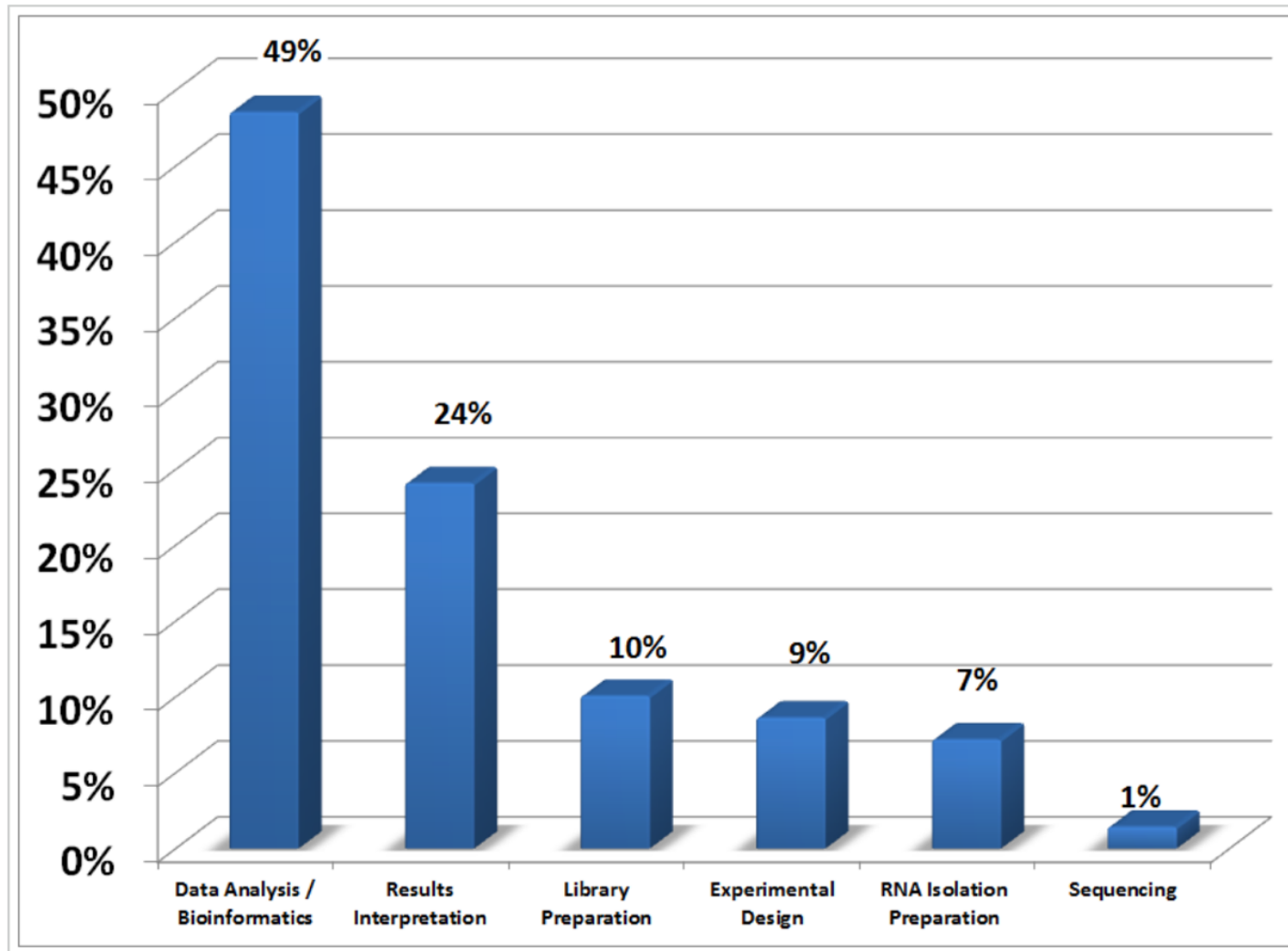




WE TRANSFORM
DATA INTO KNOWLEDGE



We asked: which step of an RNA-seq project do you find most intimidating?



Total Respondents = 287

Rnaseq blog
questionnaire 2017

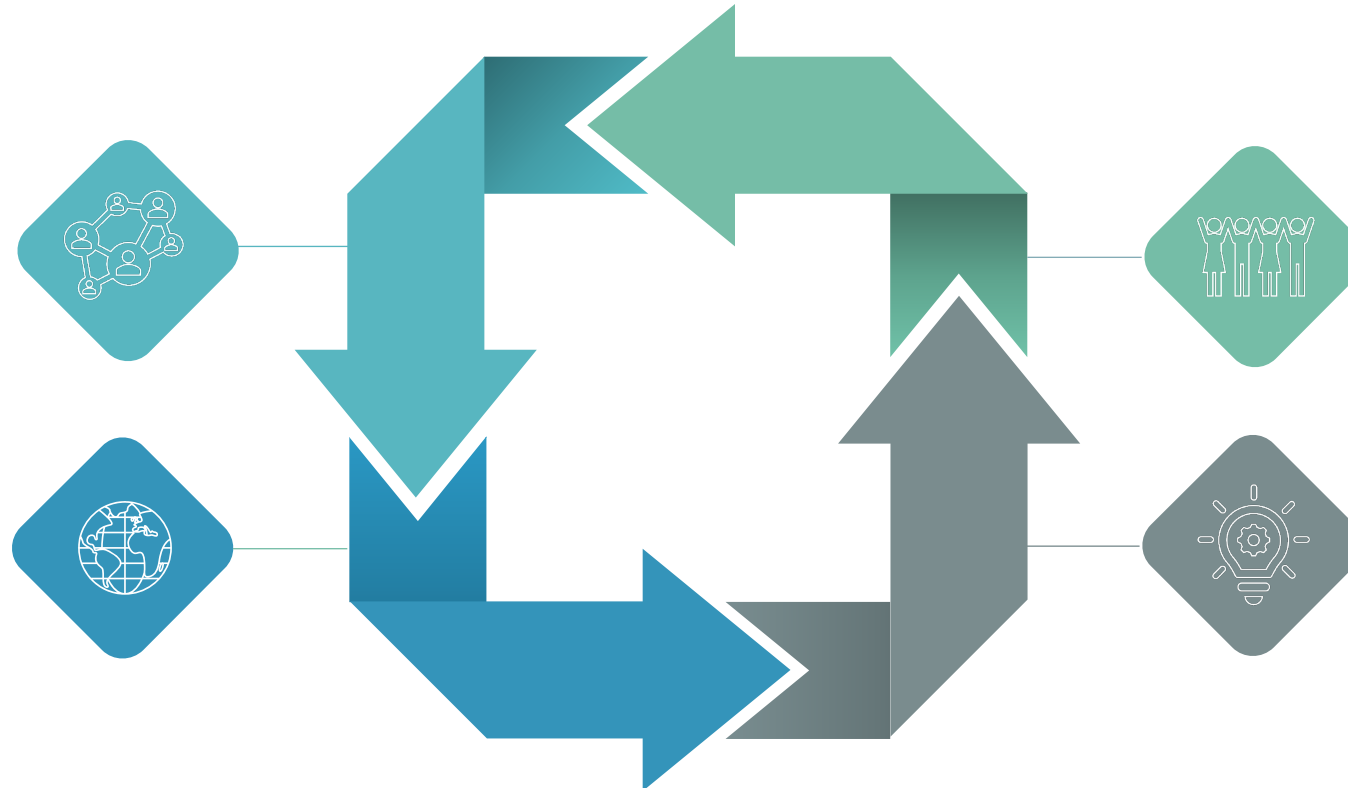
We make *omics* data more accessible for everyone

Our vision

Bring innovation and disruptive technology in the omics and biotechnology fields helping our customers to accelerate their research outcomes

Our mission

Make everyone benefiting from the extraordinary potential that omics has in changing our lives

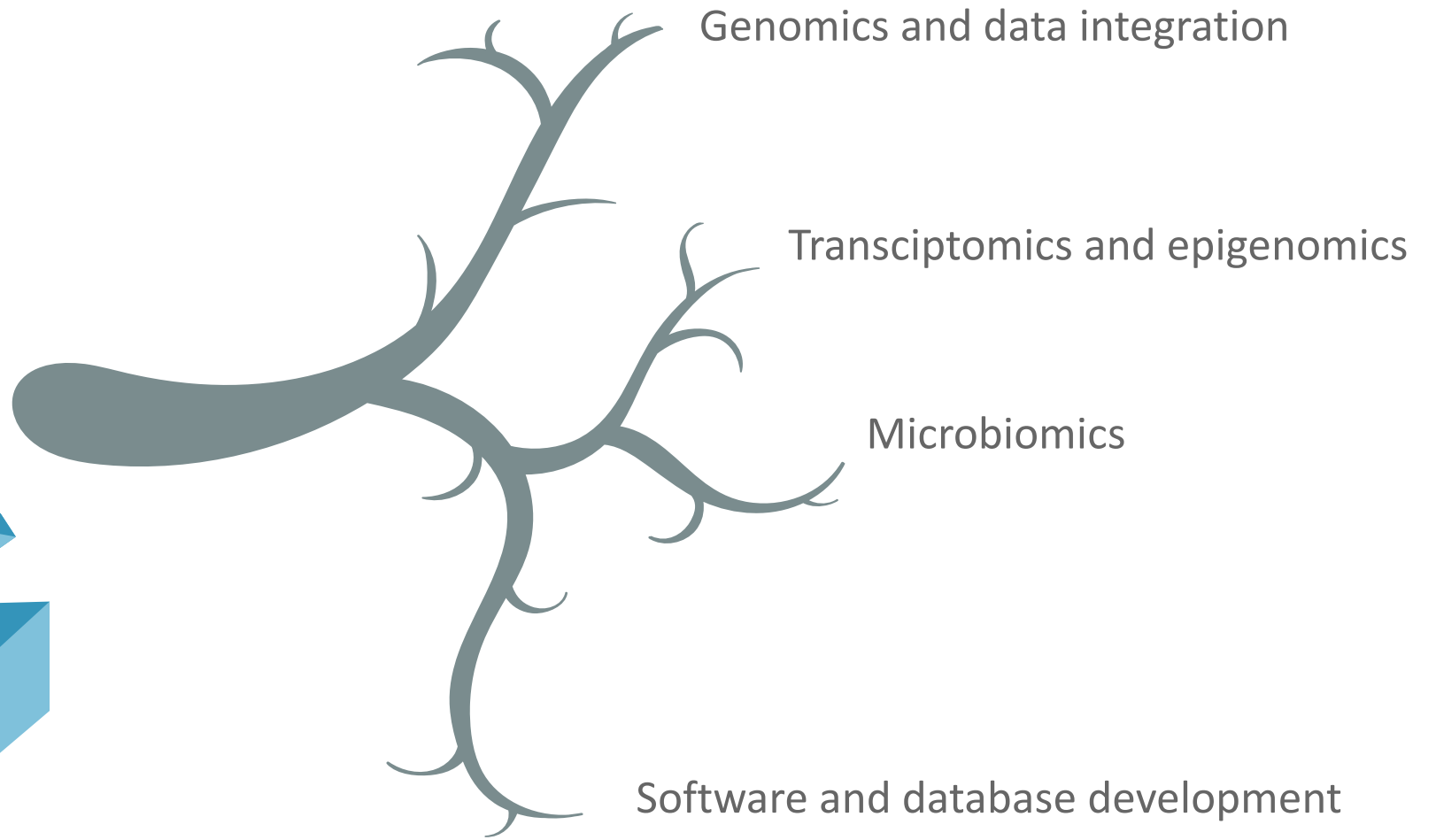


We are

Scientists, we understand the dilemmas our customers face.
Innovative, we have strong commitment and investment in cutting-edge technologies

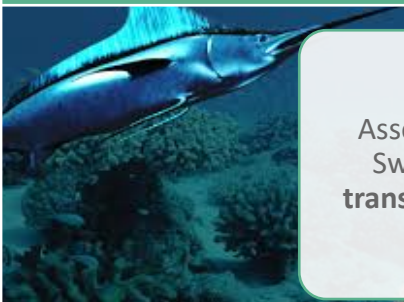
We offer

End-to-end solutions and consulting to customers working in basic, applied and industrial sciences



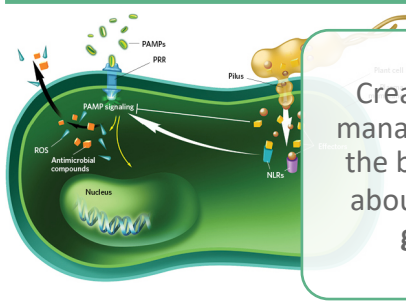
Consulting projects in agrogenomics

De novo transcriptome



Assembly of
Swordfish
transcriptome

Data Mining



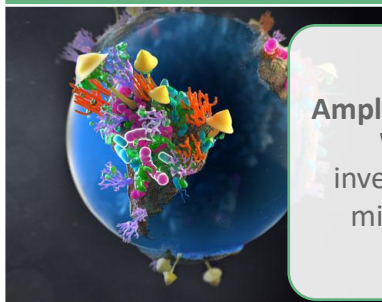
Creation and
management of
the biggest DB
about **Plant R
genes**

Transcriptomics



Create a **Single
cell RNAseq**
pipeline

Metagenomics



**Amplicon seq and
WGS** to
investigate soil
microbiome

Metagenomics



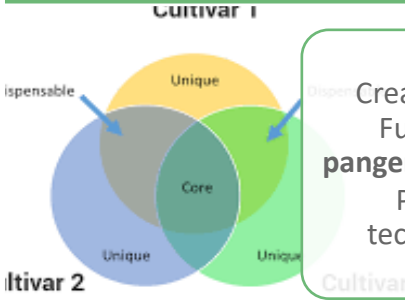
Detect **food
contamination**
using NGS data

Epigenomics



**Omics data
integration** to
study tomato
senescence
process

Genomics



Creation of a
**Fusarium
pangenome** using
PacBio
technology

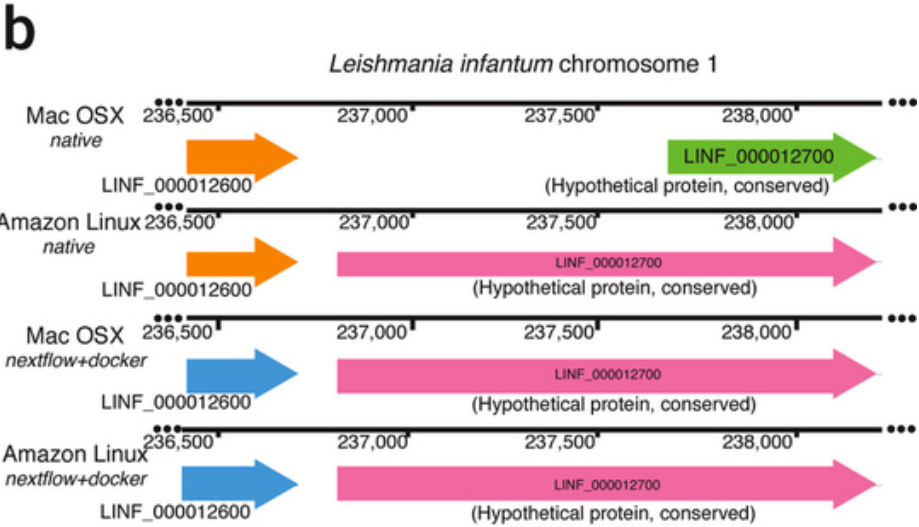
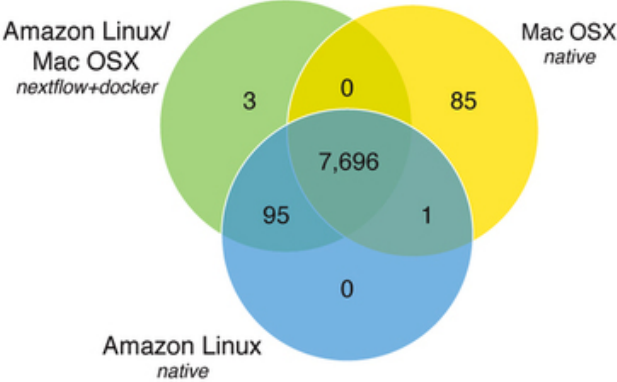
Genomics



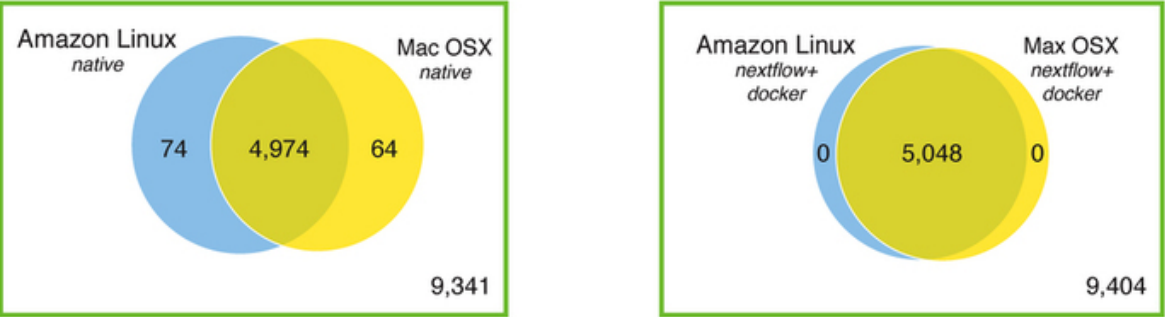
**De novo
sequencing** of a
cold resistant
potato

Hundreds of bioinformaticians performing analyses on different platforms and with different methods

a Gene annotation of *Leishmania infantum* with Companion

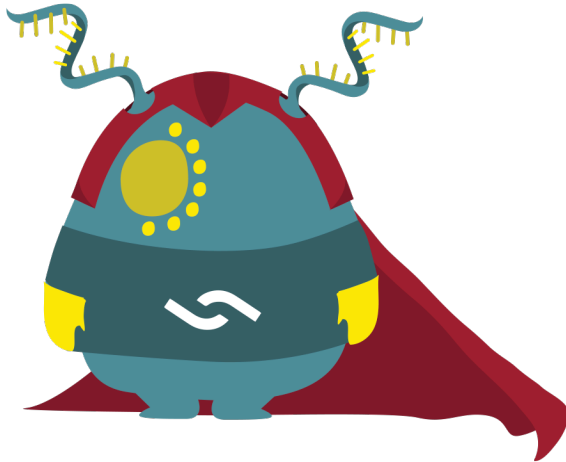


c Transcript quantification and differential expression with Kallisto and Sleuth



Nature Biotechnology 35, 316–
319 (2017) doi:10.1038/nbt.3820

AIR – Friendly platform for frightened researchers



01

FAST - Automated analysis, same day results

02

VALID - For any sequenced species

03

UPDATED - Using the latest algorithms

04

Scientifically robust and reproducible

05

NGS Platform agnostic

ion torrent
△ ★ ○ × □ + ≈

pb PACBIO®

illumina®

Oxford
NANOPORE
Technologies

SEQUENTIA

Technologies behind it

IT

Docker Technology

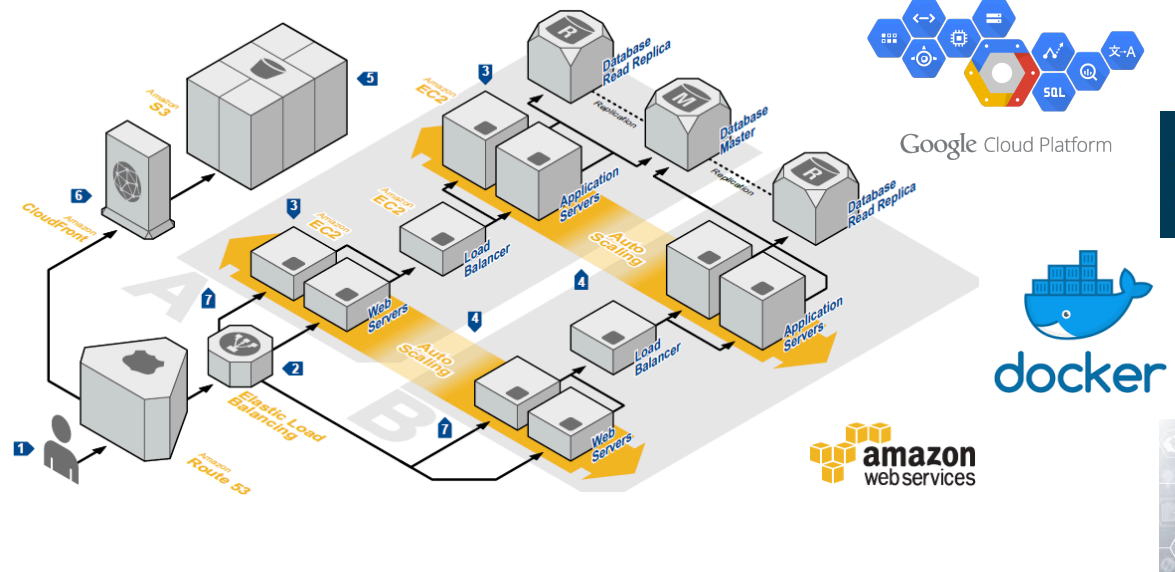
Cloud and hybrid solution

Fault tolerant

Unlimited computational resources

Proprietary flow infrastructure

Informatics layer: Seaflow our proprietary flow system



THANK YOU!

www.sequentiabiotech.com

