC horizontal activity to provide necessary knowledge and/or services to EU policy makers in this complex and heterogeneous field

European Commission



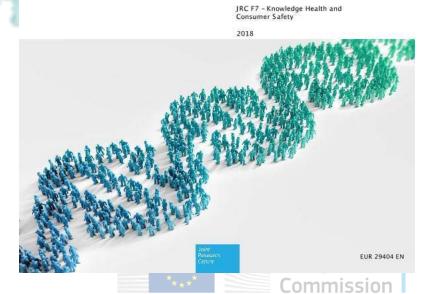


For the policy maker:

To understand and communicate the impact of new omics technologies (e.g. large scale genome sequencing), ensure the development of regulations that encourage innovation in line with bioethics and through a proper citizen dialogue



Overview of EU National Legislation on Genomics



- Identification of the challenges to implement Omics technologies
- Social science genomics & GWAS
- NGS x AMR across ≠ ecological compartments
- AMR genetic determinants databases

Collaborations

- The Global Alliance for Genomics and Health
- the European Molecular **Biology Laboratory (EMBL)**
- **EMBL-EBI**
- ELIXIR





EMB



For the research community: Provide high quality data & analysis tools for the design of solid evidence-based policies



Data for policy: data need to be reliable

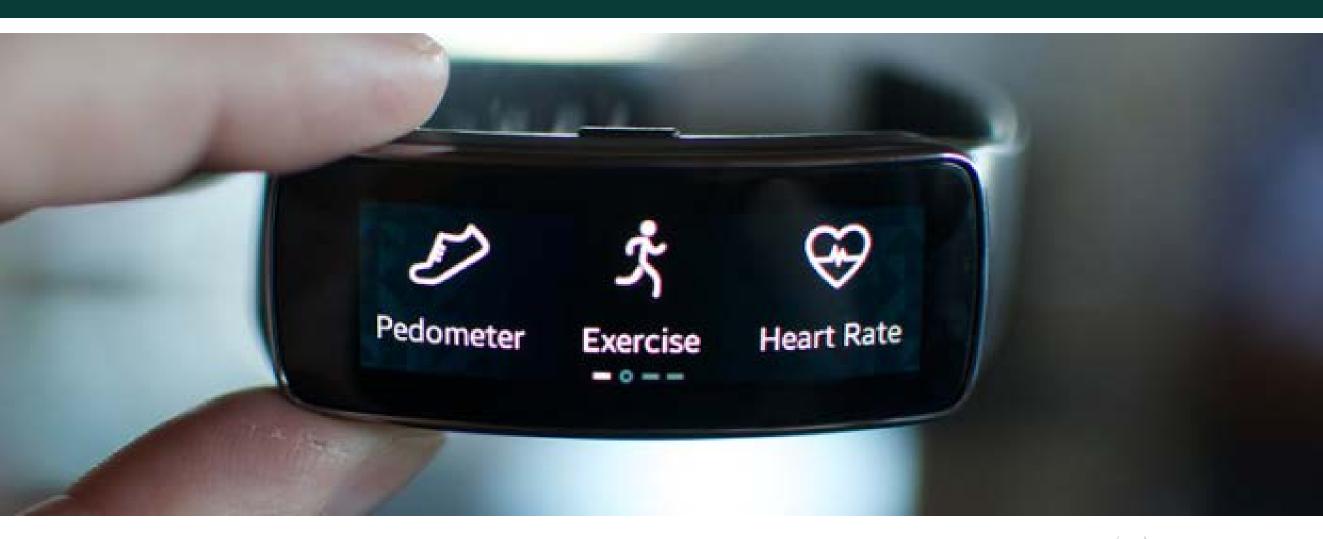
Two options:

Everyone does the same (no black boxes); simple problems – golden standards (cfr. diagnostics);

Everyone does things that perform the same (black boxes allowed); complex problem - different strategies - no clear "right" or "wrong".



Data for policy: health data & the digitisation of health





Different collections of different data sets







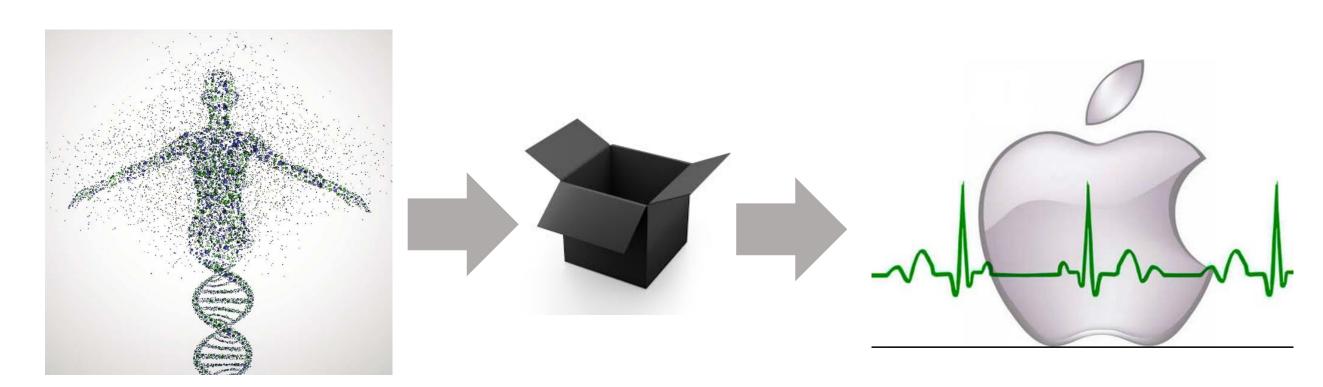












Patient benefit? Healthcare benefit? Equity? Ethics? Sustainability? Economics?





Commission Communication on enabling the digital transformation of health and care in the Digital Single Market; empowering citizens and building a healthier society (e.g. electronic health records)

Declaration of cooperation towards access to at least 1 million sequenced genomes in the European Union by 2022

"By pooling health data, using artificial intelligence and blockchain and promoting innovation, Europe can significantly improve people's lives. Earlier and better diagnosis of diseases, safer roads – this is only a glimpse of what embracing digital change can look like"





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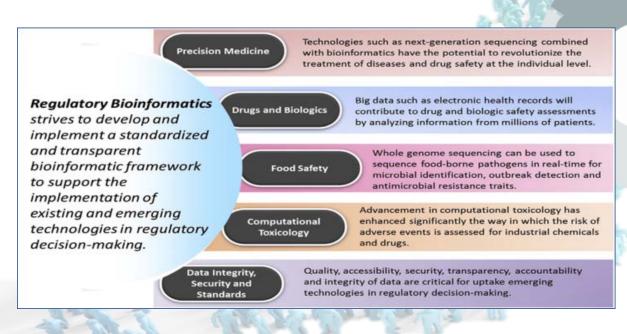
Declaration for delivering cross-border access to **genomic database**

- 1 million genomes accessible in the EU by 2022
- Linking access to existing and future genomic database across the EU
- Providing a sufficient scale for new clinically impactful results in research



Regulatory Bioinformatics

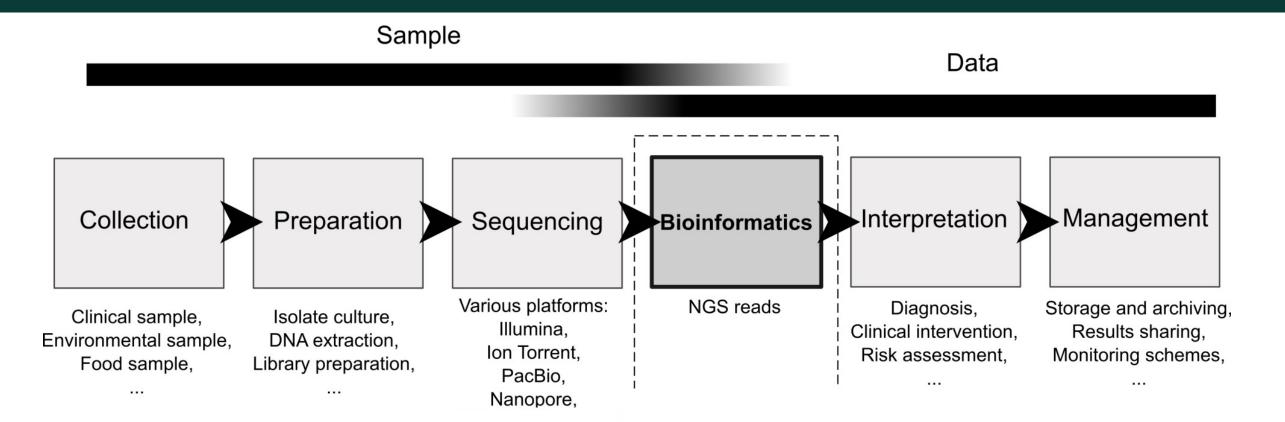
To develop and implement standardised bioinformatics frameworks fit for use in regulatory decision-making contexts



DOI: 10.1016/j.yrtph.2016.05.021

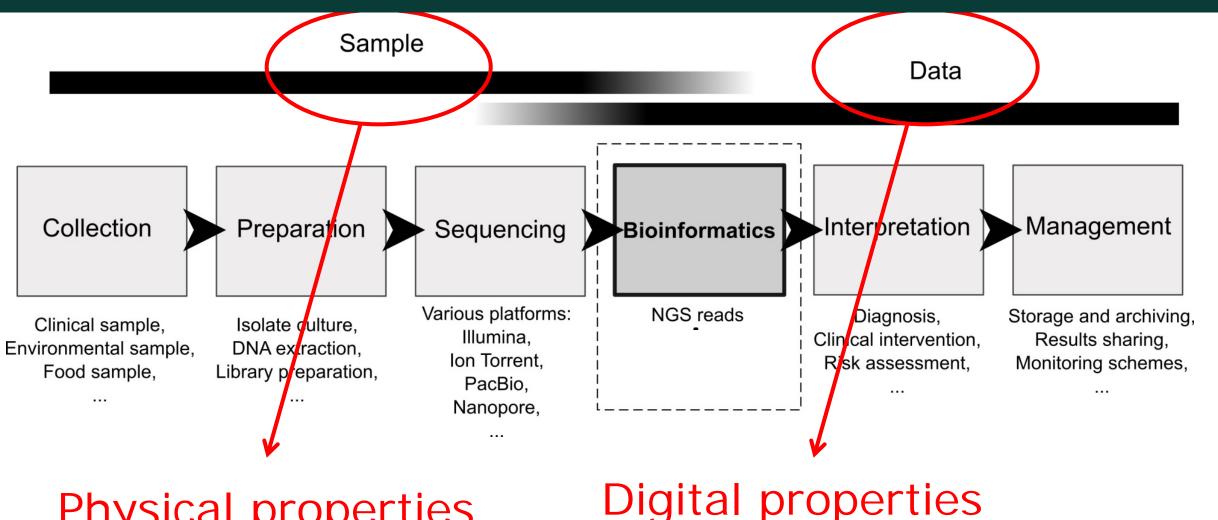


Analytical performance of NGS-based methods





Analytical performance of NGS-based methods



Physical properties



Analytical performance of NGS-based methods

A review of methods and databases for metagenomic classification and assembly

Florian P. Breitwieser, Jennifer Lu and Steven L. Salzberg

Briefings in Bioinformatics, 2017, 1-15

An even bigger issue than incorrect species labels is contamination.

The vast majority of genomes in GenBank today are 'draft' genomes. In any draft genome, some of the contigs might be contaminants, i.e. they might not belong to the species that was presumably sequenced, even though every contig is assigned to the same species. Common contaminants include human DNA, which creeps into many sequencing projects by accident. If the laboratory that created the assembly did not screen out these contaminants, they are submitted to GenBank as part of the organism. GenBank itself runs a contaminant screen on all assemblies, and contigs that appear to be contaminants are reported back to the submitter, who is encouraged to remove them and resubmit.

Despite the best efforts of GenBank curators, though, thousands of contaminants have already made their way into the draft genome data.



Do we have the equivalent for NGS?

LAMBERT ET AL.: JOURNAL OF AOAC INTERNATIONAL VOL. 100, No. 3, 2017 721

FOOD BIOLOGICAL CONTAMINANTS

Baseline Practices for the Application of Genomic Data Supporting Regulatory Food Safety

"Laboratories should use these validation and verification activities to determine whether bioinformatics workflows conform to the requirements of a given activity and whether the software satisfies its intended use and user needs."







OPINION ARTICLE

The challenges of designing a benchmark strategy for bioinformatics pipelines in the identification of antimicrobial resistance determinants using next generation sequencing technologies [version 1; referees: 2 approved]

Alexandre Angers-Loustau ¹ , Mauro Petrillo ¹ , Johan Bengtsson-Palme ¹ , Thomas Berendonk⁴, Burton Blais⁵, Kok-Gan Chan^{6,7}, Teresa M. Coque⁸, Paul Hammer⁹, Stefanie Heß⁴, Dafni M. Kagkli¹, Carsten Krumbiegel⁹, Val F. Lanza⁸, Jean-Yves Madec¹⁰, Thierry Naas¹¹, Justin O'Grady¹², Valentina Paracchini¹, John W.A. Rossen¹³, Etienne Ruppé¹⁴, Jessica Vamathevan ¹ , Vittorio Venturi¹⁶, Guy Van den Eede¹⁷





Thank you for your attention!!

