

# Building cyberthreat models around genomic security

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# Genomics and Computing

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Primary question of the talk: How has the innovation in Next-Gen sequencing and in Synthetic Biology affected our cybersecurity risk models?

Avoid giving a recipe for mayhem, while still illuminating the realistic risks.

Where does the biological community fit in?

# What is a cybersecurity risk model?

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Identify

Protect

Detect

Respond

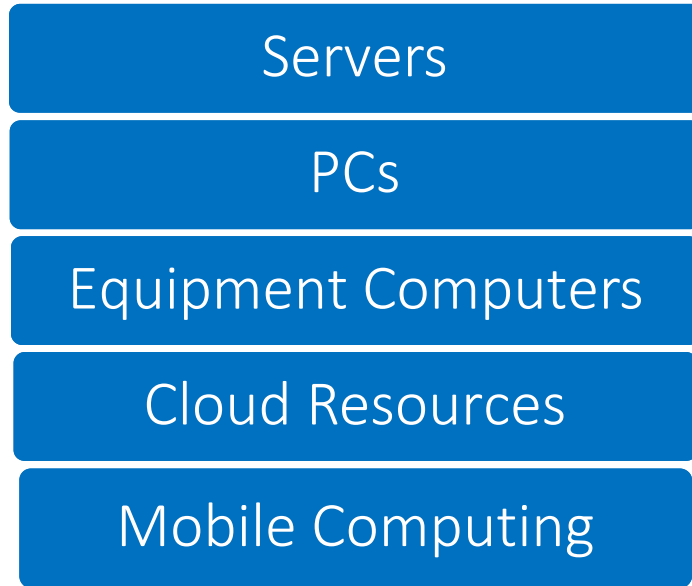
Recover

## Risk models

- Used to specify likely modes of attack
- Highlight areas for sensing attack
- Recovery mechanisms in the event of an attack.

# System profiling

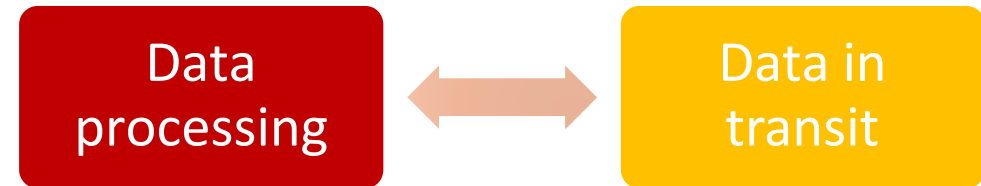
## Computational resources



Equipment

Data at rest

Data resources



Personnel

# What are the risks?

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Risk = Vulnerability x Threat x Impact x Probability

Risks at a facility doing sequencing genomics:

- Failure to complete work
- Release of protected data
  - Intellectual property
  - Personally identifying information
  - Secret information
- Destruction of data integrity
- Release of operational security and adversarial surveillance

# What are the risks?

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Risk = Vulnerability x Threat x Impact x Probability

Risks at a facility doing sequencing genomics – Hacking equivalent

- Failure to complete work – DDOS
- Release of protected data – Man-in-the-middle exfiltration and hacking theft
  - Intellectual property
  - Personally identifying information
  - Secret information
- Data manipulation – Social Media Hactivism
- Release of operational security and adversarial surveillance – APT

# Issue at hand: Genomics has moved from a scientific technique to an industry

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The <\$1000 genome has changed sequencing into a consumable.

- Illumina's NovaSeq and new technologies being developed by Complete Genomics (BGI company) suggest this will be ~\$100 by year's end.

The technology emerged before the safeguards or training in cyber-risks were in place.

Without an adequate threat model, automation exacerbates, rather than relieves the risk.

# How are genomics data different?

Comparison to credit card data  
CCs have an established threat model:

- 1) Secure data
- 2) Authenticate over encrypted network
- 3) Limit access
- 4) Regular vulnerability audit
- 5) Legal mechanisms for recuperation

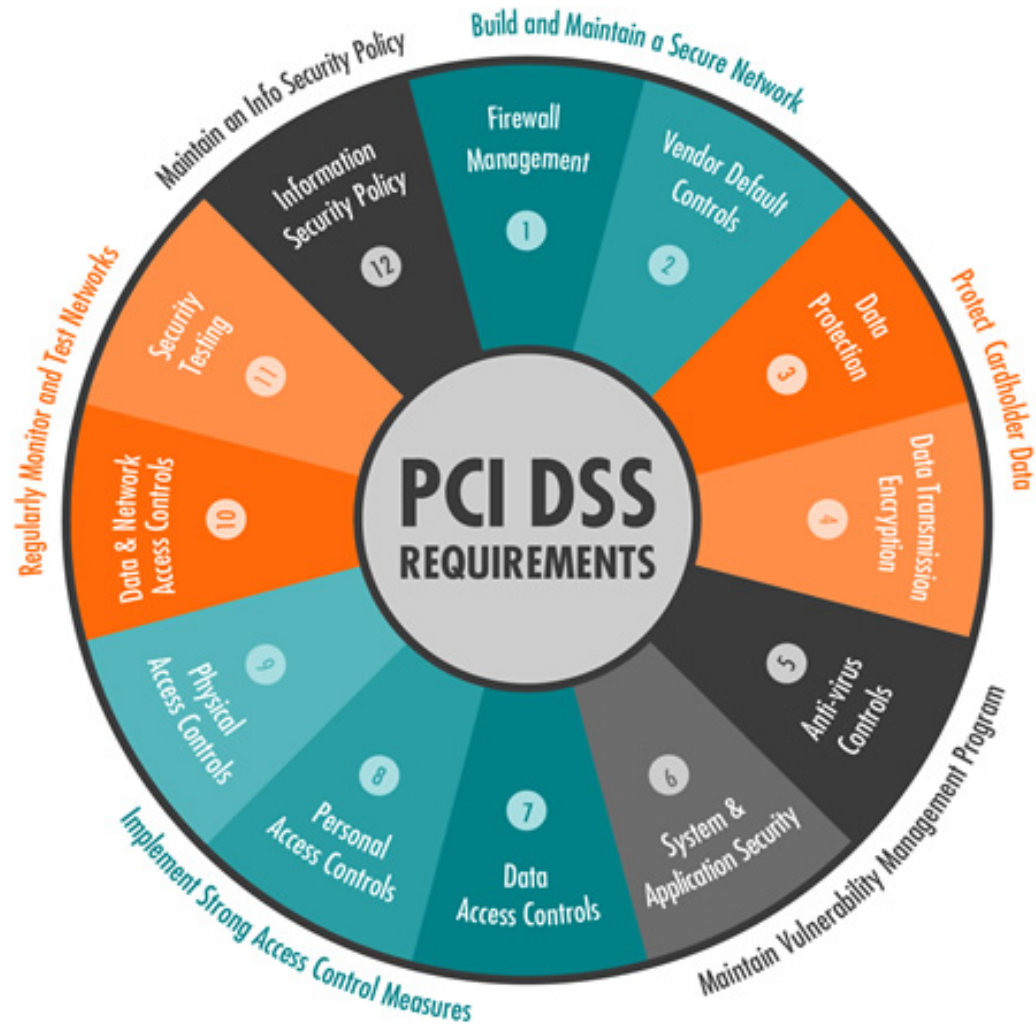
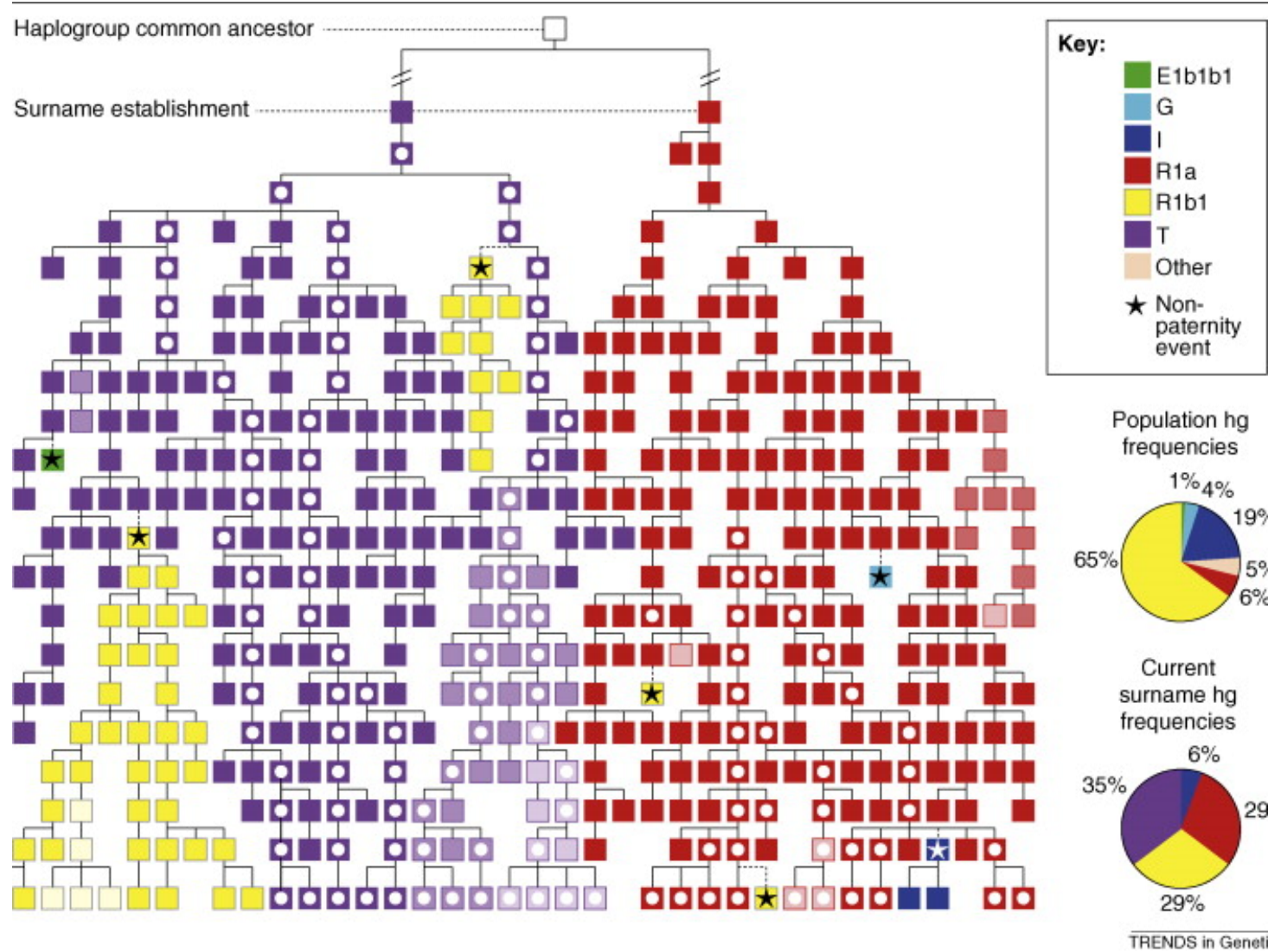


Image:  
<http://ryantech.com/>





# Genomic data are associational

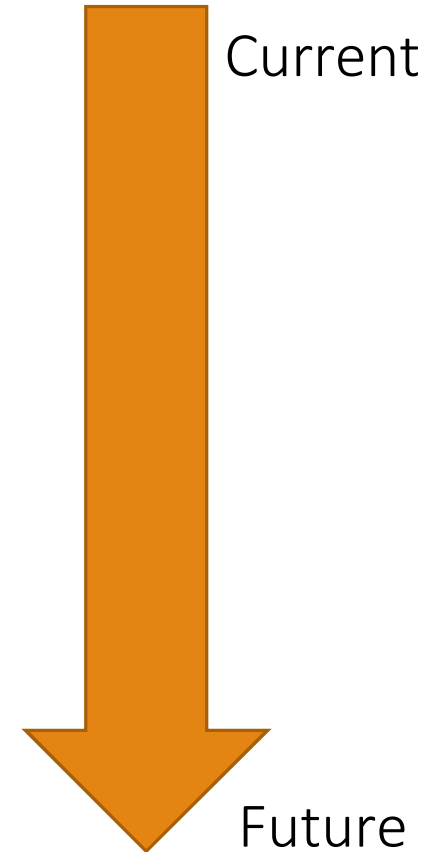
Every leaked genome leaks data about associated family members.

Asymptotically, this means that genomic data cannot be secured indefinitely.

# What is the risk space around privacy?

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- Paternity breach
- Privacy and identification
- Racial or at-risk subgroup identification
- Legal/forensic identification/manipulation
- Phenotype inference
- Genomic access controls
- Genomic targeting



# Recovering from genomic breach

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Fundamental maxim of genomic data breach: There is currently no model for recovery from genomic data release.

Genomic data are basically unchangeable through the life of the victim.

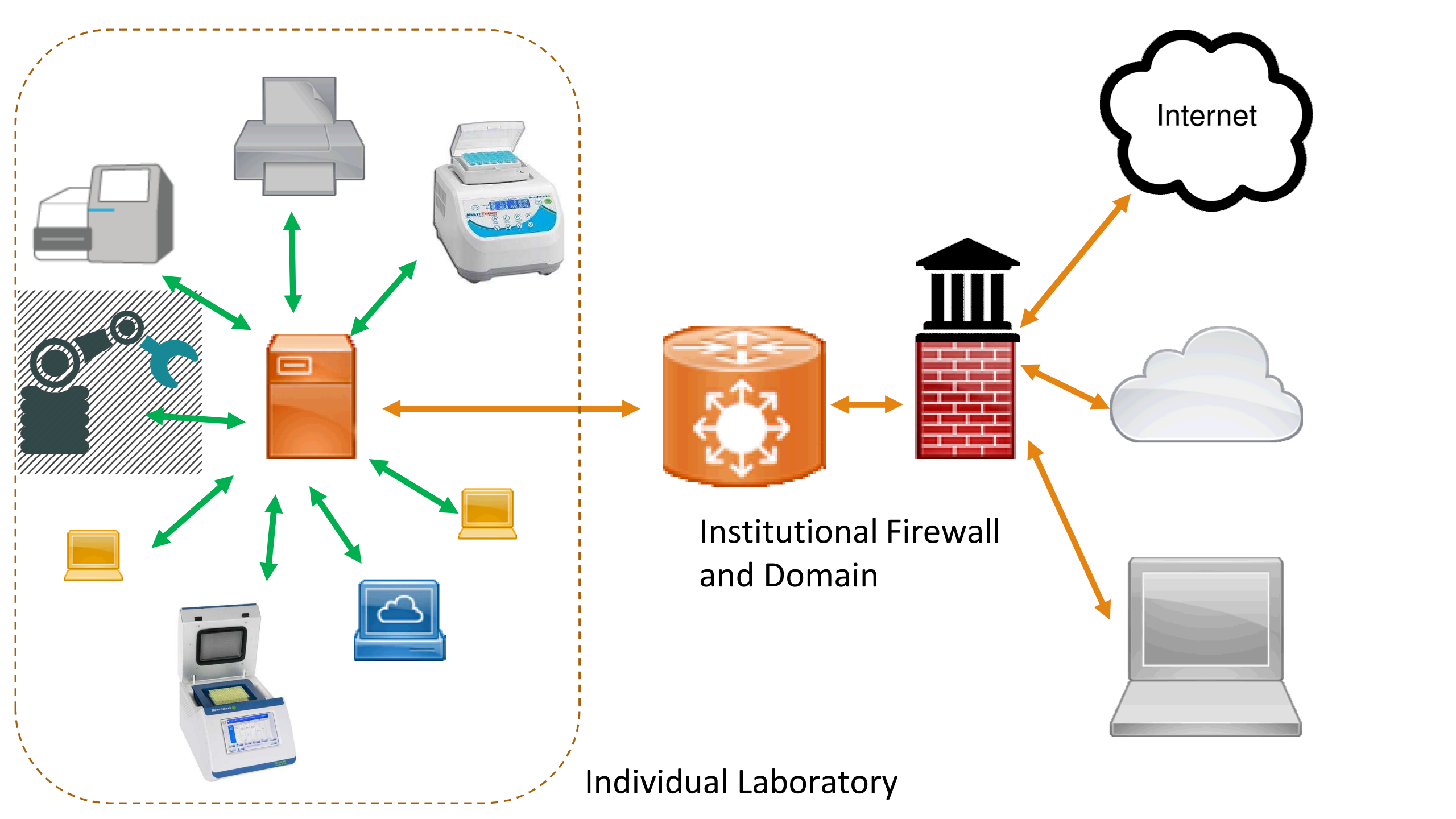
Leaked data can create new victims, through the associational nature of genomic data.

# Threat model 1: Firewall & Forget

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Mantra: Business as usual, data and operations are secure and compliant.

Use the security of the institution housing the system.



# Advantages

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- Rolling your own security system is DANGEROUS
- Institutions have IT departments
- Creates and establishes Access Control protocols
- Maintains compliance with larger institution

# Disadvantages

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- Sensors are outside realm of activity
- Machine-to-machine communication is assumed secure
- IT department may not be appraised of the level of risk they have signed on to
- To facilitate work, personnel may open unsecure channels to bypass firewall
- Many modern Next-Gen sequencing tools require cloud access

# Threat model 2: Security by obscurity

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Mantra: The system is too idiosyncratic or unsophisticated to be hacked.

Bizarre names and interactions

Idiosyncratic security protocols

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# Advantages

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- Level of reconnaissance necessary to do damage may not be worth return
- Conscious thought about threats
- Layered security is generally preferred

# Disadvantages

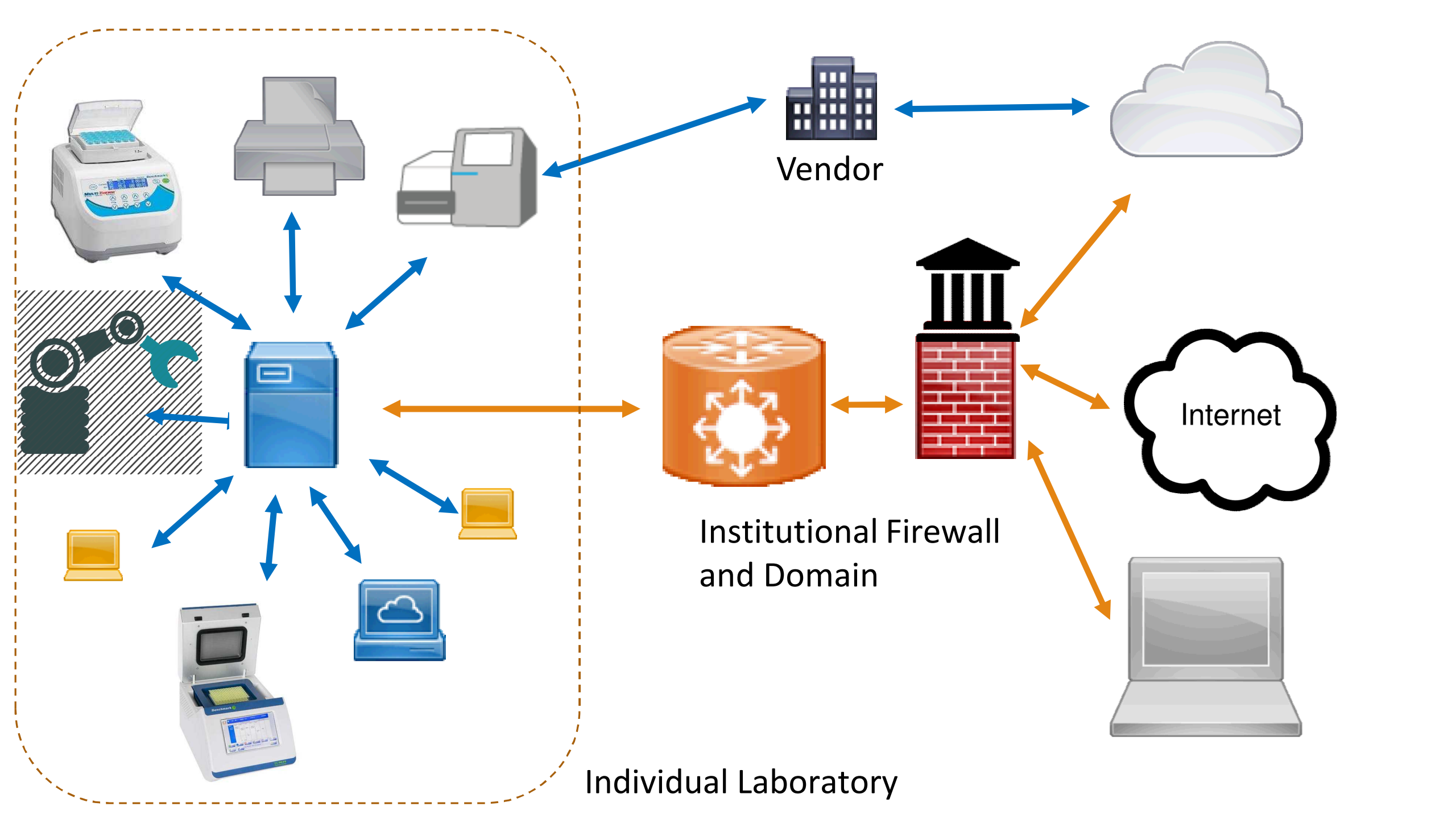
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- Rolling your own security is DANGEROUS
- May leave open huge gaps
- Threat model likely not comprehensive
- May not be compliant with institution
- Likely will not have secure ports

# Threat model 3: Leave it to the vendor

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Mantra: The less the lab interacts with security, the lower the chance that they will wreck it.



# Advantages

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- Able to simultaneously handle lab and institutional compliance
- Vendor has best understanding of machines and potential insecurities
- Allows genomics specific threat model

# Disadvantages

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- Dependent on service agreement
- A lab may involve multiple vendors
- May not be compliant with data provider's specifications
- What happens after service agreement runs out?

Question: What is the optimal network setup that perfectly solves the sequencing security problem?

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Answer: No....

All systems are vulnerable. One possible long term solution to sequence security is *fully homomorphic encryption (FHE)*.

FHE allows queries of encrypted data – but the data has to be fully assembled before it can be encrypted.

# What are the cyber-risks in synthetic biology?

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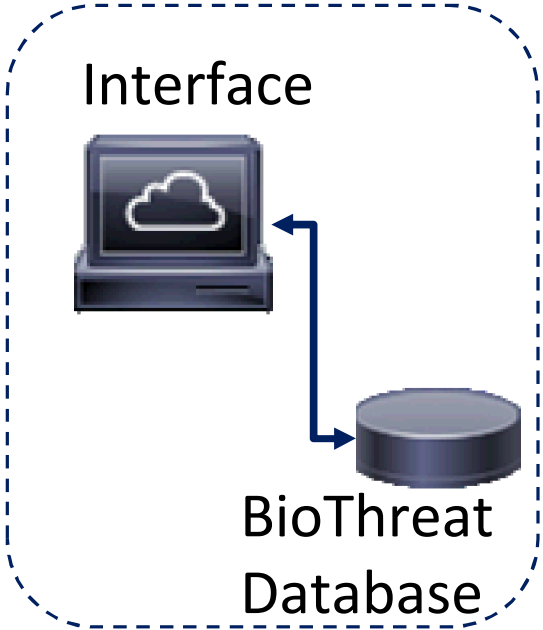
Risk = Vulnerability x Threat x Impact x Probability

Risks at a facility doing genetic/genomic manufacture:

- Sequencing risks present in manufacture as well
- Unintended manufacture



Synthesis, Assembly  
and Cloning



Design



Screening and Sequencing

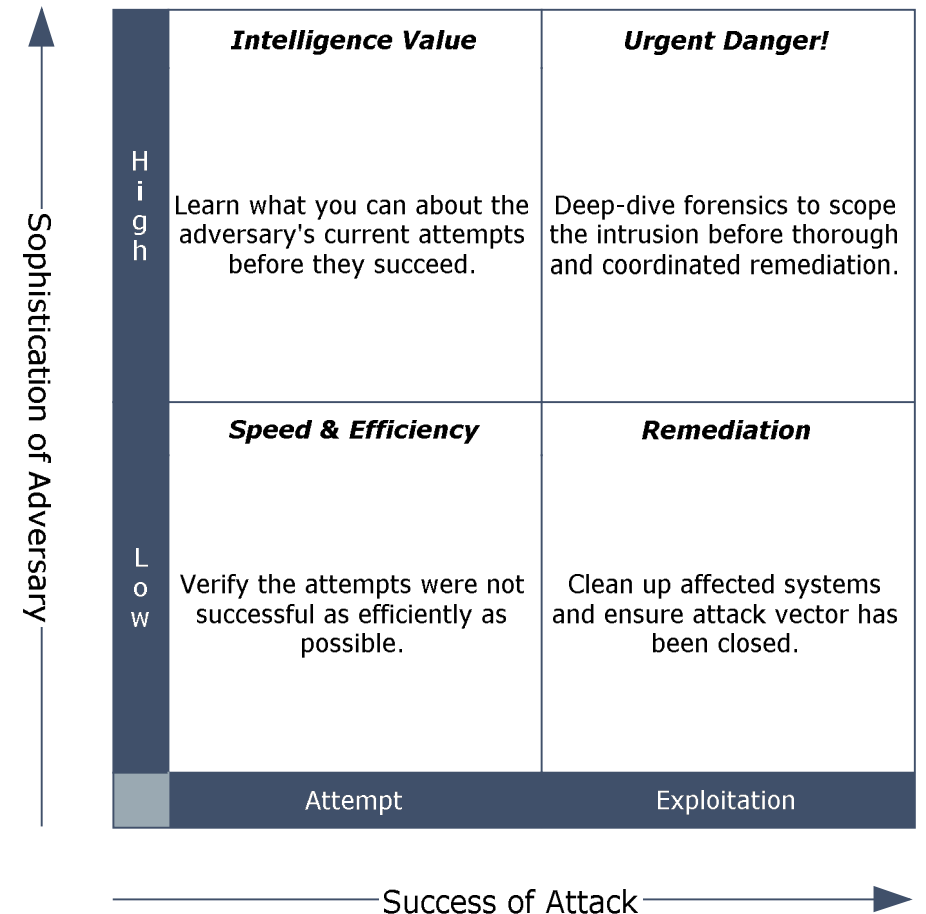


Control  
Server

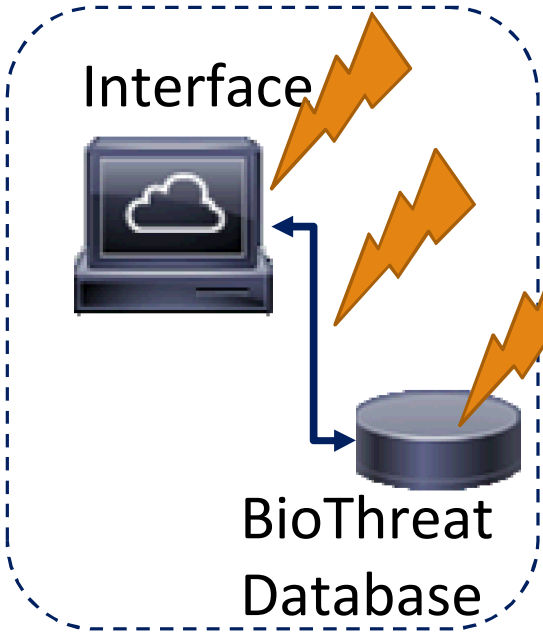
# Adversarial sophistication

Threat models typically take into account the sophistication of the adversary and the success of the attack

Once an adversary has command and control access, the sophistication of the adversary determines the response



Synthesis, Assembly  
and Cloning



Design



Screening and Sequencing



Control  
Server

# Conclusions

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- Next gen sequencing and synthetic biology has grown at a speed that has outpaced the security implications of the platform
- Desperate need for research on vulnerabilities in NGS systems
- The loss in security of genomic data has implications outside the original sequence
- There is a distinct and underappreciated risk of unintentional manufacture of synthetic biological material