

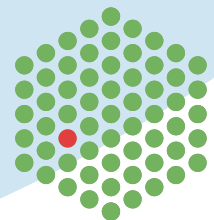


Scientific Cloud Computing Infrastructure for Europe - Strategic Plan

Rupert Lueck
Head of IT Services
EMBL Heidelberg

5th DFN Forum
May 21th 2012, Regensburg

EMBL



Helix Nebula

- “Eye of God”
- NGC 7293
- located inside Aquarius constellation
- One of the closest planetary nebula
- ~700 light-years away



Origin of the initiative

- Conceived by ESA as a prospective for providing cloud services to space sector in Europe
- Presented to the IT working group of the EIROforum where other members (CERN, EMBL) joined
- Two workshops held during 2011
 - June: hosted by ESA in Frascati
 - October: hosted by EMBL in Heidelberg



Strategic Plan

for a scientific Cloud Computing Infrastructure in Europa

- Establish a sustainable multi-tenant cloud computing infrastructure in Europe
- Initially based on the needs for the European Research Area & space agencies
- based on commercial services from multiple IT industry providers
- which adhere to internationally recognised policies and quality standards to be adopted by the governance structure involving all stakeholders

Lengert, Maryline, Jones, Robert (2011) CERN-OPEN-2011-036
<http://cdsweb.cern.ch/record/1374172/>

Objectives of the initiative

1. Set up a cloud computing infrastructure for all European Research Area
2. Identify and adopt policies for trust, security and privacy on a European-level
3. Create a light-weight governance structure involving all stakeholders
4. Define a short and medium term funding scheme

A Collaboration Initiative

**European Commission
& relevant projects**

**User organisations
*Demand-side***

**European
Cloud Computing
Strategy**

**Commercial Service
Providers
*Supply-side***

Bringing together all the stakeholders to establish a **public-private partnership**

Timeline

Set-up
(2011)

Pilot phase
(2012-2014)

Full-scale
cloud service
market
(2014 ...)

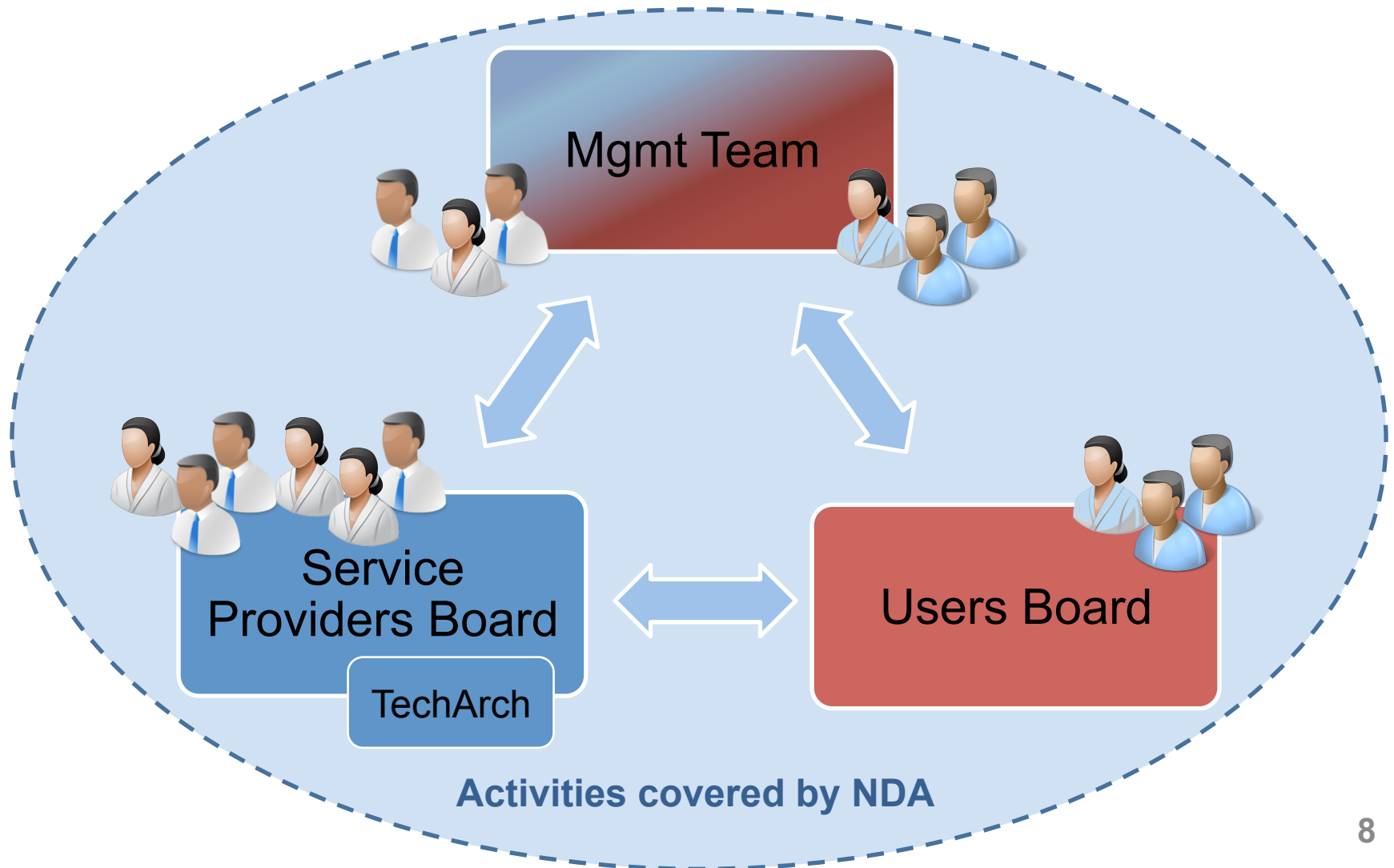
- Select flagships use cases
- Identify service providers
- Define governance model

- Deploy flagships
- Analysis of functionality, performance & financial model
- Success Stories

- More applications
- More services
- More users,
- More service providers

Governance Model

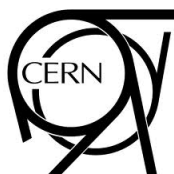
Proof of Concept stage



Consortium

Q1 / 2012

Demand-Side



Supply-Side



Activities covered by NDA

Consortium membership

- Consortium includes all participating supply-side and demand-side companies / organisations
 - Member status and adopter status
 - All sign a non-disclosure agreement

- Initial membership is defined
 - More members and adopters will be added following the Proof of Concept stage within the Pilot Phase (summer 2012)

Pilot Phase

Explore / push a series of perceived barriers to Cloud adoption:

- **Security:** Unknown or low compliance and security standards
- **Reliability:** Availability of service for business critical tasks
- **Data privacy:** Moving sensitive data to the Cloud
- **Scalability / Elasticity:** Will the Cloud scale-up to our needs
- **Network performance:** Data transfer bottleneck; QoS
- **Integration:** Hybrid systems with in-house / legacy systems
- **Vendor lock-in:** Vendor dependency once data & applications are transferred to the Cloud
- **Legal concerns:** Such as who has legal liability
- **Transparency:** Clarity of conditions, terms and pricing

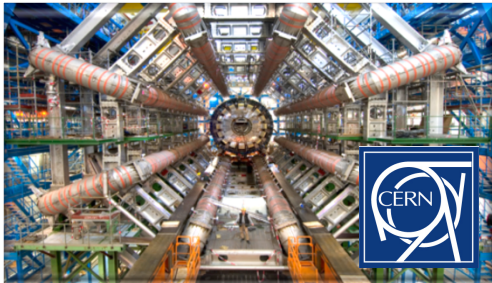
Flagship use cases

- Proposed by demand-side user organisations
- Addressing scientific challenges with societal impact
 - High-profile applications
 - Catching the public imagination & encourage others to use the services
 - Innovate in terms of functionality, performance, scope, business opportunities or impact
- Sponsored by user organisations
 - Need to contribute their own resources during the pilot phase to port application (**manpower**) and contribute to the cost of procuring required services from the supply-side (**cash**)
 - Must **participate in a costing exercise** where the total cost of deploying and operating the flagship application in-house can be compared to the cost of procuring the services via Helix Nebula
- Want to propose a flagship?
 - Send email to contact@helix-nebula.eu



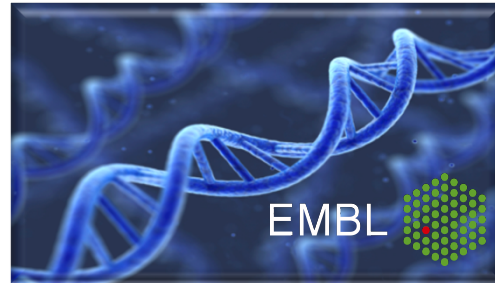
Initial flagships use cases

ATLAS High Energy Physics Cloud Use



To support the computing capacity needs for the ATLAS experiment

Genomic Assembly in the Cloud



A new service to simplify large scale genome analysis; for a deeper insight into evolution and biodiversity

SuperSites Exploitation Platform



To create an Earth Observation platform, focusing on earthquake and volcano research

Call for proposals

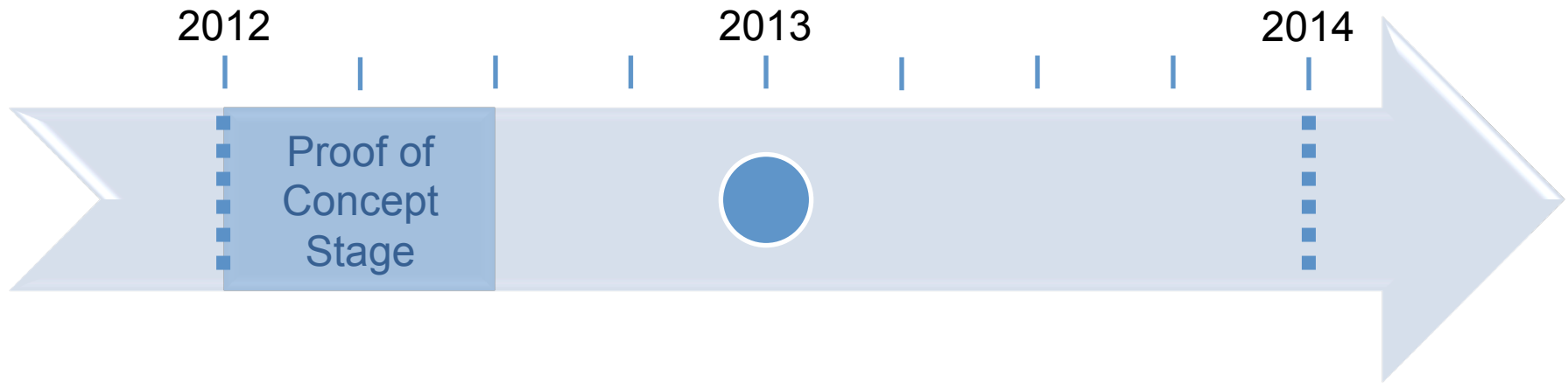
- Template agreed by demand and supply side
- Eligibility review and analysis with cloud service suppliers

Flagship use cases

	ATLAS H.E.P. Cloud Use (CERN)	Genomic Assembly in the Cloud (EMBL)	SuperSites Exploitation Platform (ESA/CNES/DLR)
Scientific goal, society impact, photogenic	✓	✓	✓
Scale of resources used	✓	✓	
Federation / aggregation of datasets		✓	✓
Long-term archiving of data			✓
On-demand processing	✓	✓	✓
Impact on community & benefits	✓	✓	✓
Potential increase of users	✓	✓	✓
Interoperability	✓	✓	✓
Data security	✓	✓	✓
Maturity	✓	✓	✓
Access to license-controlled software			✓

Flagship deployments

Pilot Phase



- Proof of Concept (PoC) stage started in January 2012
- Each flagship will be deployed with a series of providers independently
- Sequence: CERN-ATLAS, EMBL & ESA
- Initial PoC expected to be completed by summer 2012

Flagship use cases Participating Suppliers

Atos

CloudSigma 

interoute 
from the ground to the cloud

logica
be brilliant together

terradue 20 

sixsq

...T...Systems

the
SERVER
LABS

the IT architects

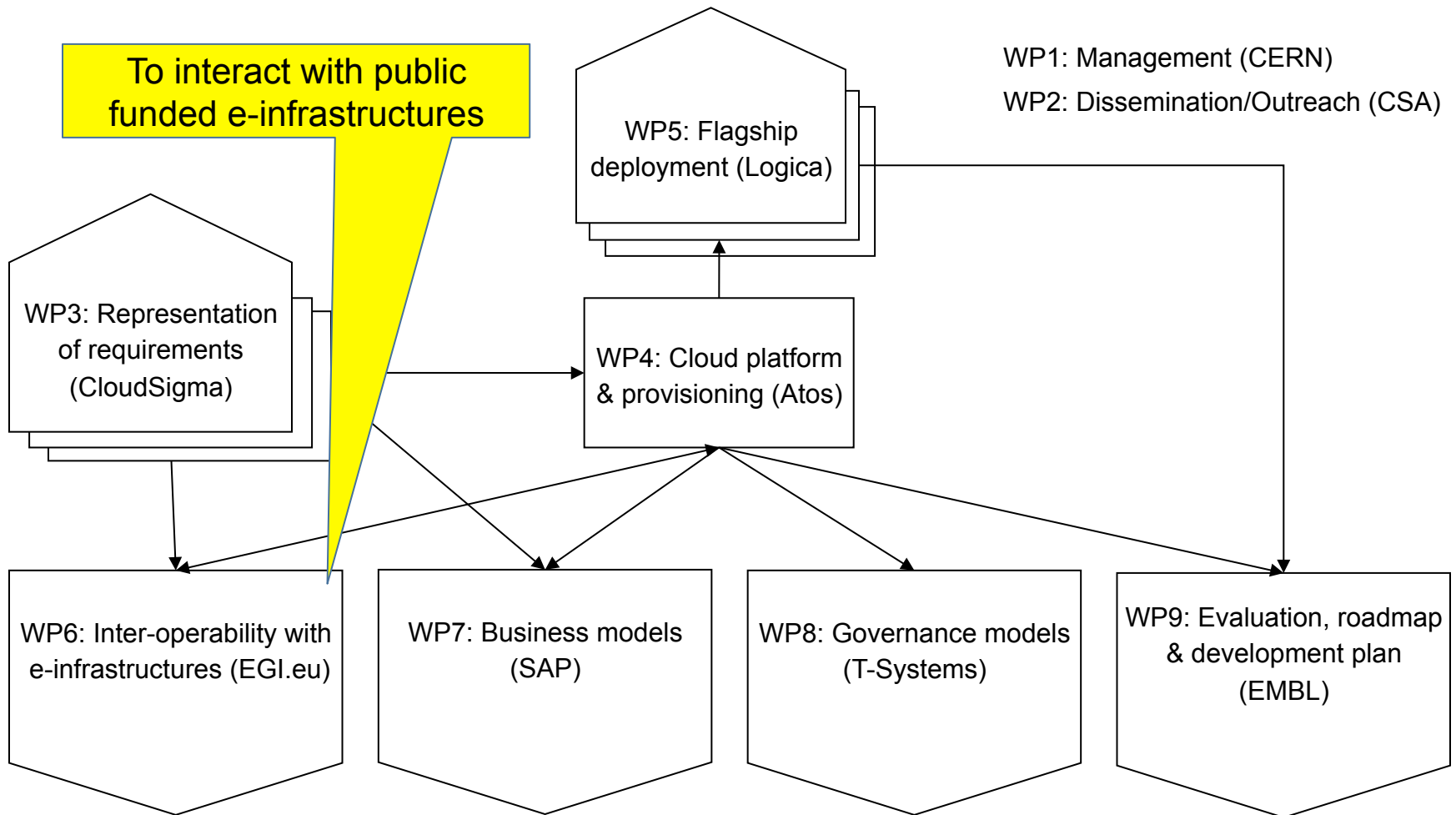
Helix Nebula EC project

Coordination action under call INFRA-2012-3.3

- Start-date 1st June 2012, duration 24 months
- Total budget ~3M€ (1.8M€ EC funding)

	Short name	Organisation	Country
1	CERN	European Organization for Nuclear Research (coord.)	CH
2	EGI.eu	STICHTING EUROPEAN GRID INITIATIVE	NE
3	EMBL	European Molecular Biology Laboratory	DE
4	Atos	ATOS	NE
5	T-Systems	T-Systems International GMBH	DE
6	CloudSigma	CloudSigma AG	CH
7	SAP	SAP AG	DE
8	Logica	Logica Deutschland GmbH & Co KG	DE
9	CNR	CONSIGLIO NAZIONALE DELLE RICERCHE	IT
10	CSA	Cloud Security Alliance Europe	UK

Helix Nebula proposal



Relevance of Helix Nebula for Network Community



- Helix Nebula to build hybrid cloud (public & private data centres)
- Research community moving towards using commercial cloud services (Helix Nebula not the only initiative in this domain)
- E-IRG response to GEANT 2020 vision paper:
Importance of participation of private research in the use of research networks
(http://www.e-irg.eu/images/stories/e-irgs_reaction_geg_a5.pdf)

e-IRG: e-Infrastructure Reflection Group

Relevance of Helix Nebula for Network Community (2)

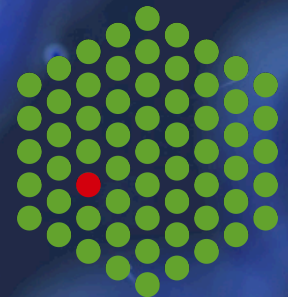
Helix Nebula pilot phase provides

- Opportunity for NRENs & network community
- to work with the research communities and commercial cloud service providers to deploy flagship applications
- to evaluate ability of NRENS to offer access to commercial data centres
- to investigate how a public-private cloud serving the research community could exist

Genomic Assembly in the Cloud

EMBL Flagship Use Case

EMBL



EMBL: European Molecular Biology Laboratory



- Intergovernmental Research Organization
- Supported by 20 Member States (+1 associated: )
- One of the world's foremost life science institutions
- EIROforum member
- 1500 staff
>70 nationalities

The Five Branches of EMBL



Heidelberg
Basic Molecular Biology
Research
Main Lab / Headquarters



Hamburg
Structural Biology
DESY



Hinxton
European Bioinformatics
Institute (EBI)
Sanger Centre

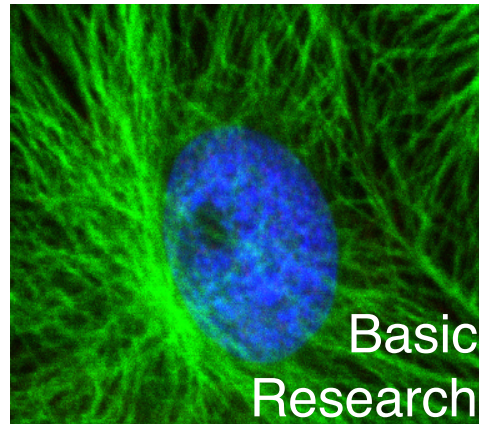


Grenoble
Structural Biology
ILL, ESRF, IBS, UVHCI

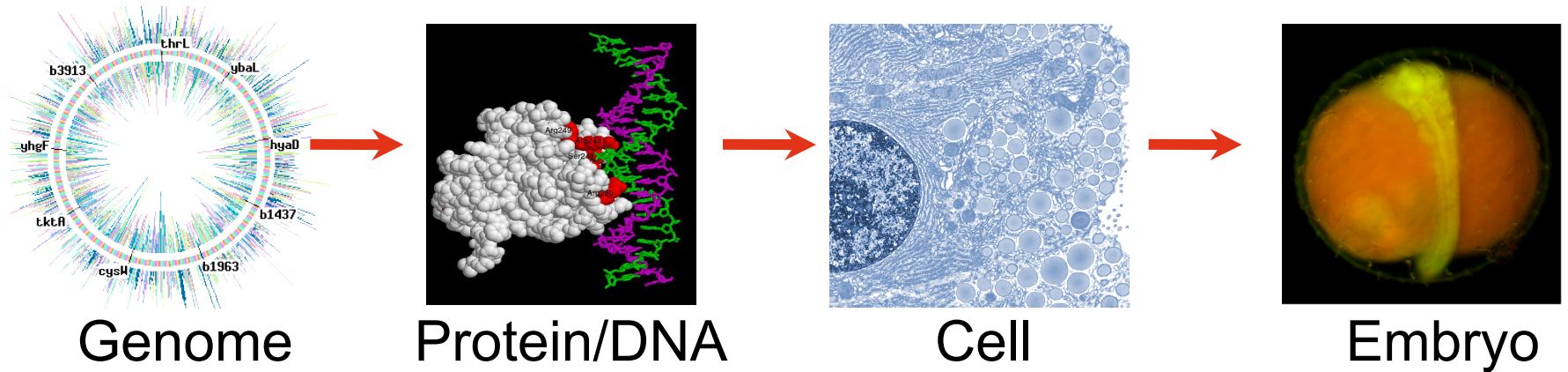


Monterotondo
Mousebiology
CNR, EMMA

EMBL's Missions



Systems Biology: From Molecules to Organisms



Development



Organisms

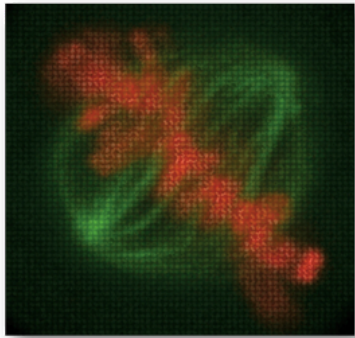


Complexity

Aging

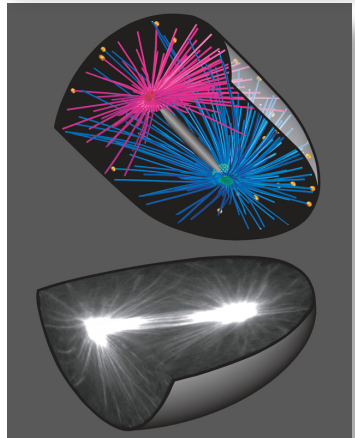
Disease

Research Directions & Key Technologies



Imaging

- Bridging scales of biological organisation: combine low- and high-resolution techniques
- Biology in four dimensions: live imaging to study dynamic processes in space and time
- Generate quantitative data



Computational Biology

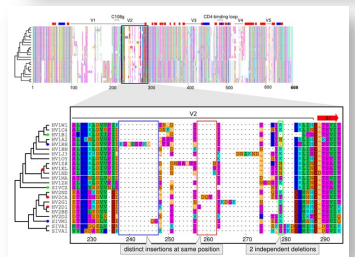
- Analysing, integrating and exploiting quantitative data
- Build predictive networks and models of biological processes

Next Generation Sequencing

- Inter-species variation: comparative sequence analysis to study evolution
- Intra-species variation: link genetic variation to phenotype

Disease Models and Mechanisms

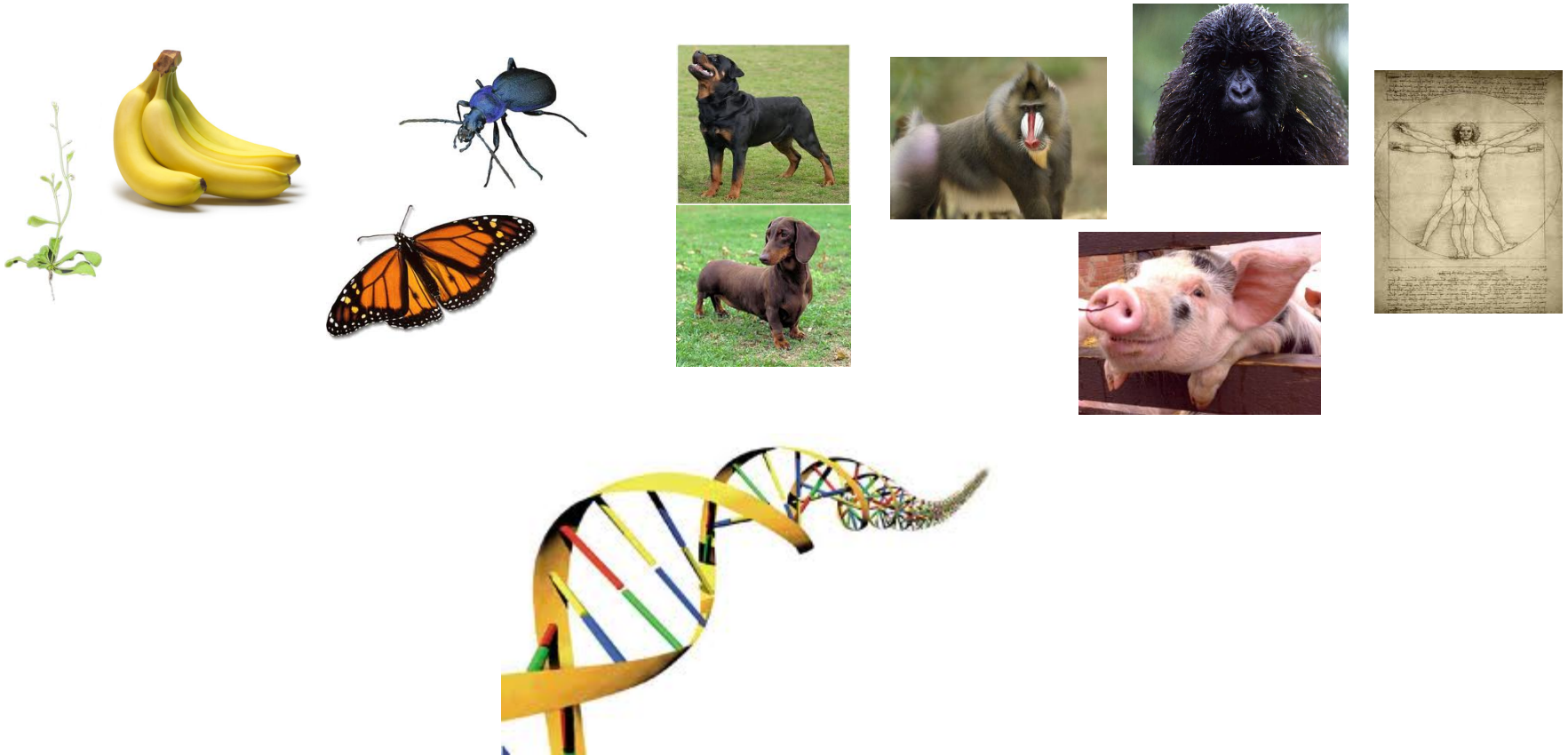
- Decipher the molecular basis of genetic and infectious diseases with the help of animal and cellular models



Exemplary Big Data Challenge

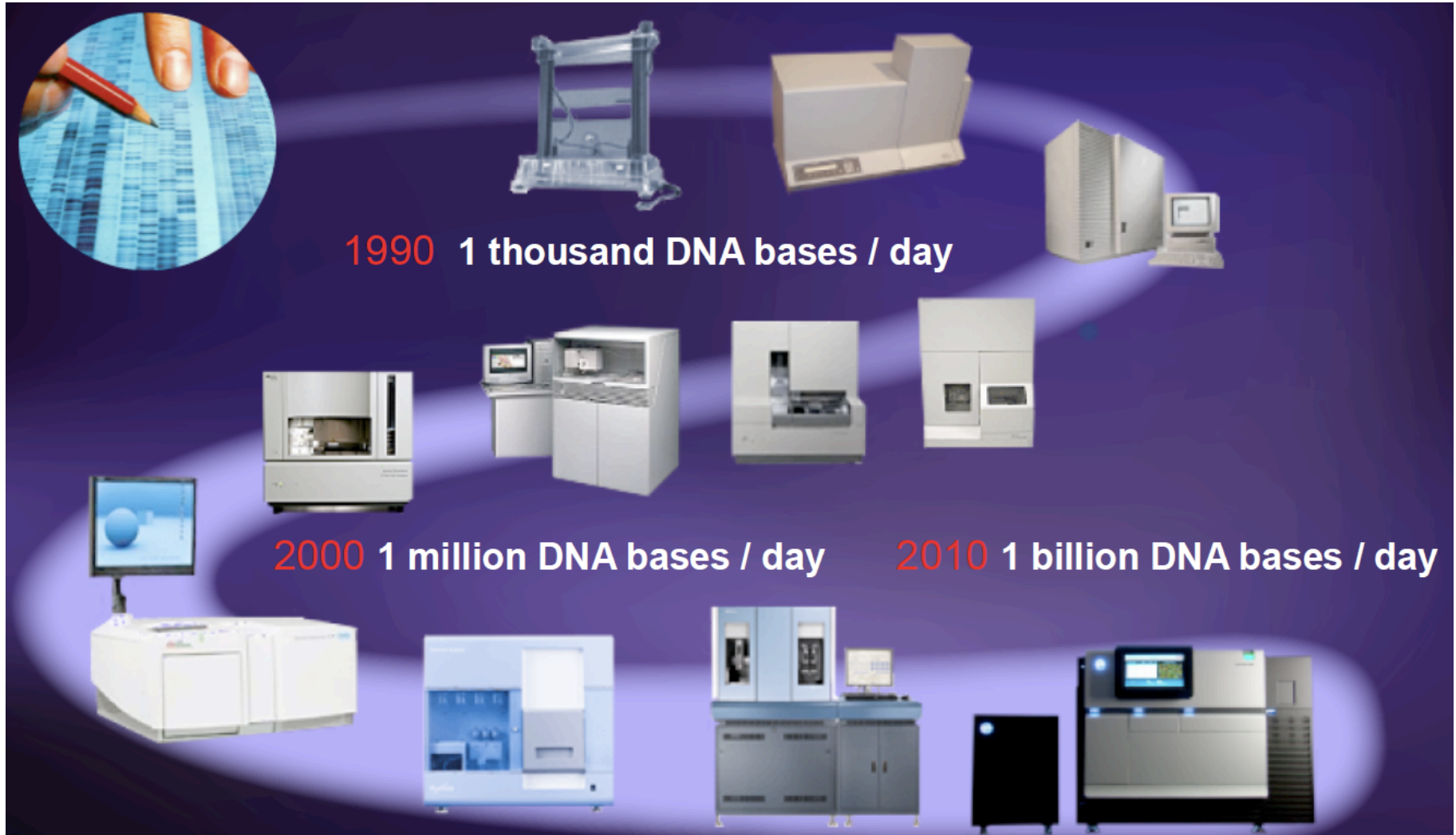
NEXT GENERATION SEQUENCING (NGS)

DNA and Life on Earth



The Sequence Holds the Code for the Organism

Next Generation Sequencing (NGS) Revolution



NGS Impact on Human Genome Sequencing

- Human genome project
 - 10 years
 - Large International Consortium
 - Thousands of Sequencers
 - \$3,000,000,000

- Sequencing today
 - < \$10,000
 - A few hours
 - One machine

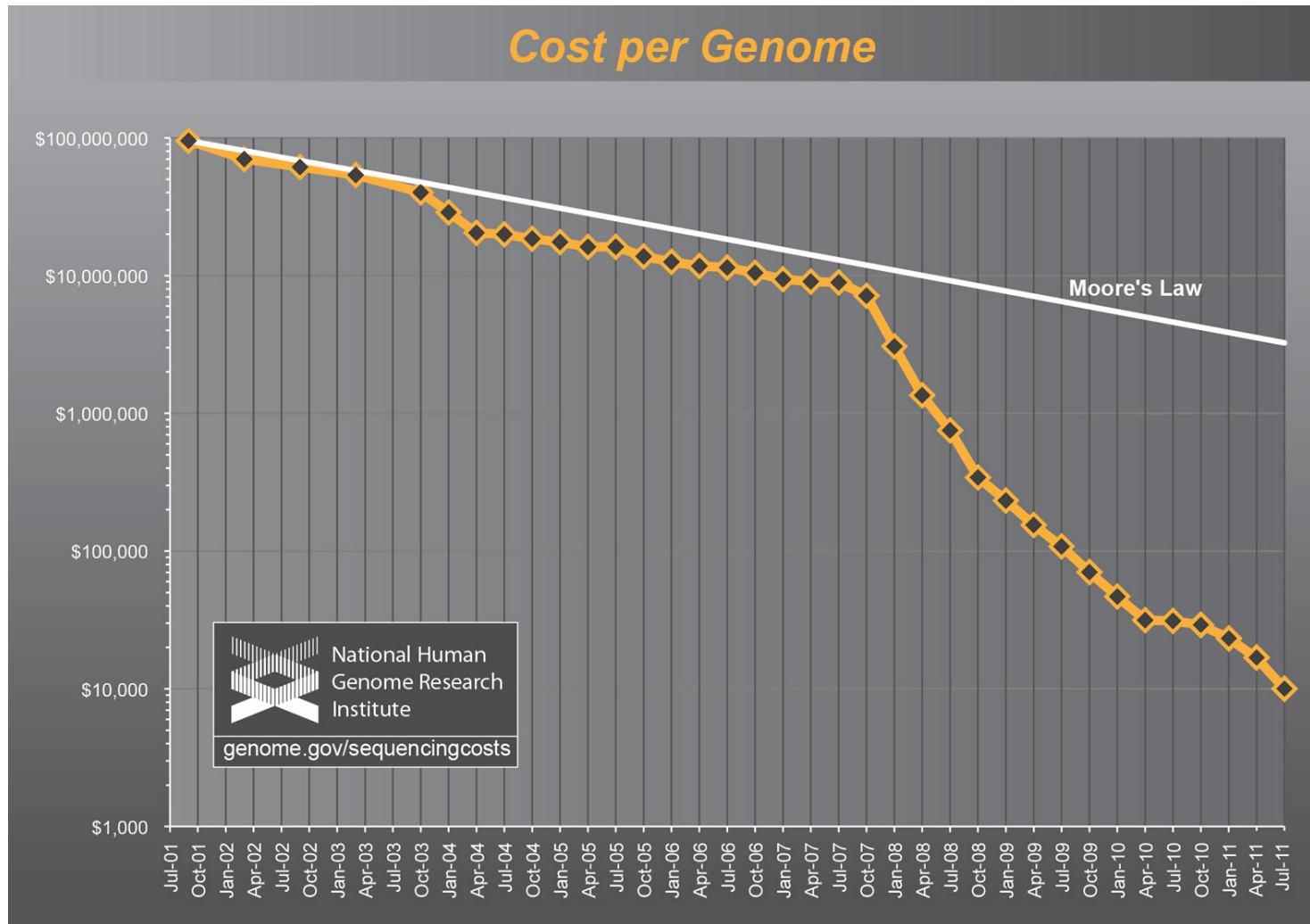
2000



2010



Cost of Sequencing Decreasing Rapidly

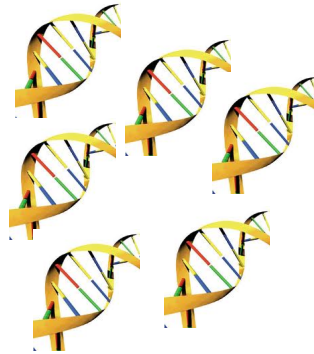


Read the Sequence to Study the Organism

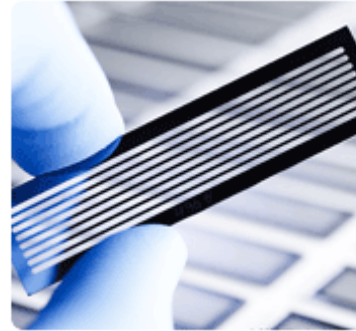
Extract DNA



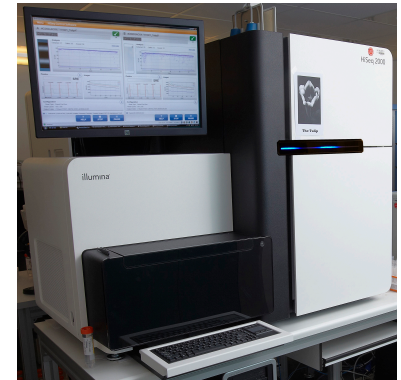
Fragment



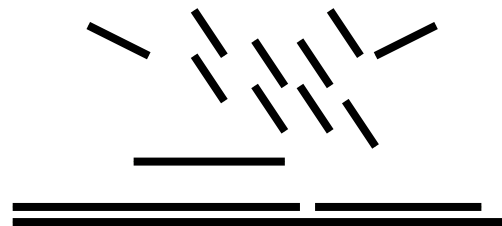
Prepare



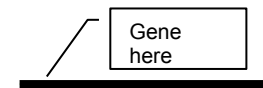
Sequence



Assemble



Annotate



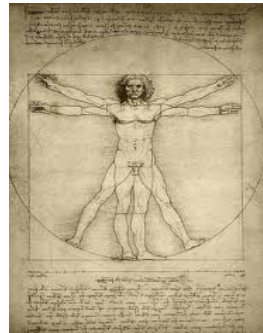
Requires Computing Infrastructure & Expertise

Genomic Sequencing is Now an Affordable Solution

Academic
Research
Groups



Medical
Research



Pharmaceutical
Companies



Agricultural
Research



Genomic Sequencing is Now an Affordable Solution

Academic
Research
Groups

Medical



The screenshot shows the ICGC website with the following elements:

- Header:** "1000 Genomes" and "A Deep Catalog of Human Genomes".
- Navigation:** "Home", "About", "Data", "A".
- Search:** "Enter keywords" input field and "Search" button.
- ICGC Logo:** International Cancer Genome Consortium.
- Menu:** "Home", "Database & Species Lists", "News", "Events", "Publications", "Participants", "For G10K Organizers (restricted)".
- Main Content:** "GENOME 10K. Unveiling animal diversity". Includes a search bar with "Search:" and "Go" buttons.
- Text:**

Genome 10K Project

To understand how complex animal life evolved through changes in DNA and use this knowledge to become better stewards of the planet.

The Genome 10K project aims to assemble a genomic zoo—a collection of DNA sequences representing the genomes of 10,000 vertebrate species, approximately one for every vertebrate genus. The
- Buttons:** "Join us", "Become a G10K affiliate".
- Sidebars:** "ICGC Committee", "Sort by:", "Bladder United States".

Genomic sequencing is
now an affordable solution

but ...

Problem – 1: Assembly

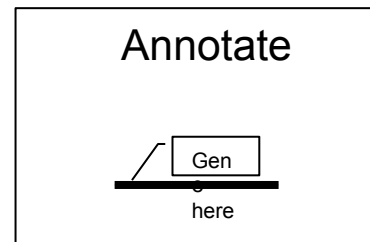
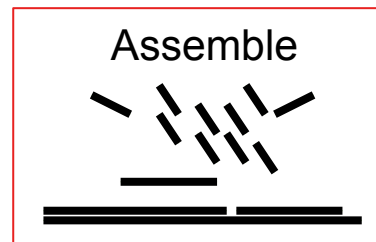
ATGCATT... 200,000,000 ...TGCGGATC

Genomes contain long strings of bases

ATGCATT... 105 ... GTATTCC

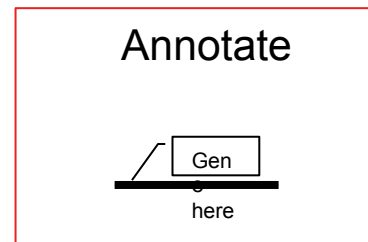
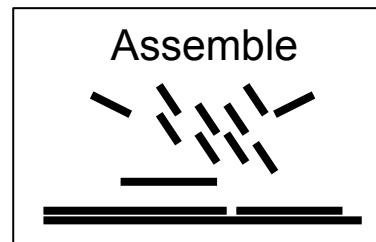
NGS output: millions of very short sequence reads

- The short reads have to be assembled into genomes
- Up to 1TB RAM required to solve puzzle

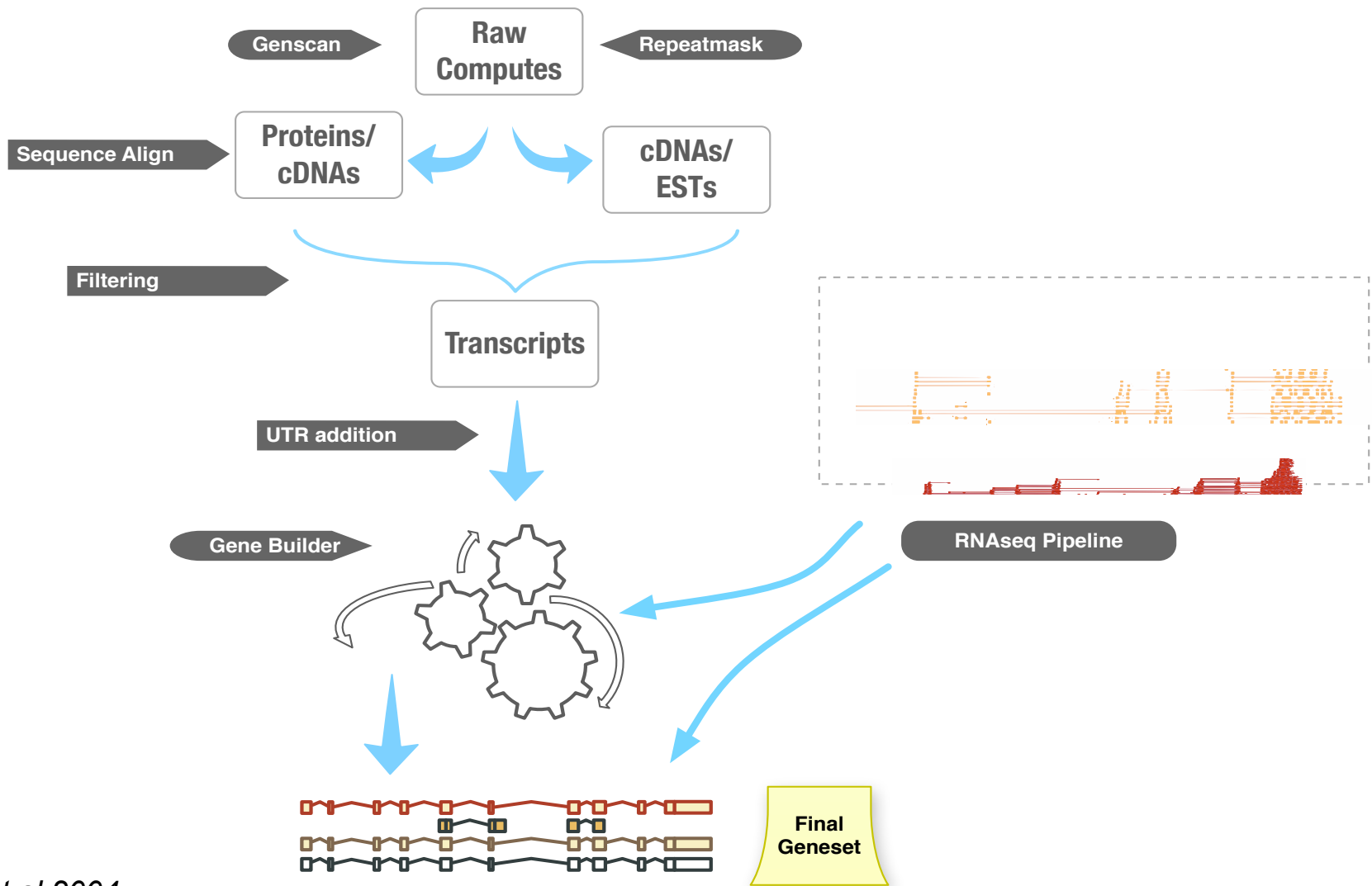


Problem – 2: Annotation

- Strings of assembled bases need to be annotated
- Document features inside the code
- 3 billion bases ~25k genes
- Looking for genes, gene and promoter sequences
- Requires multiple pipelines and databases



e! Ensembl Annotation Pipeline



Potter et al 2004

Computational Steps Involved

Upload Data

1TB sequence data

→ QC and Filter data

Assemble Sequences

→ Large Graphs **1TB RAM**

Multicore processing

Repeat
With different
parameters

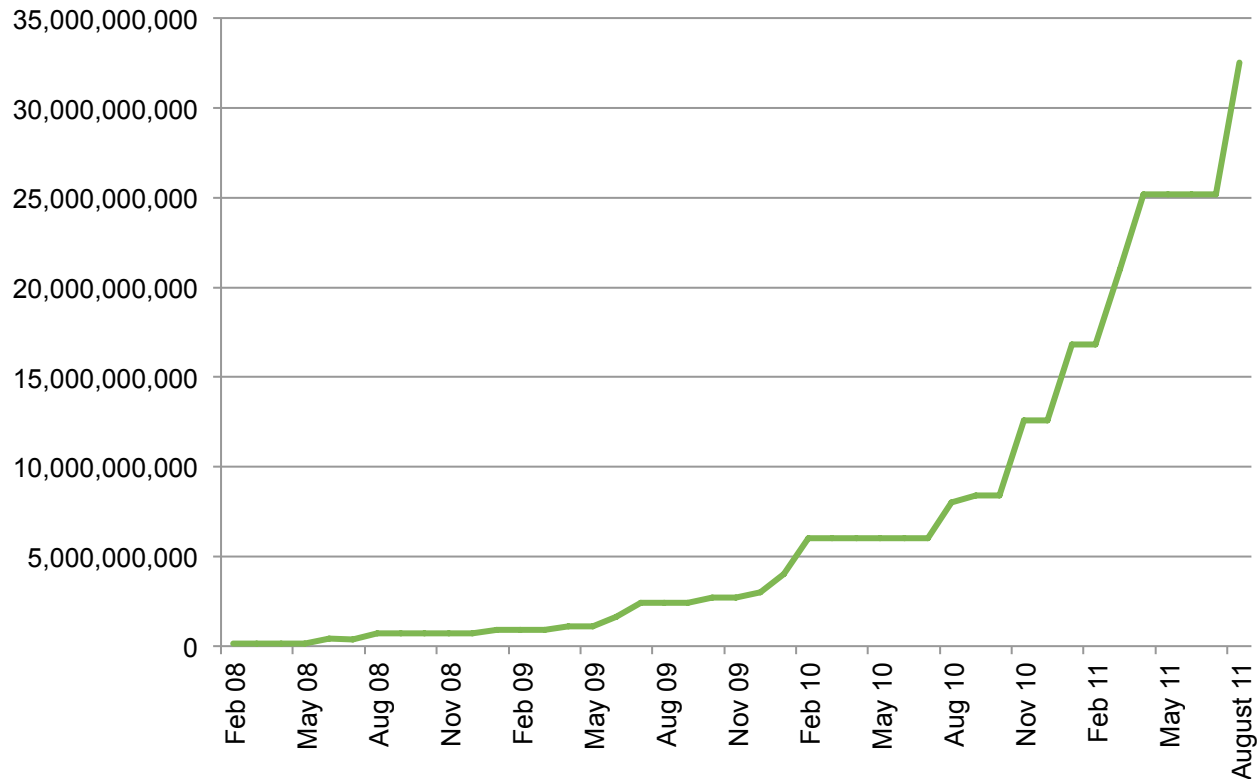
x Weeks

→ Annotate Assembly

→ Download Annotated
Assembly 100GB

Problem - Technology Explosion with NGS

**Bases Sequenced / Sample / Run @ EMBL
(Illumina)**

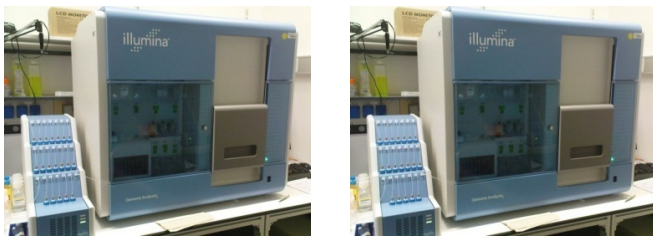


Sequence Production & IT Infrastructure at EMBL

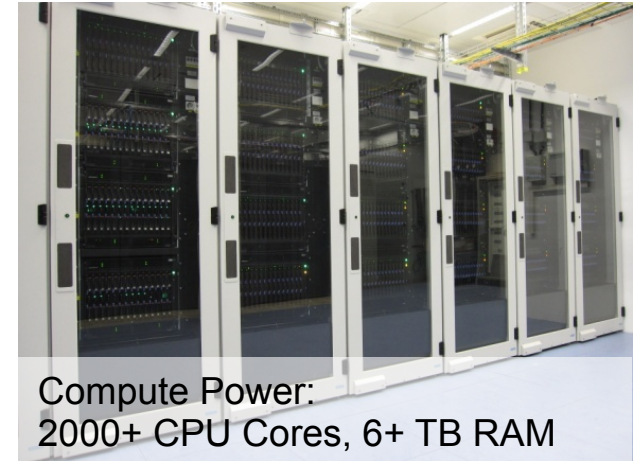
4 x Illumina HiSeq2000



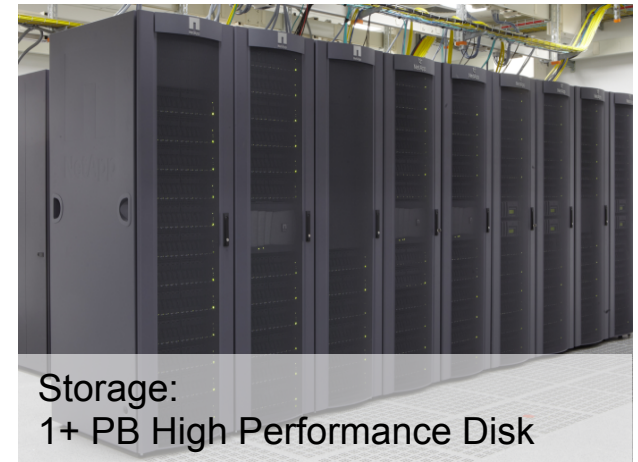
2 x Illumina GAIIx



25 TB data
each week



Compute Power:
2000+ CPU Cores, 6+ TB RAM

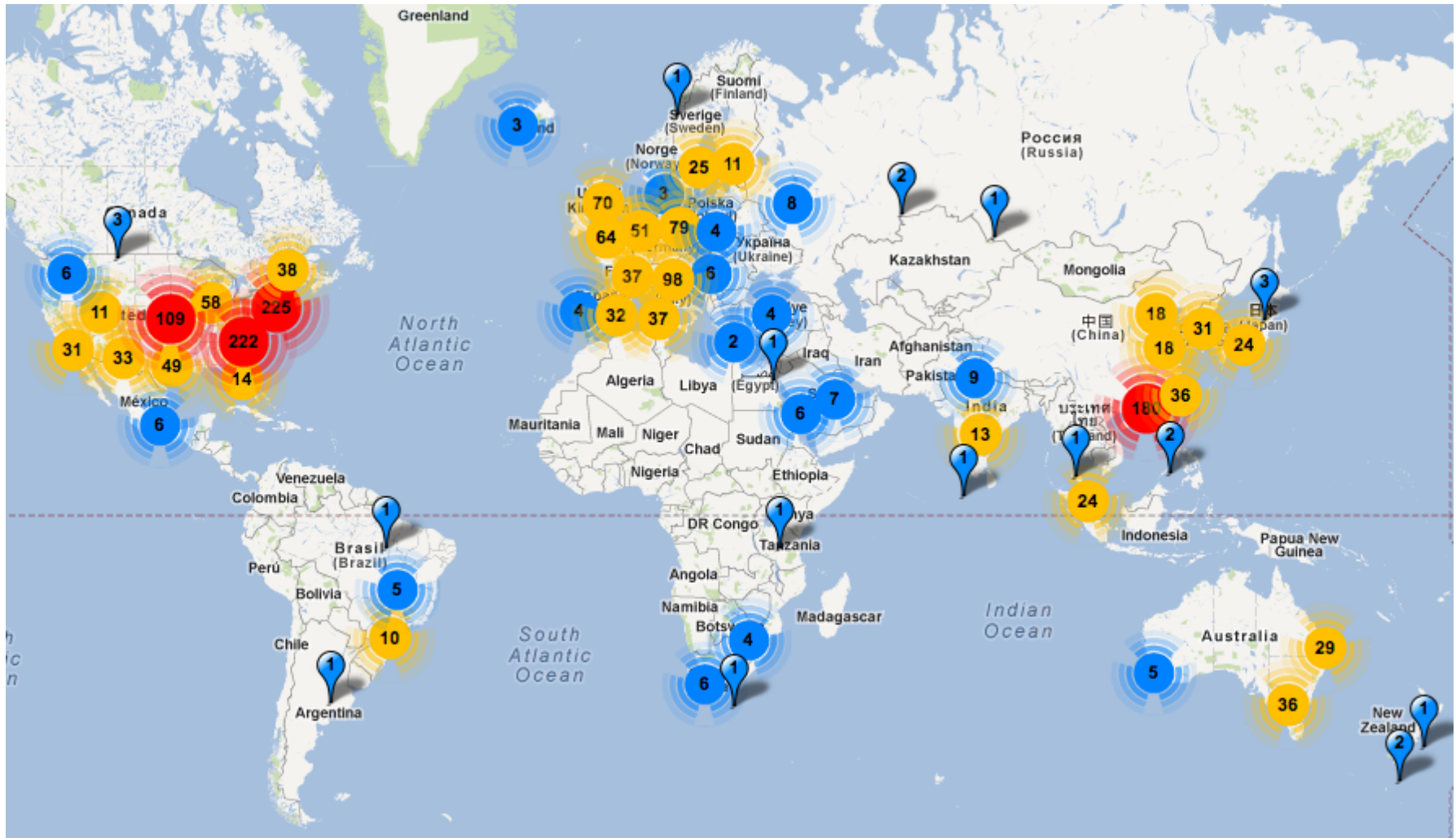


Storage:
1+ PB High Performance Disk

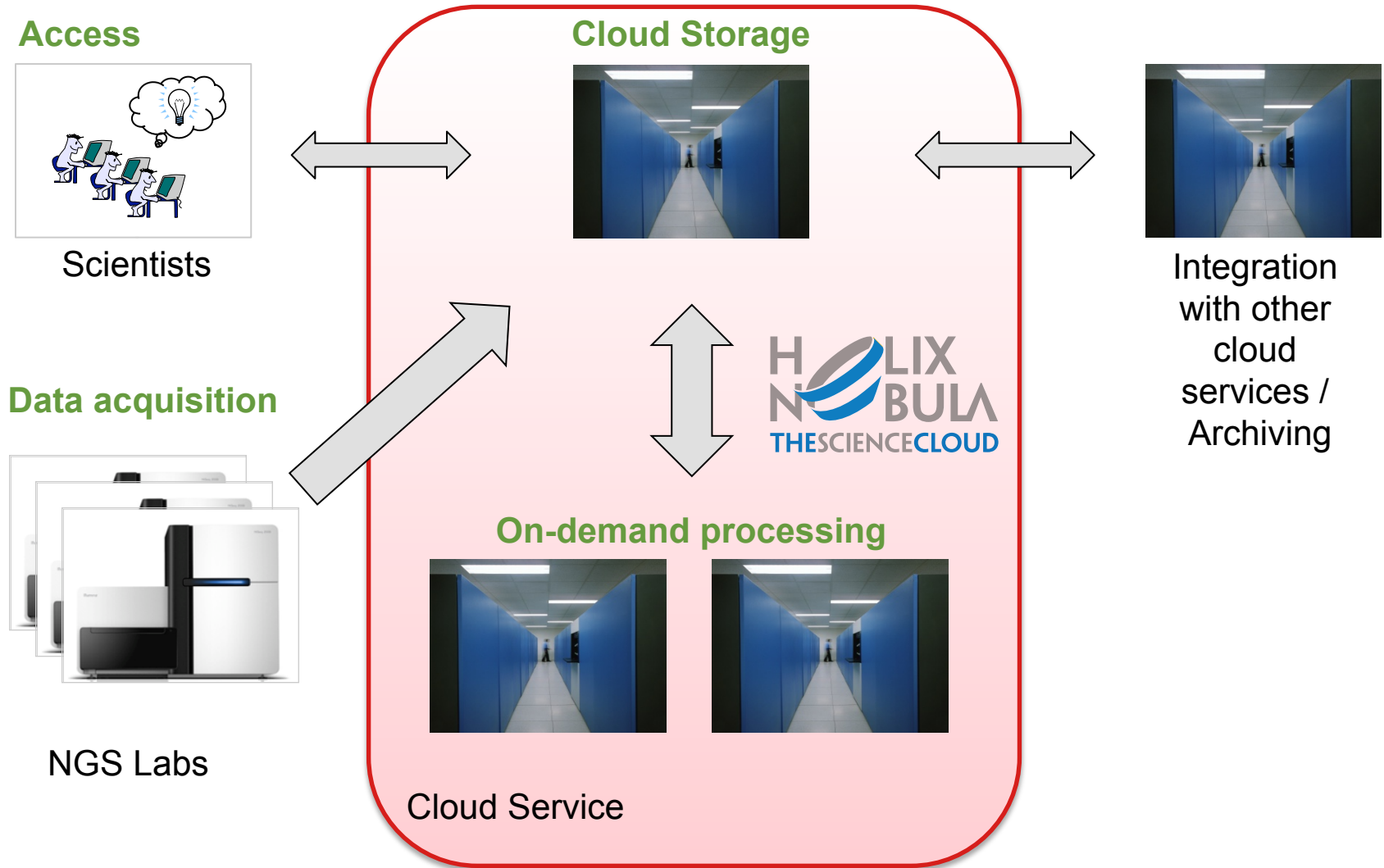
NGS - The Big Picture

- ~ 8.7 million species in the world (estimate)
- ~ 7 billion people
- Sequencers exist in both large centres & small research groups
- 200+ Illumina HiSeq sequencers in Europe alone
 - capacity to sequence 1600 human genomes / month
- Largest centre: Beijing Genomics Institute (BGI)
 - 167 sequencers, 130 HiSeq
 - 2,000 human genomes / day
- 500-1000 HiSeq devices worldwide today
 - 3-6 PB /day
 - 1.1 – 2.2 ExaBbytes / year

World Map of High-throughput Sequencers



EMBL Flagship project: Whole-Genome Assembly



A European cloud computing partnership: big science teams up with big business



Strategic Plan

- ▶ Establish multi-tenant, multi-provider cloud infrastructure
- ▶ Identify and adopt policies for trust, security and privacy
- ▶ Create governance structure
- ▶ Define funding schemes



To support the computing capacity needs for the ATLAS experiment

EMBL



Setting up a new service to simplify analysis of large genomes, for a deeper insight into evolution and biodiversity



To create an Earth Observation platform, focusing on earthquake and volcano research

Atos

Capgemini
CONSULTING. TECHNOLOGY. OUTSOURCING

CloudSigma

esi

interoute
from the ground to the cloud

logica
be brilliant together

OpenNebula.org
The Open Source Toolkit for Cloud Computing

orange Business Services

SAP

the SERVER LABS
the IT architects

sixsq

Telefonica

terradue 2.0

THALES

Systems



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