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SHOP  
GARR  
2022

NET  
MAKERS

Human Technopole  
from startup phase to  
a large-scale research  
infrastructure

Albino Zamboni

Human Technopole



# HT Foundation

- The Human Technopole Foundation was established by financial law n. 232, 11 December 2016. The founding members of the Foundation are the **Ministry of Economy and Finance, the Ministry of Health and the Ministry of Education, University and Research** which are responsible for supervising the Foundation.
- The purpose of the Foundation, as indicated in art.1, c. 116 of the above mentioned law, is the **creation of a multidisciplinary scientific and research infrastructure of national interest**, integrated in the fields of health, genomics, nutrition, data and decision science and in the implementation of the Human Technopole scientific and research project (“HT Project”).



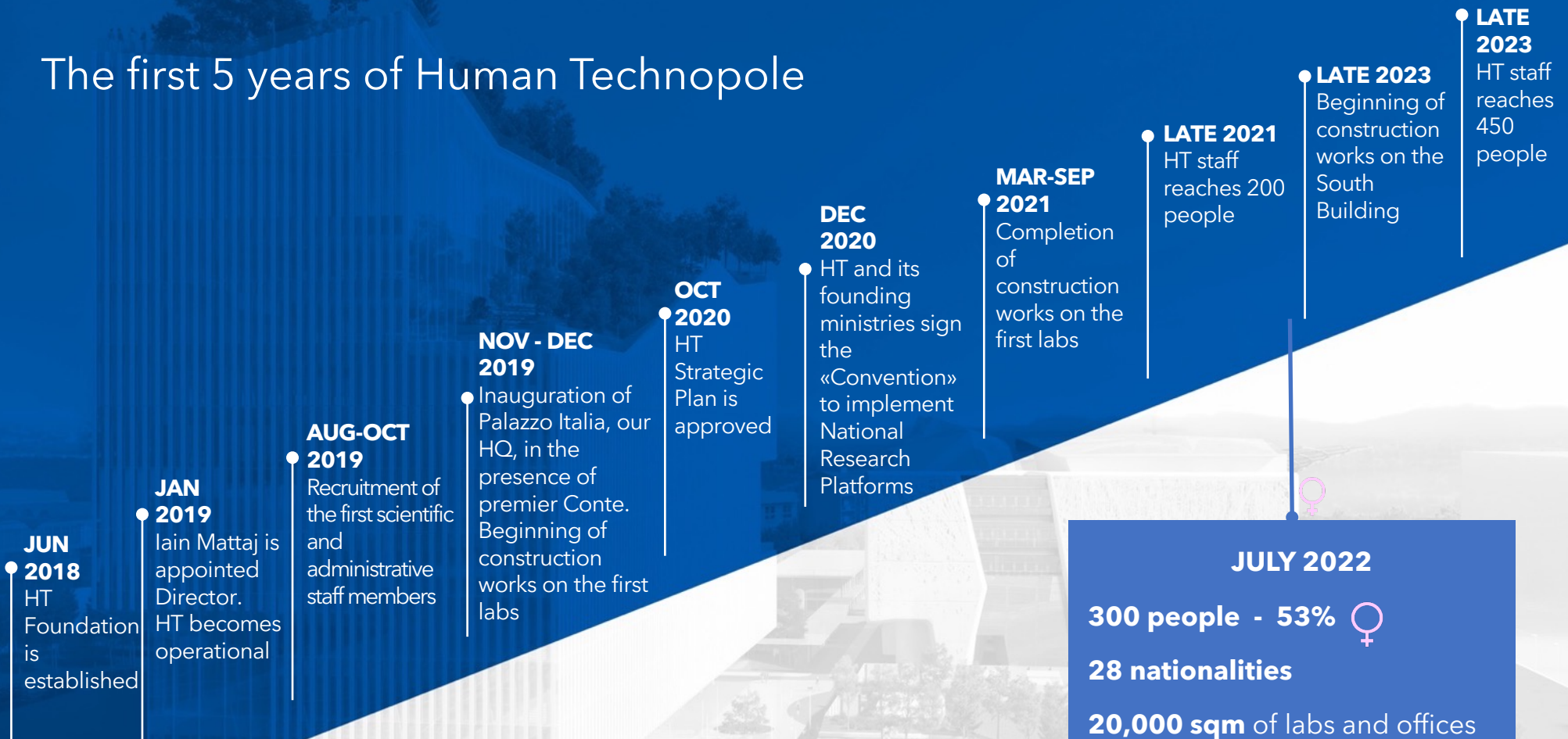
# Our Mission

- Improve human health and well-being, including healthy ageing.
- Carry out frontier research to improve people's health and well-being.
- Set up and operate a large-scale research infrastructure with interdisciplinary laboratories for the development of precision medicine.
- Act as an open hub to support the growth of the Italian life science research community.
- Engage in industrial cooperation and technology transfer support activities.
- Employ 1,000 scientists including biologists, bioinformatics, chemists, engineers, mathematicians and computer scientists.



# The first 5 Years of HT

## The first 5 years of Human Technopole



# HT Today

- **63%** of the scientific team from **international institutions**.
- **70 Italians back from abroad**.
- **Fast scientific recruitment** (1 scientist per week for the next 3 years).
- **8.5-million-euro grant** (Jan. '21- Mar. '22).
- **20,000 sqm** of labs and offices.

*Cryo-Electron Microscopy  
Light Imaging & Image Analysis  
Genomics  
Data Centre*



# Our Research Centers

## OUR LINES OF RESEARCH

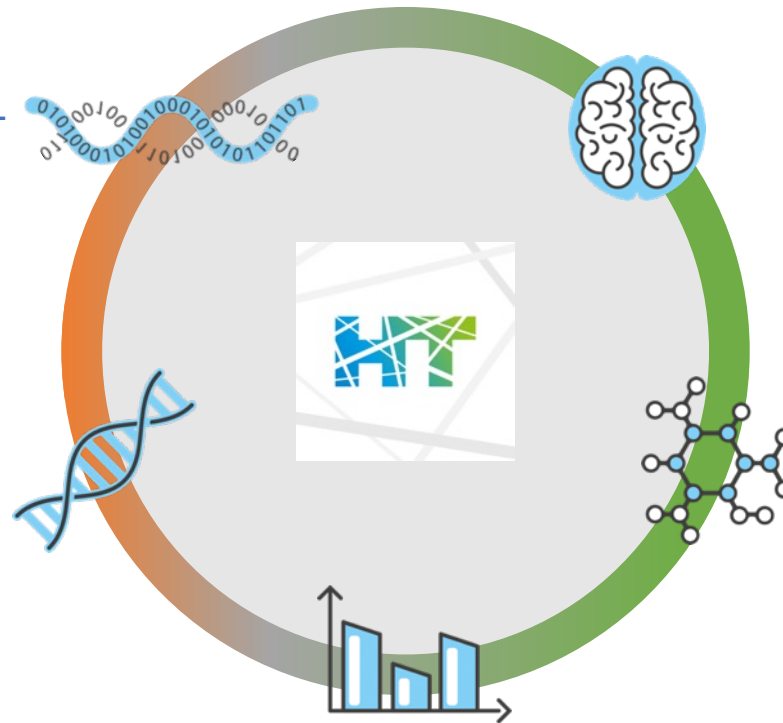
**GENOMICS**  
FUNCTIONAL  
POPOULATION & MEDICAL

**NEUROGENOMICS**

**STRUCTURAL  
BIOLOGY**

**COMPUTATIONAL  
BIOLOGY**

**HEALTH DATA SCIENCE**



# Our Research Centers

Our biomedical research aims at developing predictive and personalised medicine to treat cancer, cardiovascular and neurodegenerative diseases.



## Genomics

The Centre studies genomics characteristics and traits to identify how heritable genetic information is shared in view of identifying more personalised treatments



## Neurogenomics

The Centre studies neuropsychiatric and neurological diseases to probe the structure, function and development of the nervous system.



## Structural biology

The Centre aims at gaining precise knowledge of the structure of macromolecules, a fundamental step in understanding the function of cells.



# Our Research Centers

Our biomedical research aims at developing predictive and personalised medicine to treat cancer, cardiovascular and neurodegenerative diseases.



## Computational biology

The Centre develops solutions for the analysis, management and integration of data produced by other Centres, making it available to the wider scientific community.

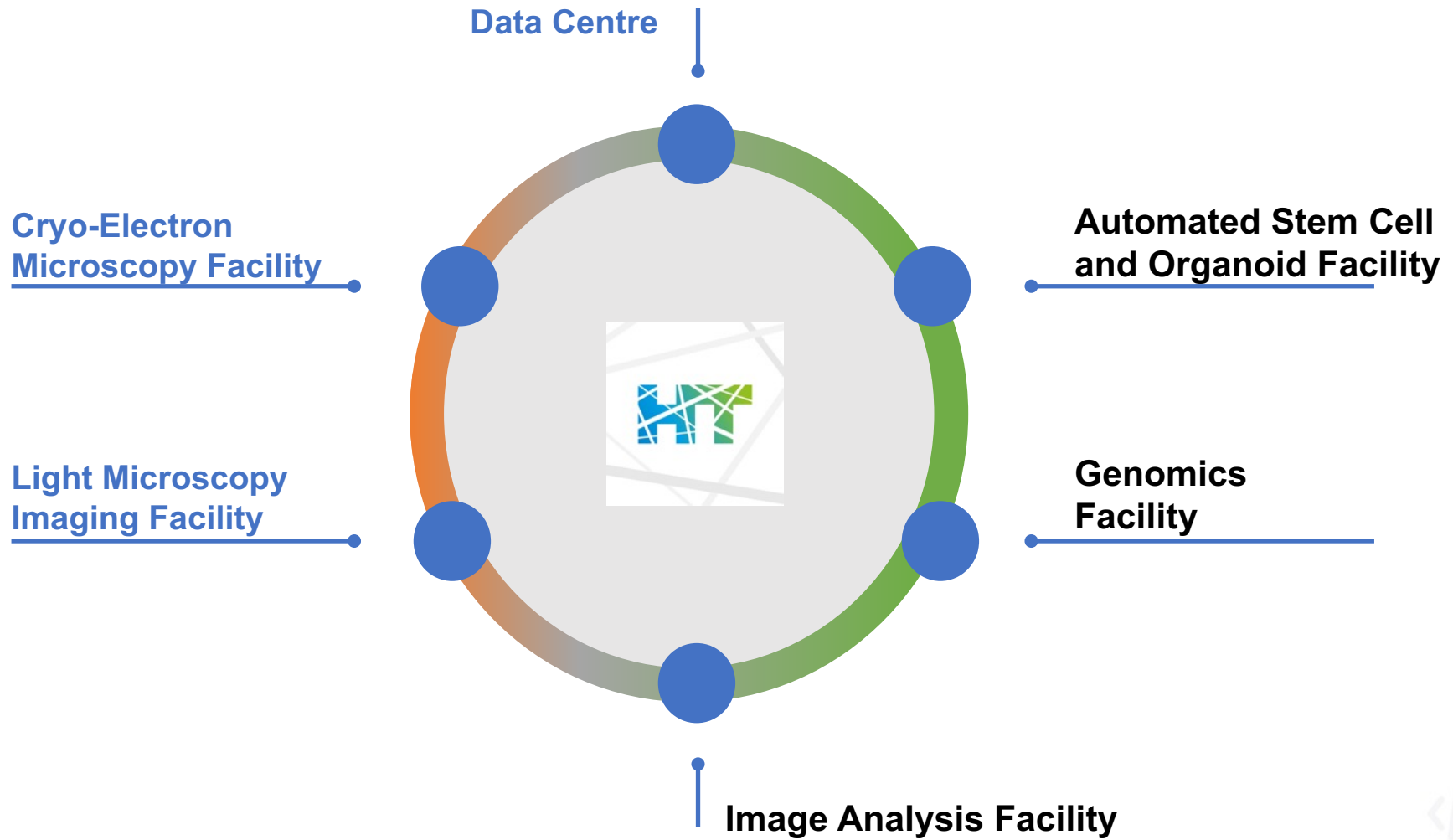


## Health Data Science

The Centre analyses clinical and socio-economic data to provide advice to different stakeholders, in particular policymakers, mainly to the national health system.



# Our Facilities

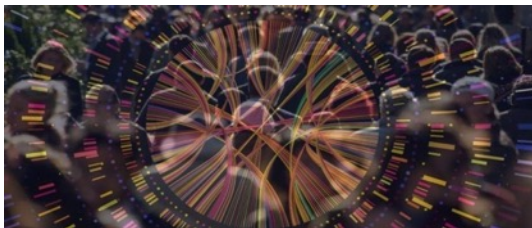


# Our Facilities

HT is a national hub and centre of reference for life science research.

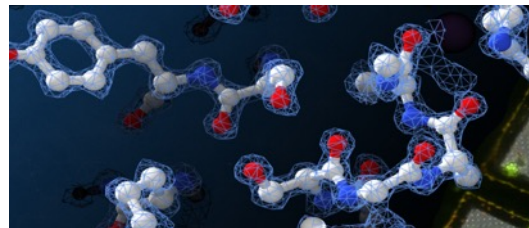
Our facilities are available to HT scientists and researchers as well as to the external scientific community who will access them through open selection procedures based on merit.

## Six research facilities:



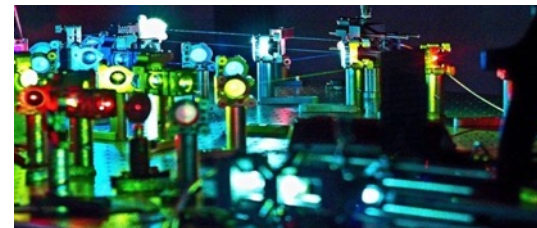
### Genomics

Large-scale DNA/RNA sequencing infrastructure to conduct population studies and support national screening initiatives.



### Cryo-Electron Microscopy

Italy's most comprehensive CryoEm infrastructure: five state of the art microscopes to freeze molecules and observe them at atomic level.



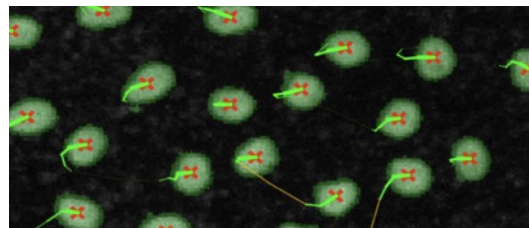
### Light Imaging

With a focus on 3D imaging it will photograph rare, dynamic and constantly evolving processes.



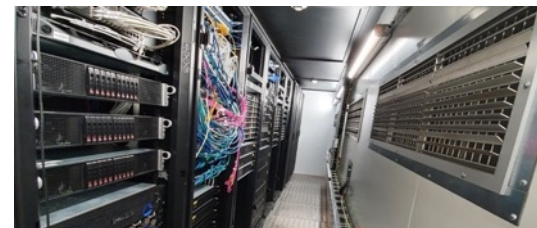
### Automated Stem Cell and Organoid Facility

It will engage in cell re-programming, genome editing and organoid culture.



### Image Analysis

Solutions for for image restoration, real-time image analysis, big data management and visualization.

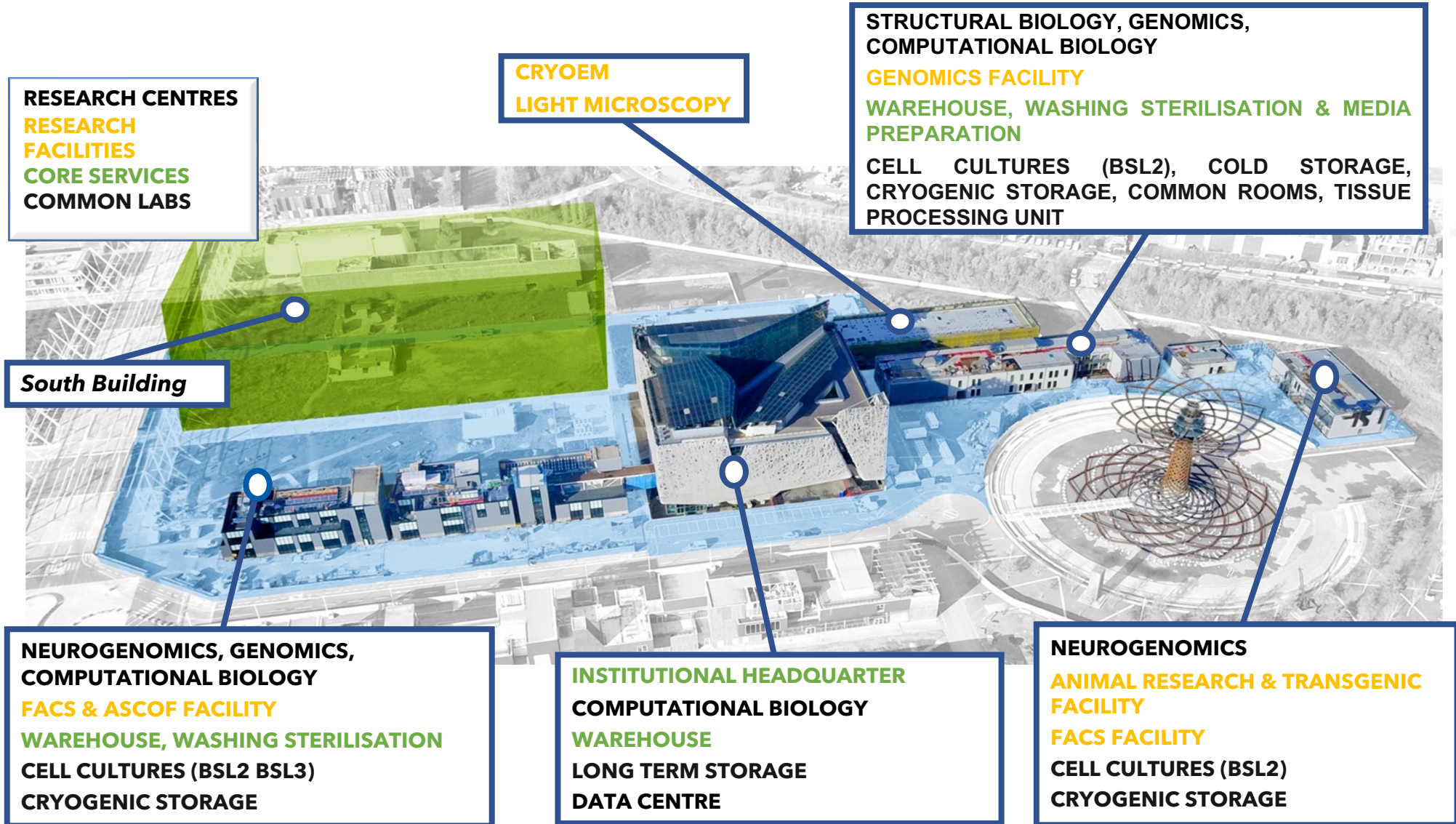


### Data Centre

High storage and computing capacity to support researchers in the storage and analysis of huge amounts of data.

# Our Campus

Ongoing works to build a large scale research infrastructure





# South Building

**Once completed, it will host labs for 800 scientists as well as offices, event spaces, workshops, and training courses.**

The winning project has been awarded in April 2020, and the building, which is to be constructed, will be ready in 2027.





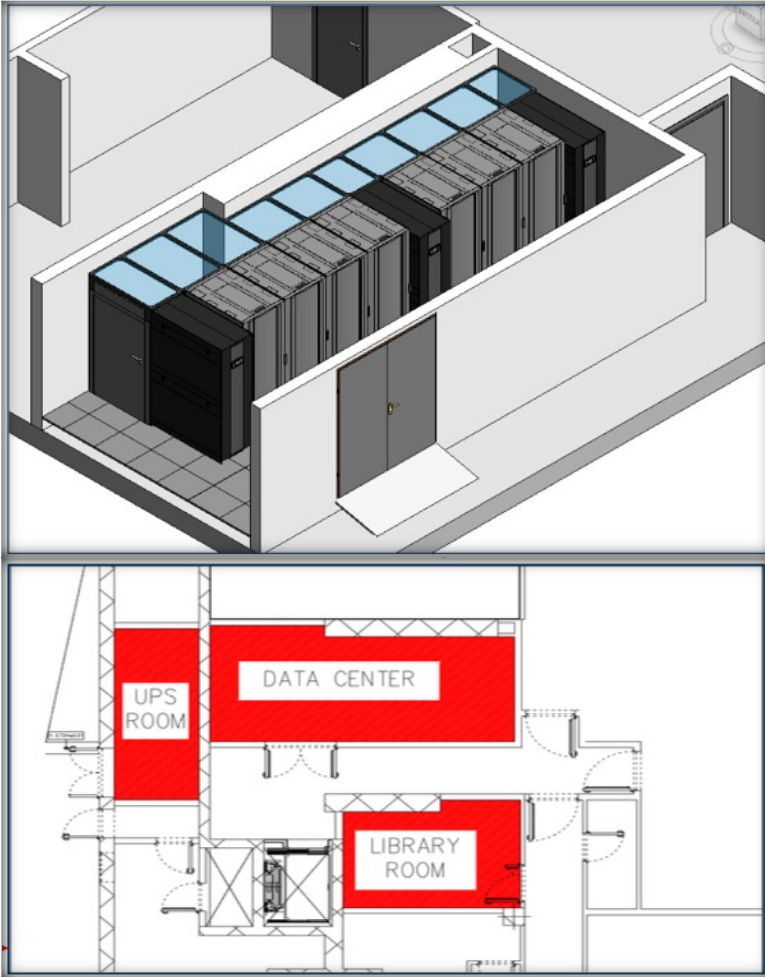
# Our Campus

## IT Data Center Container



# Our Campus

Extending resources of External shelter





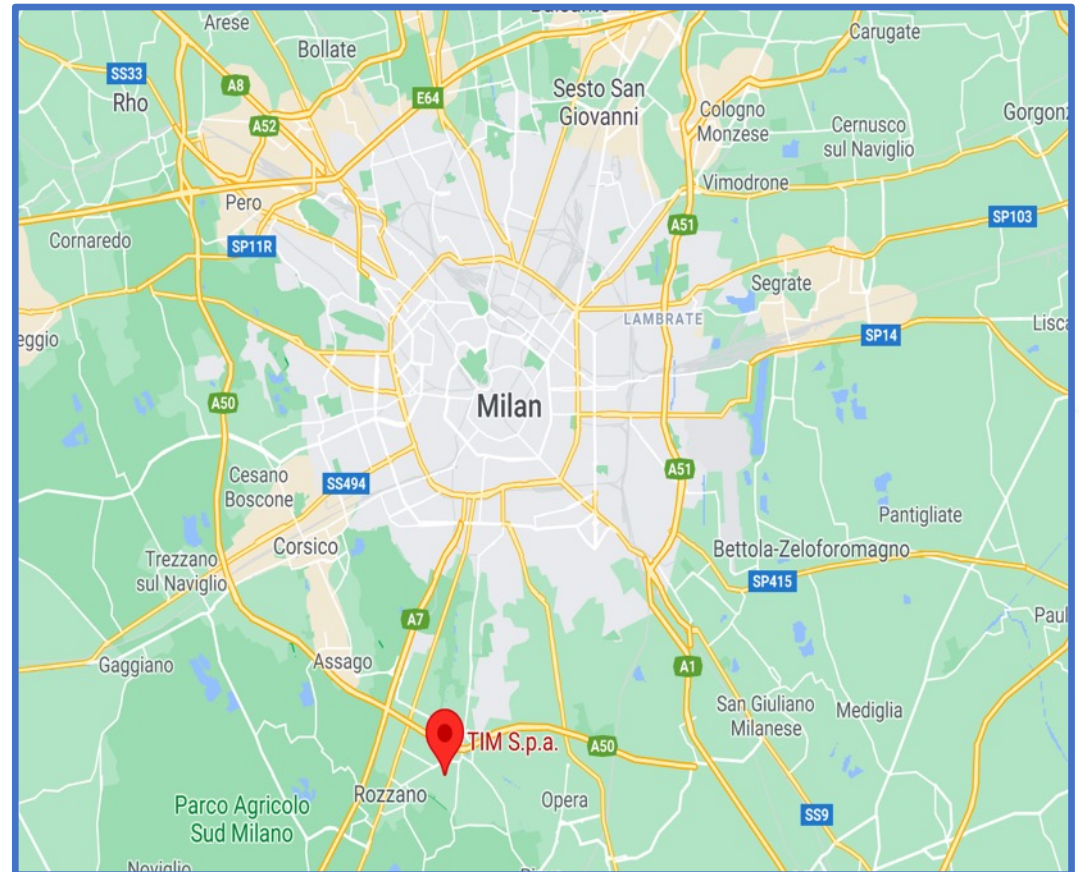
# Our Colocation

## Offsite backup and data processing

### Redundant services:

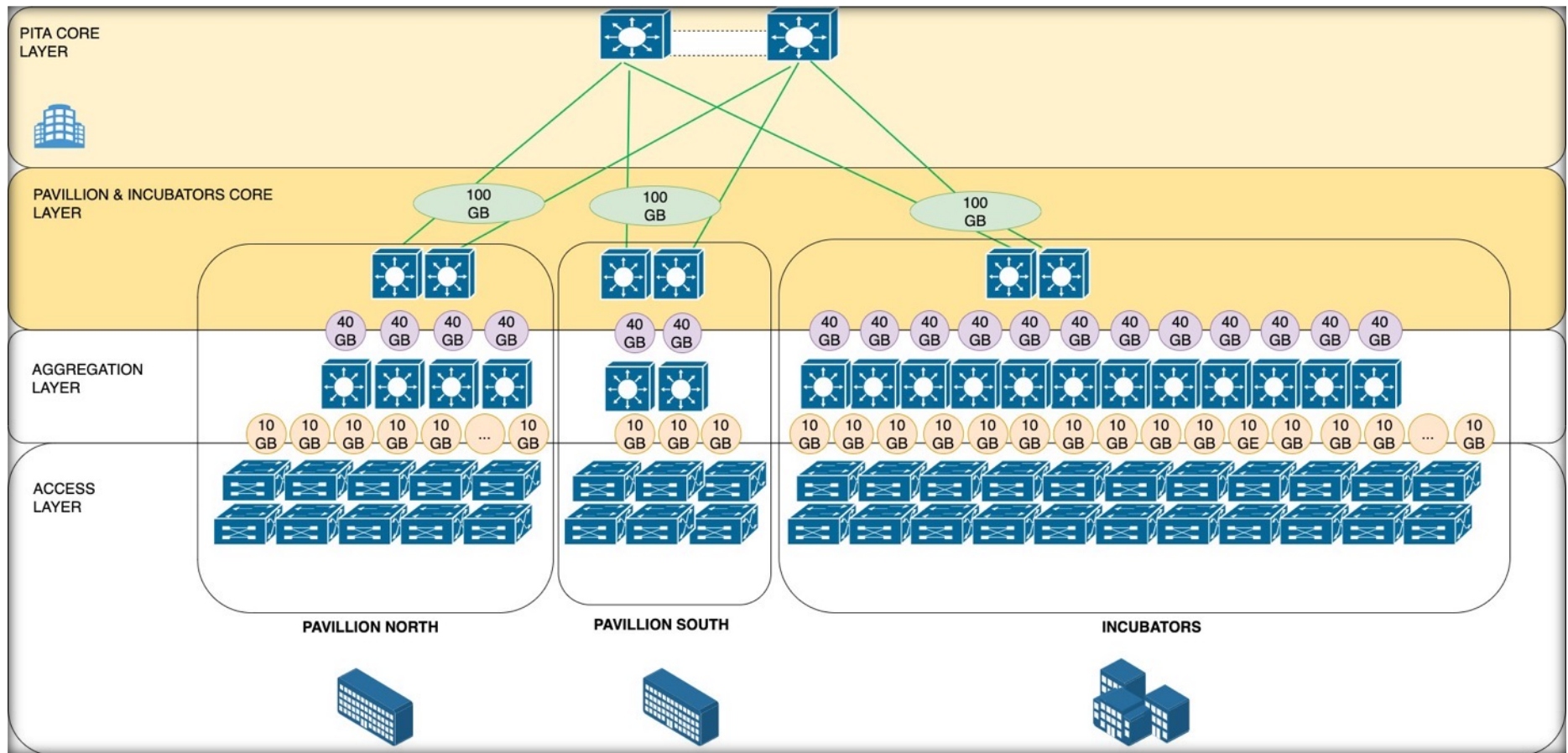
- **HPC**
  - 12 compute nodes
  - 1 PB attached storage
- **Virtualisation**
  - 3 nodes
  - 30 TB Fiber Channel storage
- **Storage:**
  - 1PB central scientific storage
- **Connection:**
  - 10Gbps DWDM redundant

### Low latency network connection



# Our Campus LAN connections

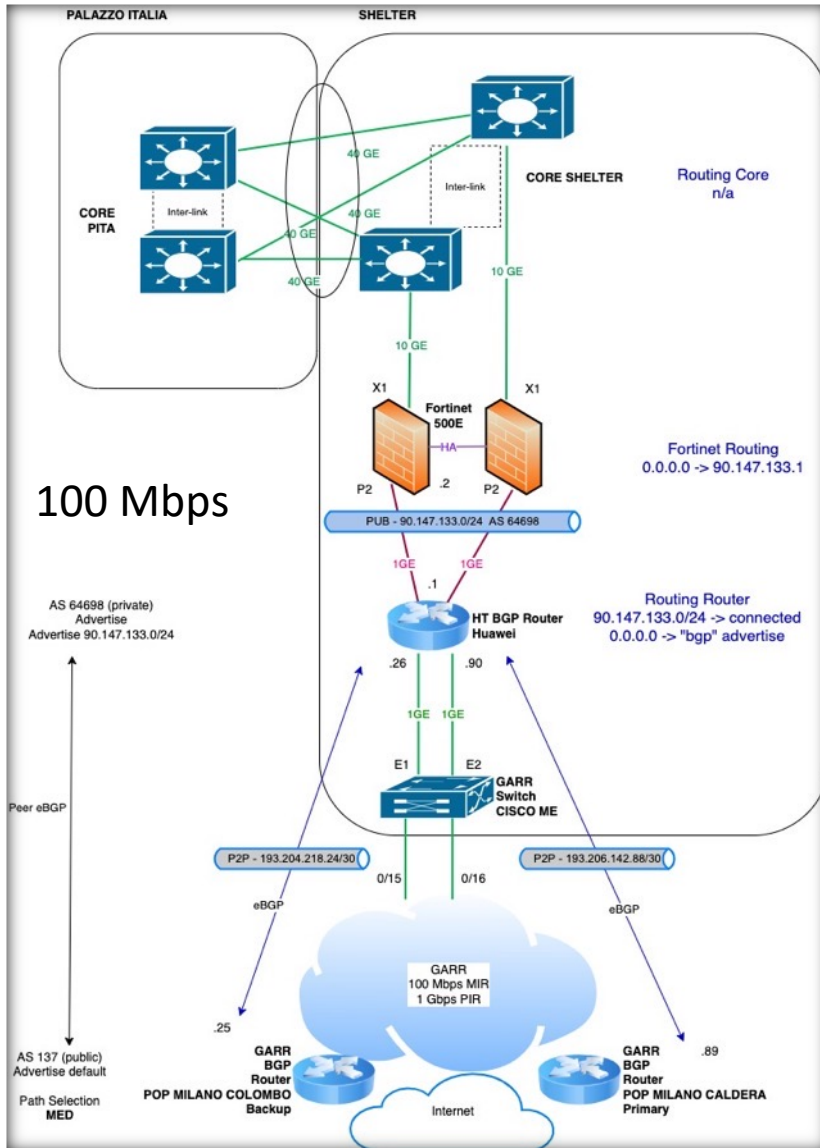
x 100Gbps inter-buildings connections



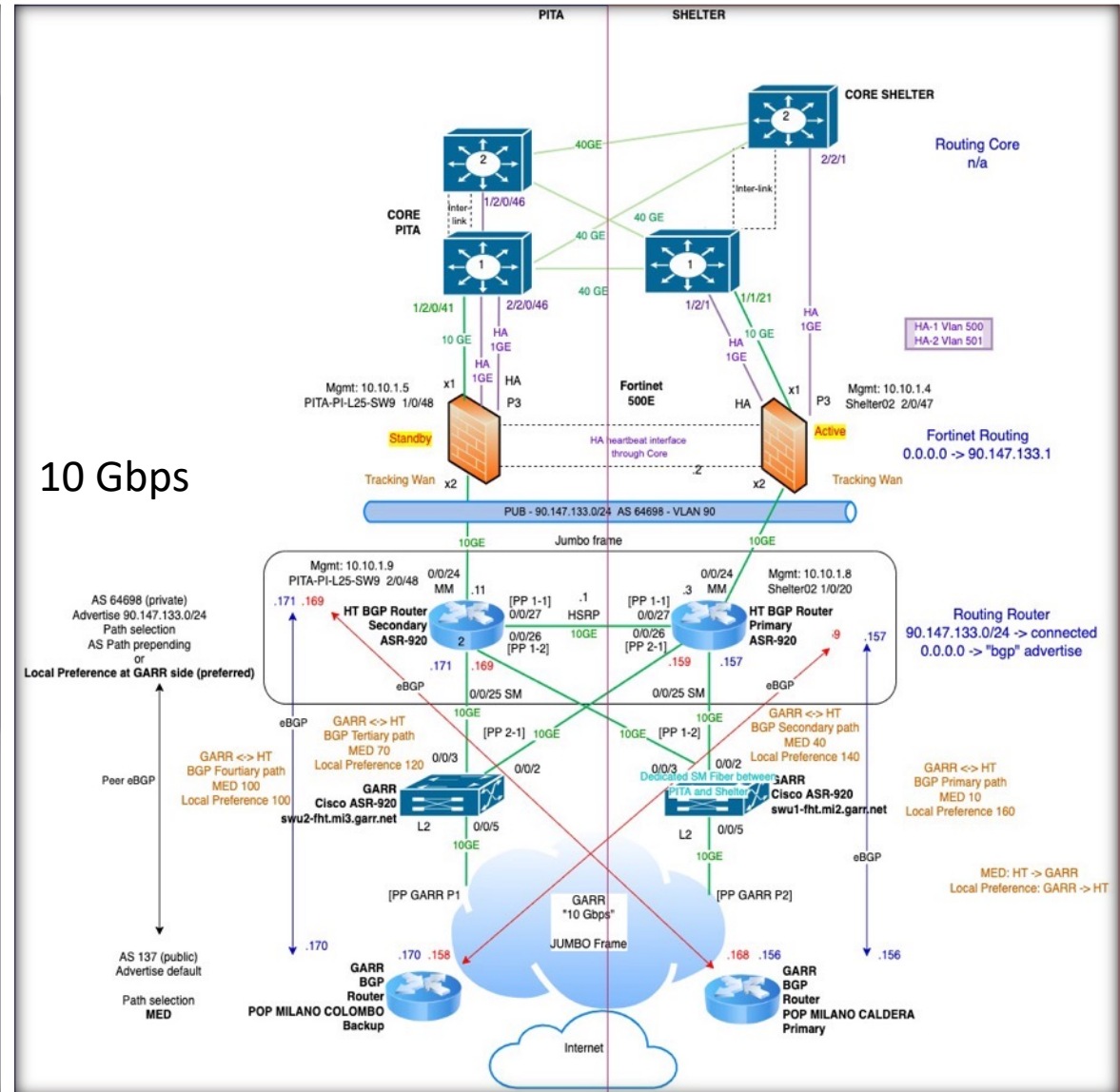


# Our Campus WAN connections

2021



2022





# High Performance Computing

## DELL HPC

**Usage:** Computational processing

**Operating System:** CentOS 8

**Scheduler:** SLURM

**Cluster Interconnect :** 100 Gbps

	25 CPU Nodes	10 GPU Nodes	5 FAT Nodes
CPU GHz	2.9	2.2	2.2
N° Core	32	36	36
RAM	576 GB	576 GB	1.1 TB
GPU		4 x V100	4 x V100
Total CPU Core Count			1340
Total RAM			26 TB
Calculated performance			<b>150 TFLOPS</b>

### Available Queues:

- **CPU Standard** - CPU nodes (CPU Intensive)
- **GPU Standard** - GPU nodes (GPU intensive)
- **High Intensive** - Specially allocated nodes (Memory or GPU Intensive)

**Access:** 2 x Login Nodes

# Storage

HPC access

**High Performance storage**

**Usage:** High Performance IO Operations

**Filesystem:** Parallel BeeGFS with HA

**Capacity:** 2.1 PB

**Throughput:** 18 GB/s with sequential R/W

**Cluster interconnect:** 100 Gbps Infiniband HDR



**BeeGFS®**



# Virtualisation

## Traditional Virtualisation + High Performance GPU-Driven VDI

Core Services	
Physical Servers	5
Cores	560
Memory (GB)	3840
Storage (TB)	30

VDI - Double Precision GPUs - Intel Based	
Physical Servers	5
Cores	320
GPUs (NVIDIA V100s)	20
Memory (GB)	3840
Storage (TB)	30

VDI - Single Precision GPUs - AMD Based	
Physical Servers	2
Processors	64
GPUs (NVIDIA A40)	12
Memory (GB)	4096
Storage (TB)	30

Q1 2023

VDI - Single Precision GPUs - AMD Based	
Physical Servers	12
Processors	384
GPUs (NVIDIA A40)	72
Memory (GB)	24576
Storage (TB)	180

## Mellanox low latency switches

- 300nsec for 100GbE port-to-port

- Flat latency across L2 and L3 forwarding

# Storage

## Central Scientific Storage

**Usage:** Tier 1 Storage for Group Shares and User Homes

**Protocols:** NFSv4, SMB 3.1, and S3

**Group Share (Default) Size:** 10TB

**Home Share (Default) Size:** 200GB

**Backup Policy:**

- Hourly Snapshots – 1 Day Retention
- Daily Snapshots – 1 Week Retention
- Weekly Snapshots – 1 Month Retention



**RAW** Central Storage Total Capacity

# 9PB

Available on Campus

## 7PB approx

8/52 nodes moved to colocation

## > 2PB



# Backup

Second line of defence for recovery

**Backup System:** Bacula Enterprise

**Capacity:** 2.5 PB

**Technology:** Disk Backup

**Retention:**

- **Group Shares:** 1 year
- **VMware VMs:** 3 months
- **Laptops:** 1 month
- **Office 365:** 1 month



**Upgrade Plans (2022 - 2023):** Long Term Archive (10 years) on Tape

**Initial deployment plan capacity:** 60 PB based on LTO8

**Scale as needed strategies:**

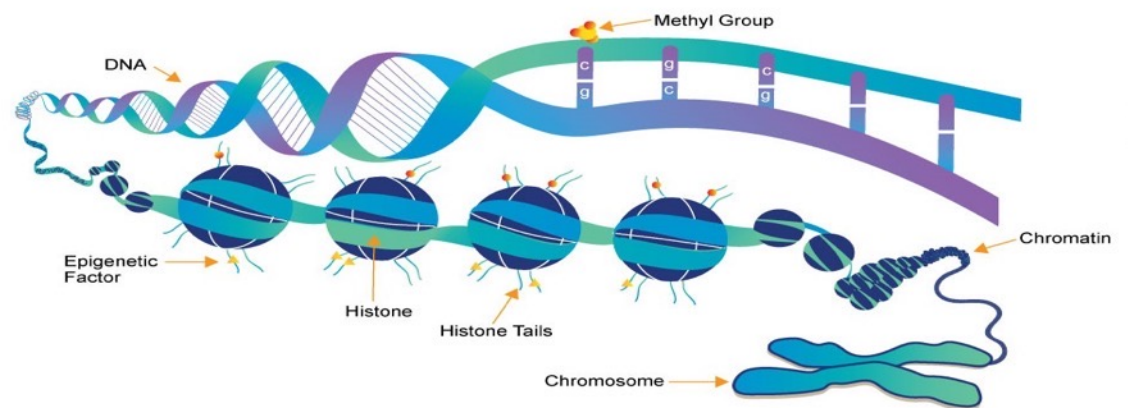
- New Expansion shelf
- Tape Generation



# Genomic-Sequencers

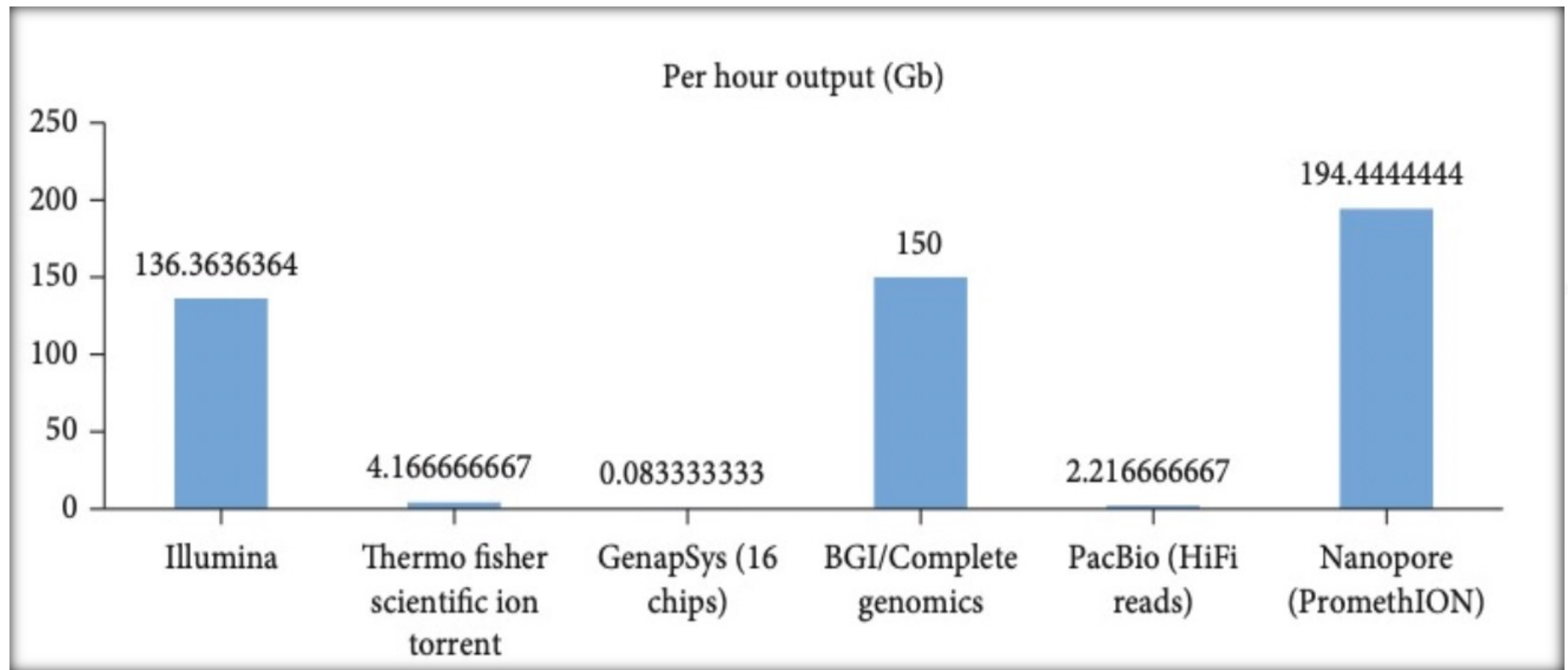
New sequencers generation:

- High Throughput - Gbps
- Long run experiment - hours
- High cost run – tens of k €
- Cloud AI monitoring



# Genomic-Sequencers

Sequencers can produce up to 200 Gbyte data/h



# Genomics-Sequencers

Experiments can take up to 72 run hours and produce 14 Tbyte data

Manufacturer	Read length	Data output	Max. run time (hours)	Chemistry	Key applications**
Illumina (NovaSeq 6000)	300 PE	6 Tb (6000 Gb)	44	Sequencing by synthesis	SS-WGS and TGS, TGEP, 16sMGS, WES, SCP, LS-WGS, CA, MS, MGP, CFS, LBA
Thermo Fisher Scientific Ion Torrent (Ion GeneStudio S5 Prime)	600 SE	50 Gb	12	Sequencing by synthesis	WGS, WES, TGS
GenapSys (16 chips)	150 SE	2 Gb	24	Sequencing by synthesis	TS, SS-WGS, GEV, 16S rRNA sequencing, sRNA sequencing, TSCAS
QIAGEN (GeneReader)	100 SE	Not available	Not available	Sequencing by synthesis	Cancer research and identifying mutations
BGI/Complete Genomics	400 SE	6 Tb (6000 Gb)	40	DNA nanoball	Small and large WGS, WES and TGS
PacBio (HiFi Reads)	25 Kb	66.5 Gb	30	Real-time sequencing	DN sequencing, FT, identifying ASI, mutations, and FPM
Nanopore (PromethION)	4 Mb	14 Tb (14000 Gb)	72	Real-time sequencing	SV, GS, phasing, DNA and RNA base modifications, FT, and isoform detection



# Genomics-Sequencers

A single sequencer can produce up to 676 Tbyte data/year

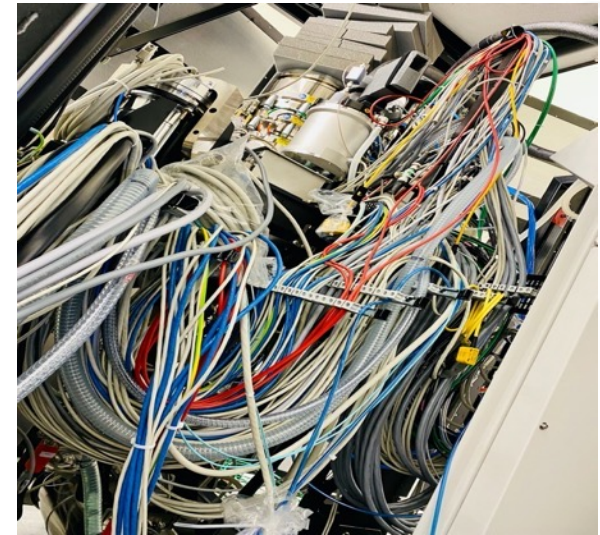
Instrument	WEEK	MONTH	YEAR
NovaSeq 1	14 Tb	56 Tb	672 Tb
NovaSeq 2	14 Tb	56 Tb	672 Tb
MiSeq	30 Gb	120 Gb	1.4 Tb
NextSeq	840 Gb	3.3 Tb	40.3 Tb
PromethION	1.1 Tb	4.5 Tb	56 Tb



# Cryo Electron Microscopes

To get high-resolution images we need to freeze and keep the sample at  $-175\text{ }^{\circ}\text{C}$

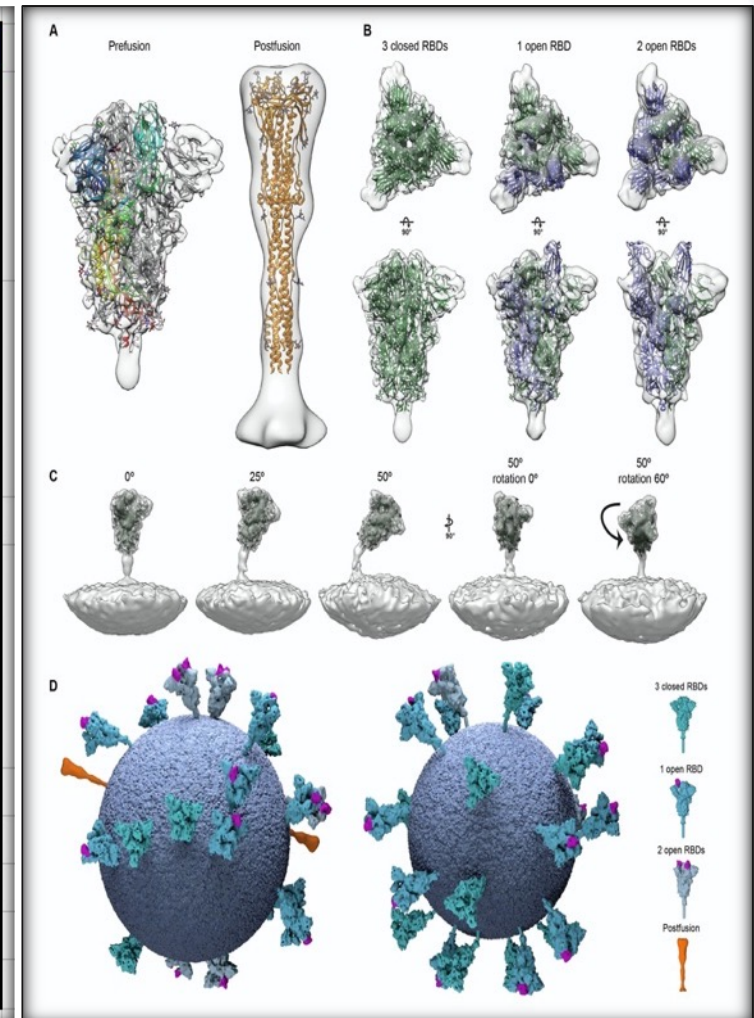
- **Cryo-electron microscopy** allows to photograph the inner section of a frozen cell and individual molecules at high resolution
- We **freeze** the sample (cells, enzymes, DNA, viruses, etc.) to preserve them and get better images
- It's a completely new level of **observation**, this is why the technique was awarded the 2017 Nobel Prize for Chemistry



# Cryo Electron Microscopes

Microscopes can produce up to 15 Gbyte image data/minute

Instrument	Maximum Throughput (MB/s)
Krios	Min: 1 image (500MB to 1.5GB) per minute. Max: 10 image (500MB to 1.5GB) per minute. NO intermediate files. Usual dataset size: 1-4 TB.
Glacios	Min: 1 image (500MB to 1.5GB) per minute. Max: 10 image (500MB to 1.5GB) per minute. NO intermediate files. Usual dataset size: 1-4 TB.
Spectra	Max: 1 image per second (64 MB/s)
Tundra	Min: 1 image (500MB to 1.5GB) per minute. Max: 10 image (500MB to 1.5GB) per minute. NO intermediate files. Usual dataset size: 1-4 TB.
Talos	Max: 1 image per second (64 MB/s)
Leica Stellaris	Max: 1 image per second (32 MB/s)
Leica Thunder	Max: 1 image per second (32 MB/s)
Aquilos 2	Max: 1 image per second (32 MB/s)
Arctis	Max: 1 image per second (64 MB/s)





# Cryo Electron Microscopes

A single microscope can produce up to 2 PB image data/year

Instrument	TB/year
Krios	770
Glacios	770
Krios	770
Spectra	1.925
Tundra	770
Talos	1.925
Leica Stellaris	962
Leica Thunder	962
Aquilos 2	1.925
Arctis	1.925

# Inter-Institutes connections

## Institutes/research centers:

- IEO  
[Istituto Europeo di Oncologia \(ieo.it\)](http://ieo.it)
- IRCCS (research hospitals in Italy)  
<https://www.hsantalucia.it/en/irccs>
- CRG  
[Centre for Genomic Regulation Website \(crg.eu\)](http://crg.eu)
- MRC  
[MRC Laboratory of Molecular Biology \(cam.ac.uk\)](http://cam.ac.uk)
- NCBI  
[National Center for Biotechnology Information \(nih.gov\)](http://nih.gov)
- EBI  
[The European Bioinformatics Institute < EMBL-EBI](http://ebi.ac.uk)
- EMBL  
[Heidelberg | EMBL.org](http://embl.org)

## Protocols:

- Globus
- sftp (ssh)
- Ncftp
- ncftpput

Whole Genome Sequencing (**WGS**)  
files size to transfer:

**1.8 PB**

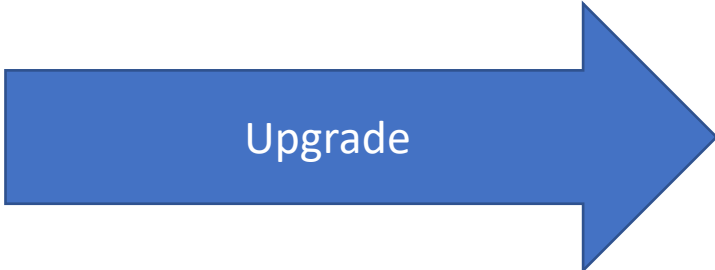
17 days at 10Gbps



# What's next!

## More resources

We have to be ready to host:

- Tens of Genomics Sequencers
  - Tens of Cryo-EM Microscopes
  - More WGS and other scientific data to transfer
  - More scientists and administrative employees (1000 – 2000)
- 
- More computing: HPC, GPU VDI
  - More storage: Tens PetaByte
  - More WAN bandwidth: 40 / 100 Gbps
  - More Security and Data Governance





# Fundamental deployment

It has to be rock solid!

## Stability

- Scientist work 24/7

and if they don't, many of their long running jobs do!

-Some researchers run irreproducible experiments

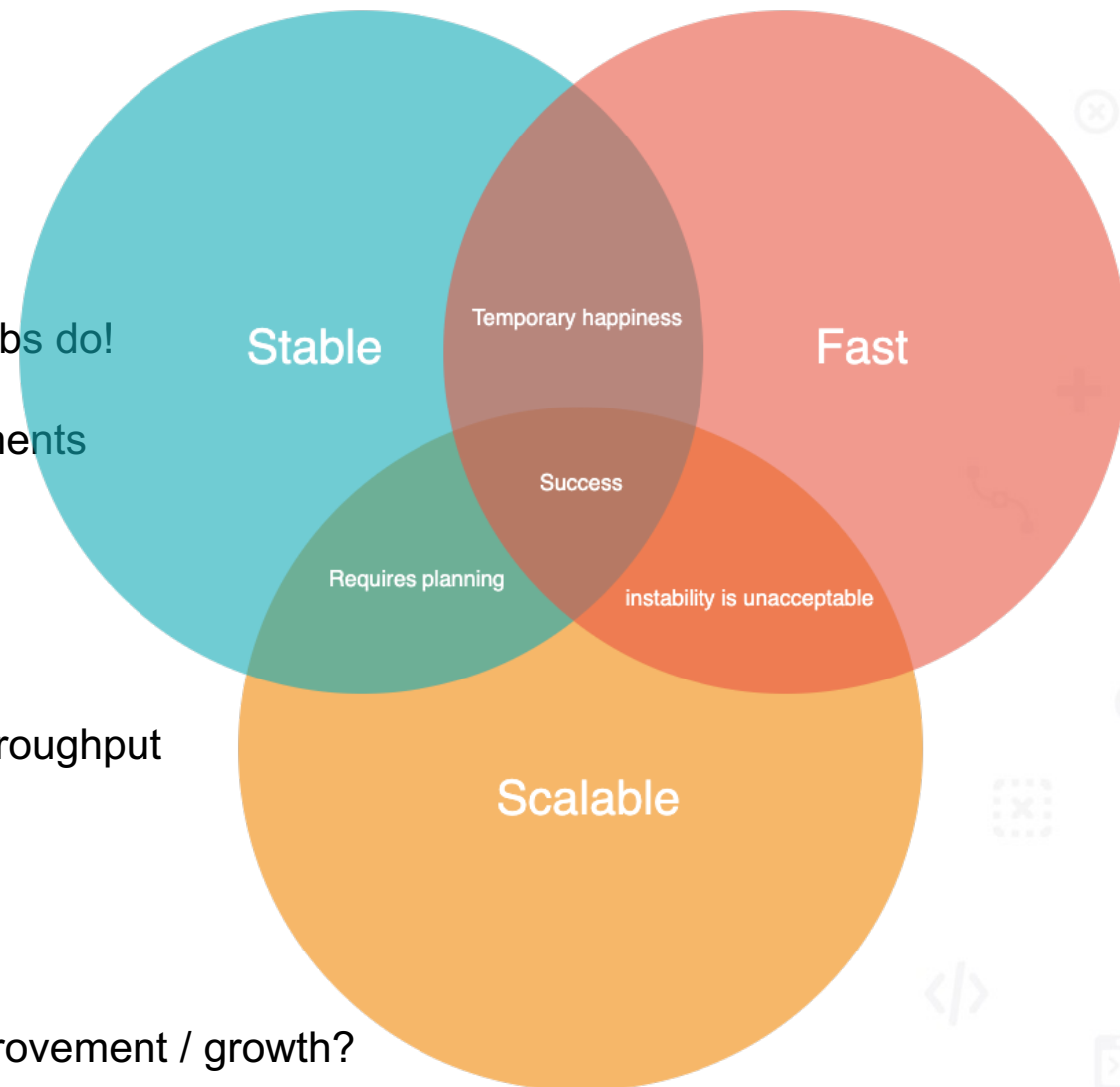
and some are expensive!

## Speed

- Some facilities require on top of stability throughput

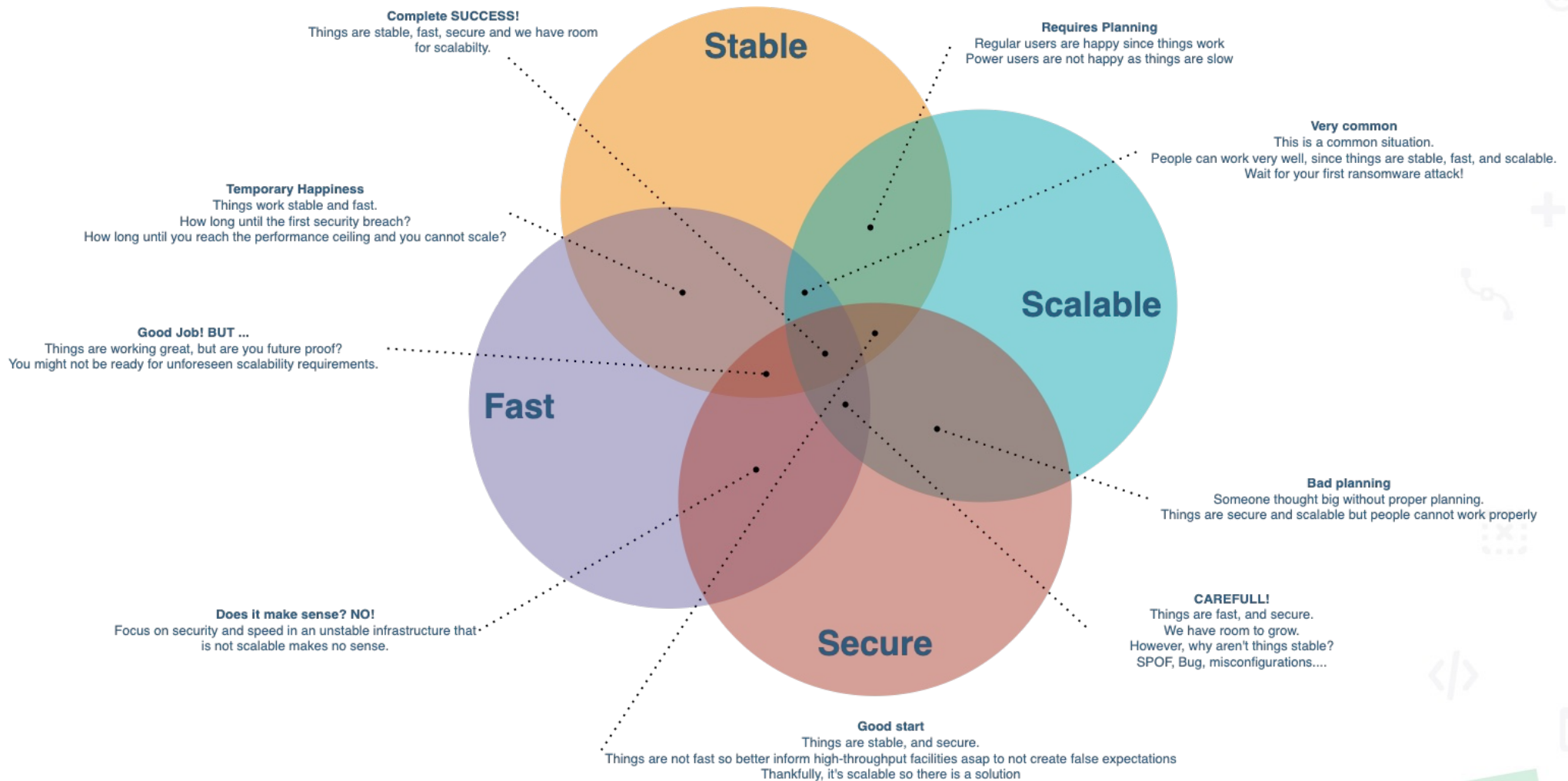
## Scalability

- Prepare for what's coming
- Are we future-proof? Is there room for improvement / growth?



# Let's add a bit of complexity

## What about Security?



# CyberSecurity

Network Segmentation

NAC

Multi Factor Authentication

Web Filtering

EDR-XDR

MDM

Vulnerability Assessment

Anti phishing Campaign

Patch Management



# Data Governance

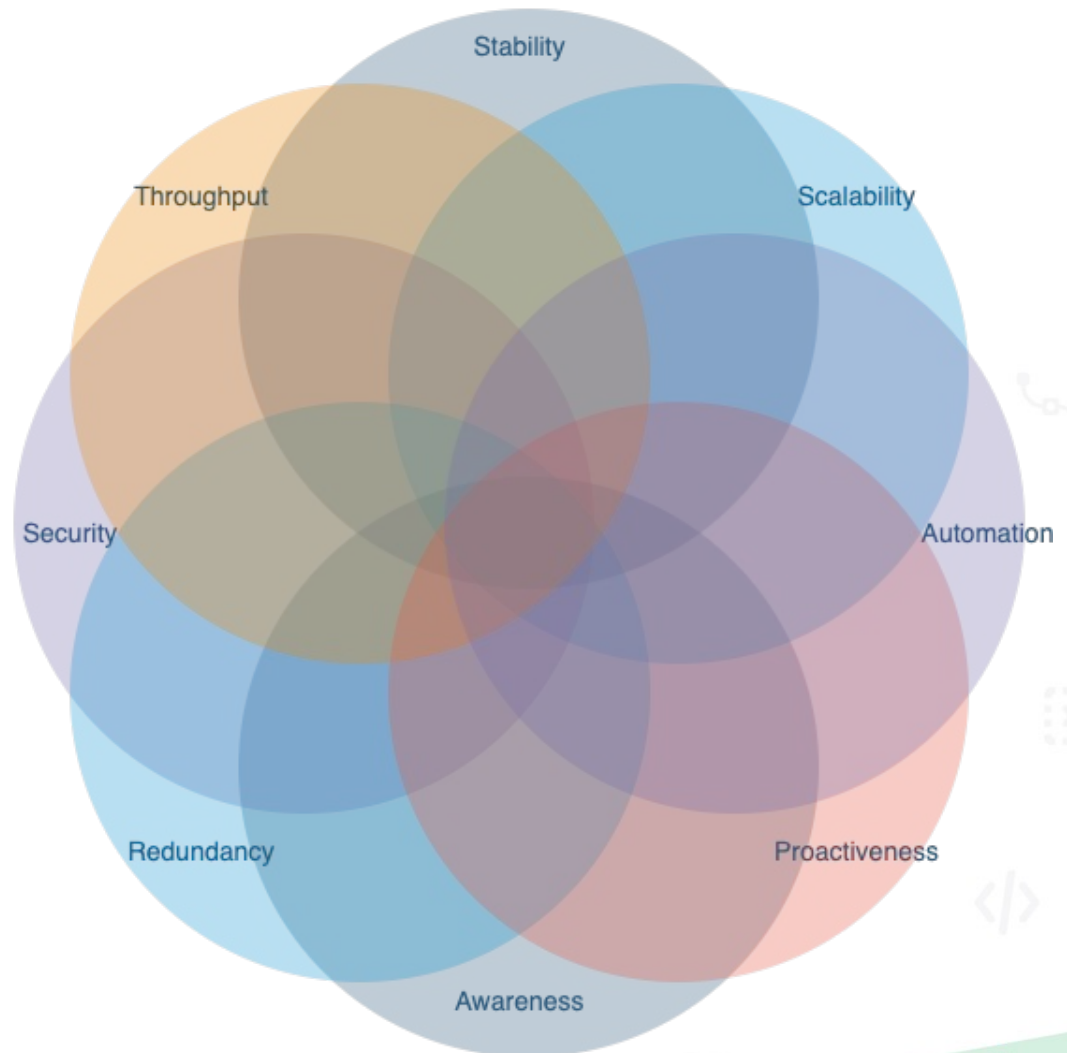
- ✓ **STRUCTURED APPROACH TO ACQUIRE EXTERNAL DATA  
(HEALTHCARE DATASETS FROM INSTITUTIONAL DATA PROVIDERS)**
- ✓ **GDPR COMPLIANCE - PROCESSING OF PERSONAL DATA**
- ✓ **STRUCTURED PROCESS TO MONITOR AND MANAGE DATA AND ITS  
QUALITY**
- ✓ **STRUCTURED DATA ARCHITECTURE ENABLING FASTER AND BETTER  
ANALYSIS**
- ✓ **REDUCTION OF THE RISKS ASSOCIATED WITH THE MANAGEMENT OF  
TREATMENTS CONTRARY TO THE LAW**

# What about more complexity

Redundancy, monitoring, alerting, automation... you name it.

$$\sum_{i=2}^n \frac{n!}{i! (n-i)!} = 2^n - n - 1$$

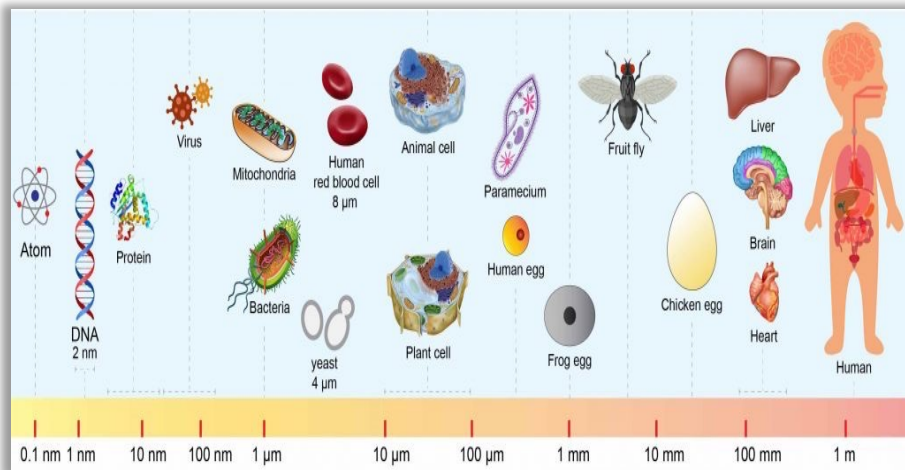
Variables	Overlapping regions
3	4
4	11
5	26
6	57
7	120
8	247



# Research Institutes WAN networks

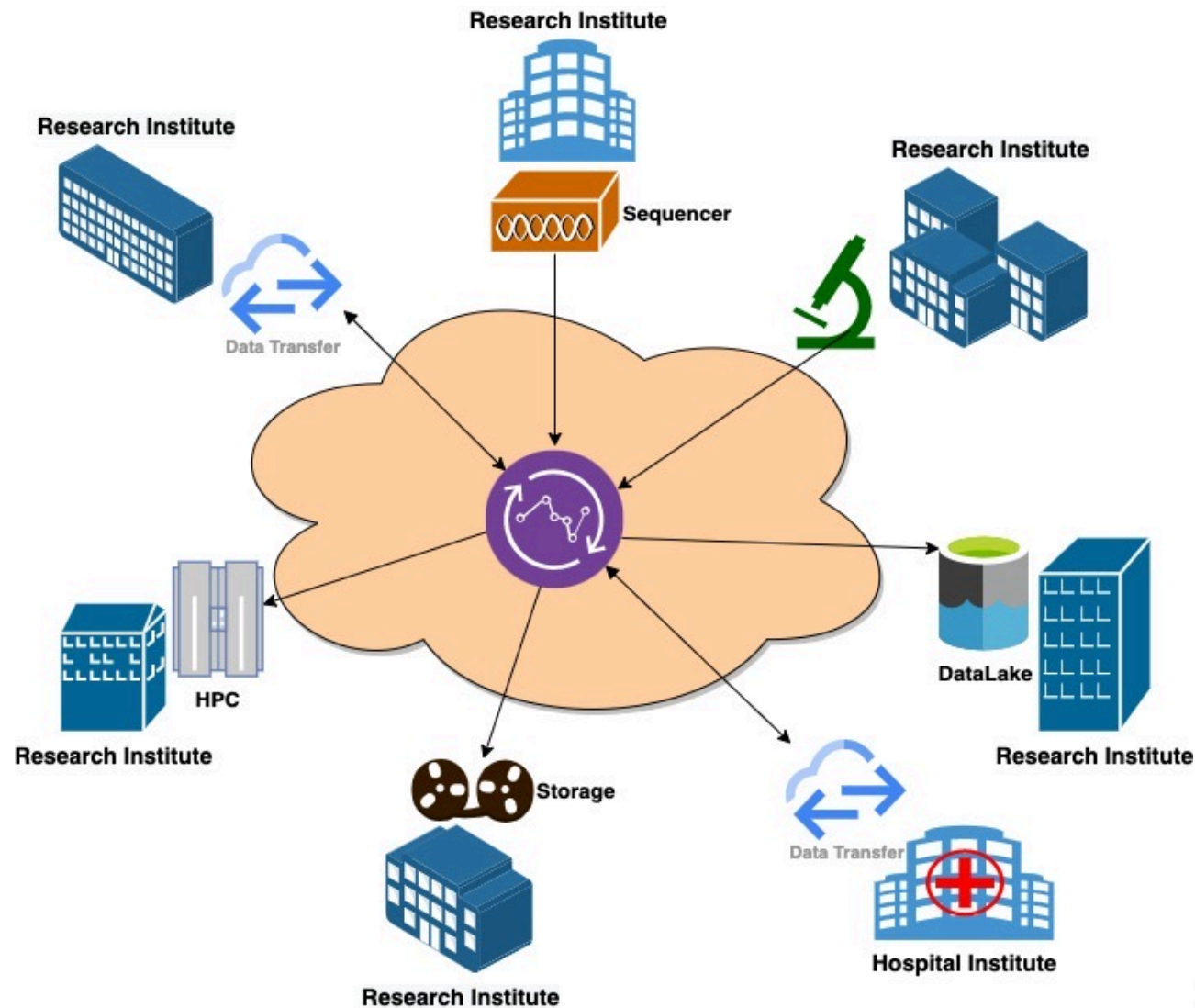
Research Institutes WAN networks are used for:

- Normal business operations including email, web browsing, O365, VPN SSL, SaaS among others. The network must also be built with security features.
- The scientific research process as scientists depend on this infrastructure to share, store, and analyse research data from many different external sources.
- Networks optimized for business operations are neither designed for nor capable of supporting the data movement requirements of data intensive science.



# Inter-Institutes connections

These connections could be possible



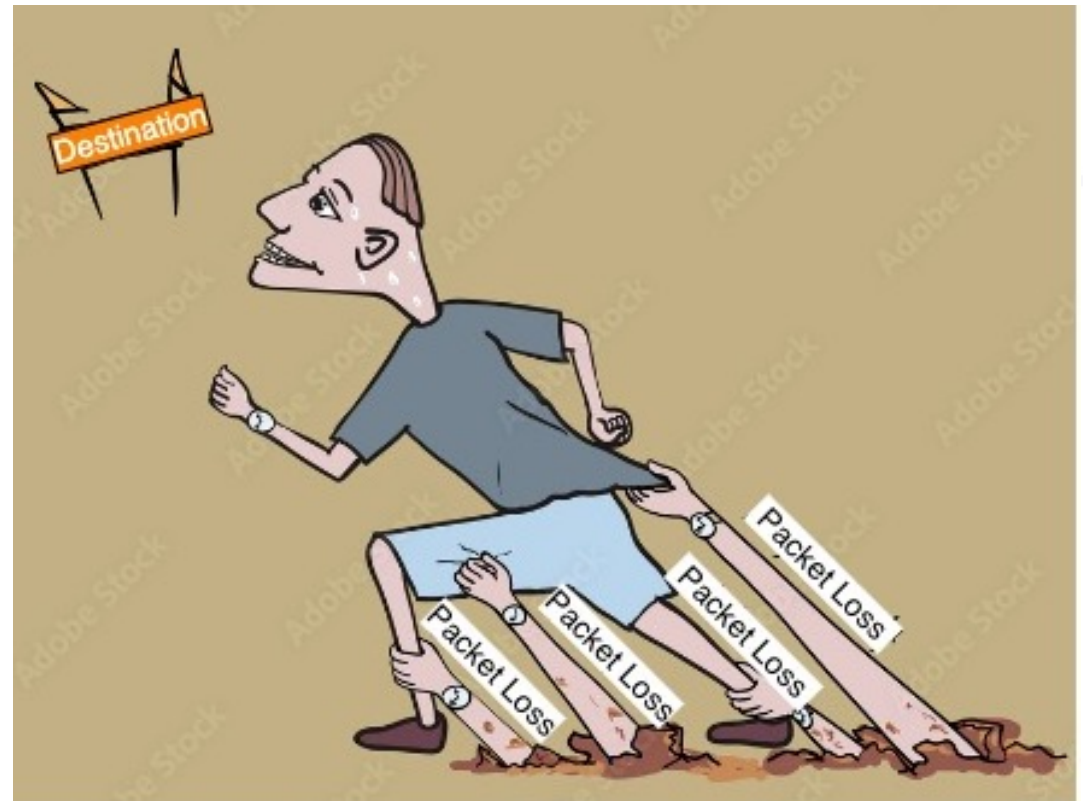


# Research Institutes WAN networks

## TCP performance issues:

- While most science applications that need reliable data delivery use TCP-based tools for data movement, TCP's interpretation of packet loss can cause performance issues..
- TCP interprets packet loss as network congestion, and so when loss is encountered TCP dramatically reduces its sending rate.
- The rate slowly ramps up again.
- This becomes more dramatic as the distance between communicating hosts is increased and with MTU path not in jumbo frames.

## TCP Marathon

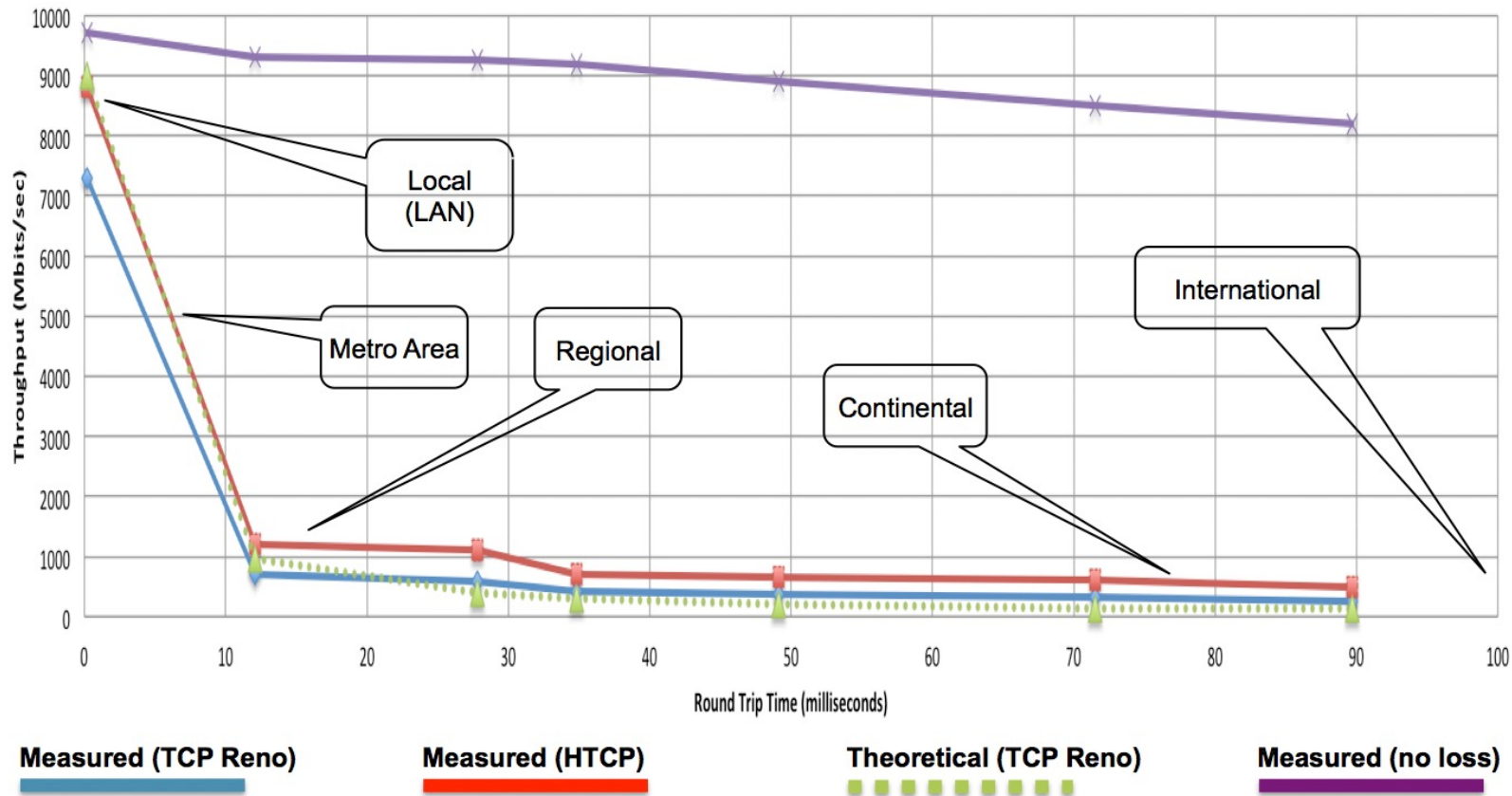


# Research Institutes WAN networks

## TCP performance issues:

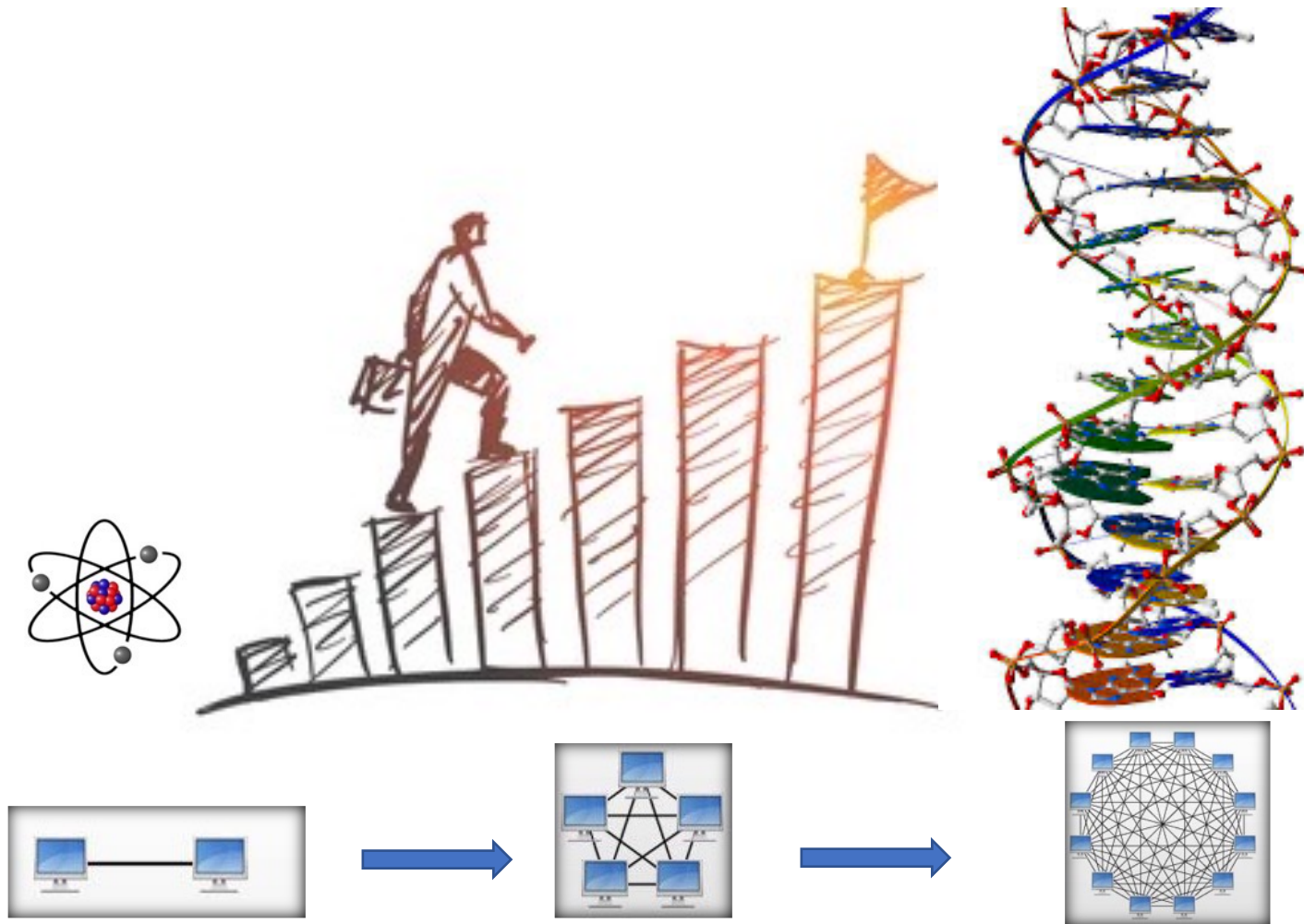
- A tiny amount of loss (much less than 1%) is enough to reduce TCP performance by over a factor of 50.

Throughput vs. increasing latency on a 10Gb/s link with **0.0046%** packet loss



# Research Institutes WAN networks

The challenge for a future DMZ Network



# Research Institutes WAN networks

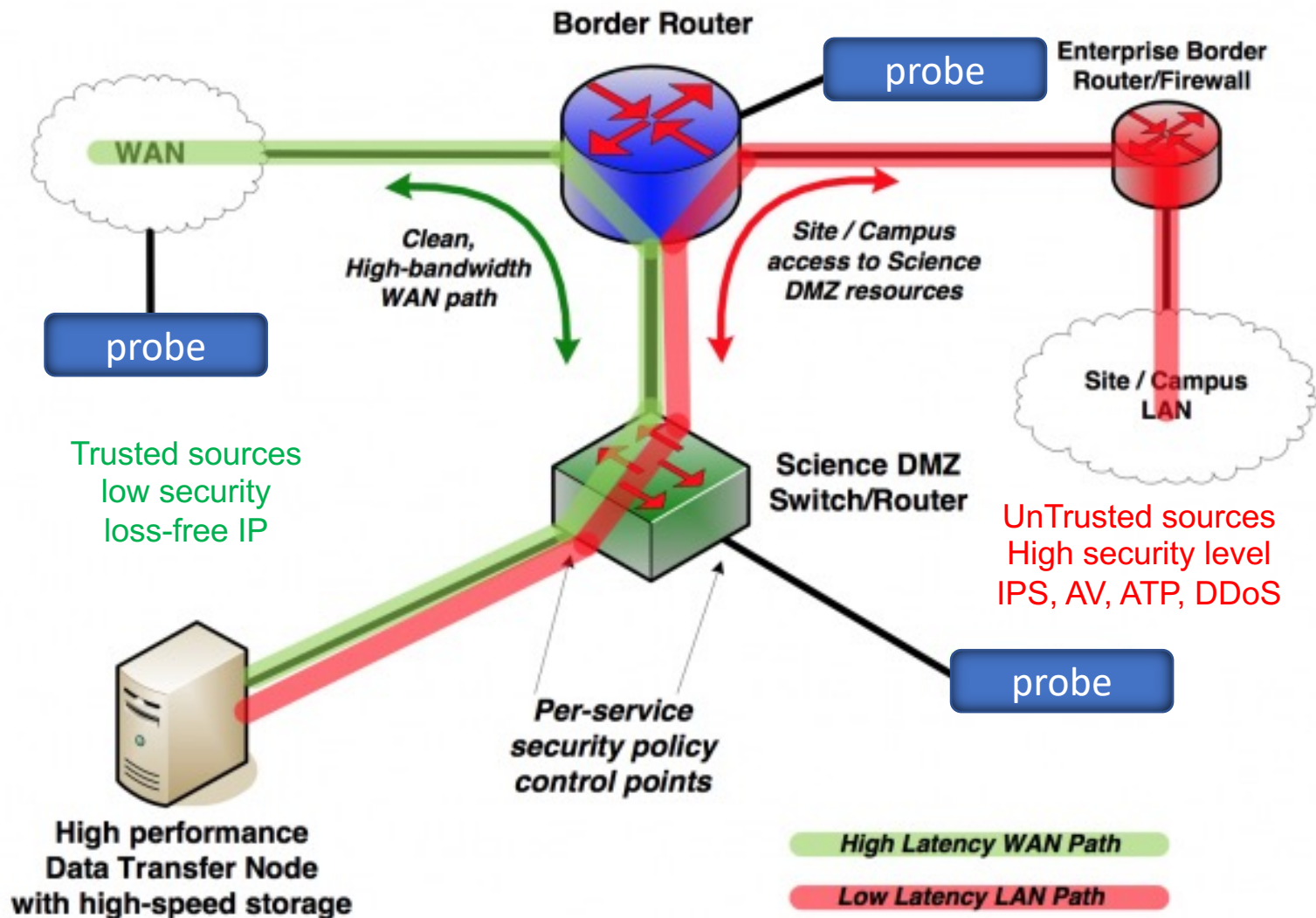
## Science DMZ

- The **Science DMZ** Model accomplishes this by explicitly creating a portion of the network that is specifically engineered for science applications and does not include support for general-purpose use.
- By separating the high-performance science network (the Science DMZ) from the general-purpose network, each can be optimized without interfering with the other.
- The Science DMZ model allows a laboratory, campus, or scientific facility to build a special-purpose infrastructure that can provide the necessary services to allow high-performance applications to be successful.



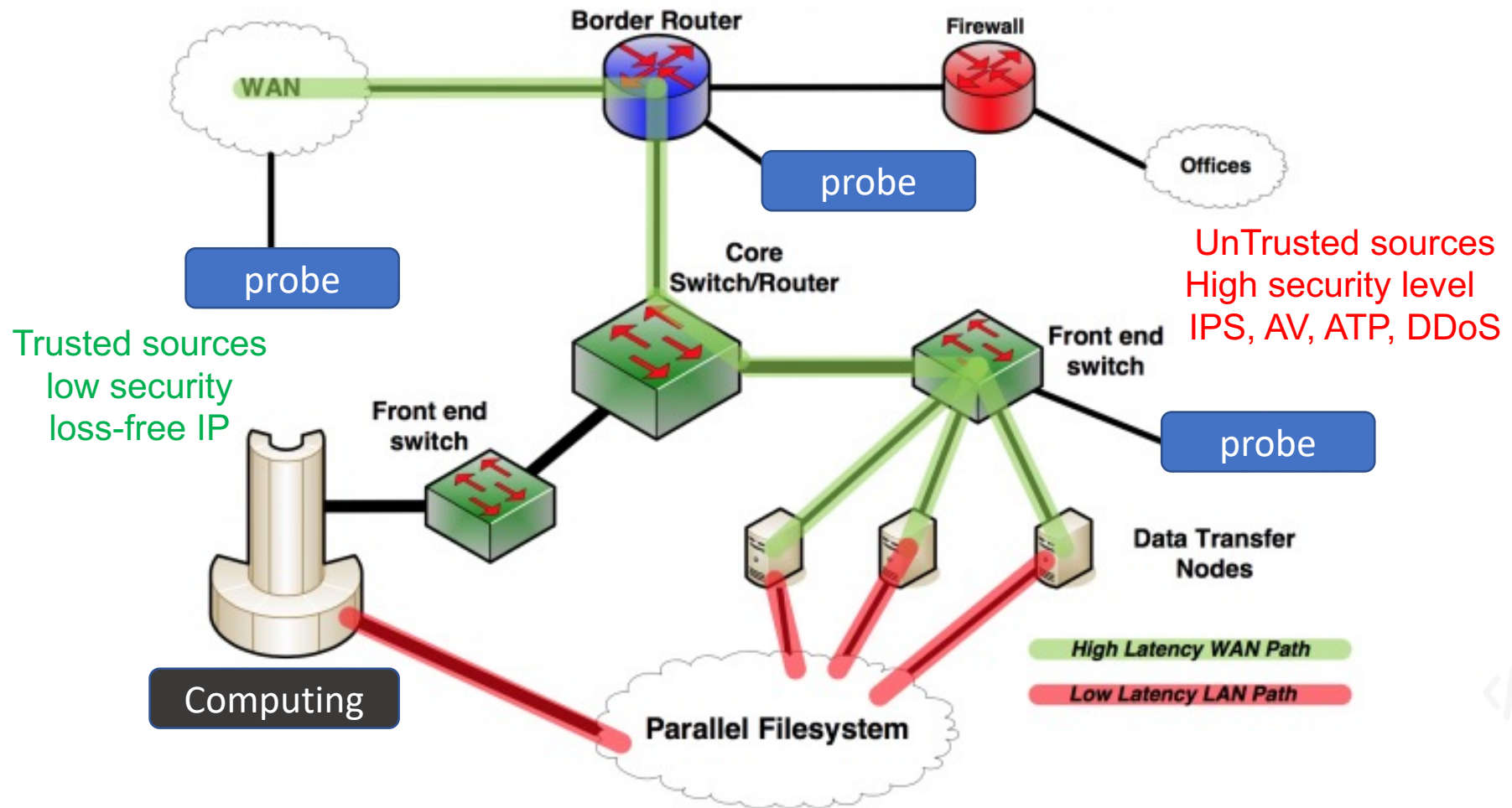
# Simple Science DMZ architecture

High-volume bulk data transfer, remote experiment control



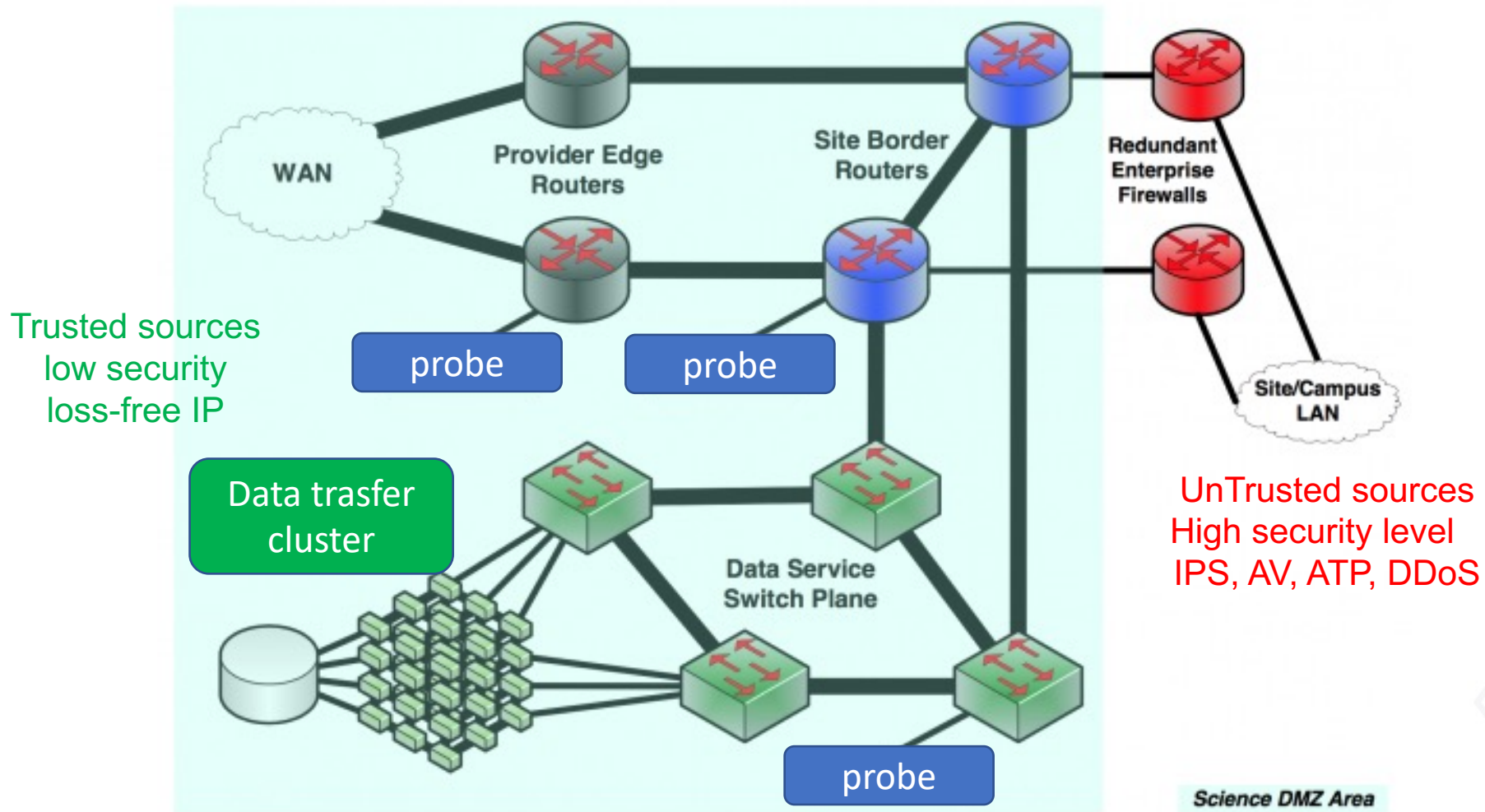
# Science DMZ architecture

## High-volume data processing transfer, remote experiment control



# Science DMZ architecture

High-volume parallel bulk data transfer, remote experiment control





Thank you

