AtlantTIC

Research Center for
Information & Communication Technologies

Secure Genomic Susceptibility Testing based on Lattice Encryption

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Outline

- Introduction
- Secure Signal Processing
- Private Genomic Susceptibility Testing
- Previous Solutions
- Proposed Scheme
- Security and Performance Evaluation
- Conclusions

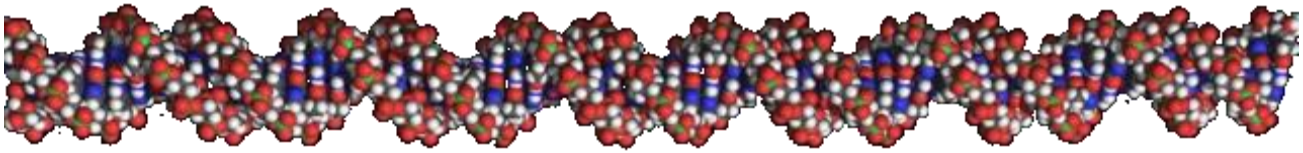
Introduction

Genomic Privacy

Privacy-Preserving Genomic Data Processing

Motivation

- Rapid advances in genomic research and sequencing
- Growing volume of data has to be outsourced
- Inherently sensitive information in DNA (individual and relatives)



Need for privacy-preserving processing!

Privacy-Preserving Genomic Data Processing

Objectives

- Personalized health-care (disease susceptibility tests)
- The most common variants in genome are SNPs (Single Nucleotide Polymorphisms)
- SNPs are suitable for running disease susceptibility tests

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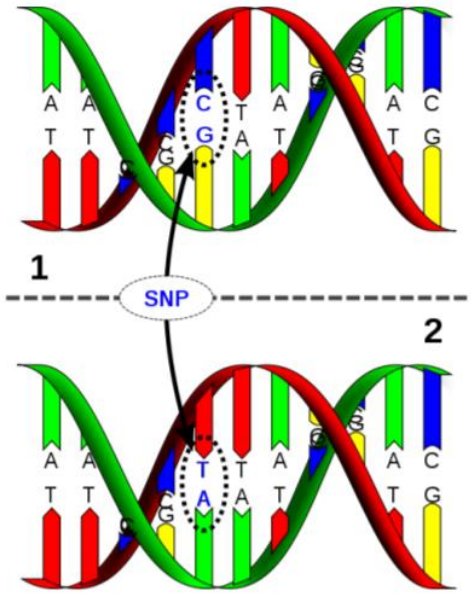
The presence and absence of SNPs give information about the susceptibility to a particular disease



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Contributions

- An efficient protocol for performing encrypted genomic susceptibility tests
- Our solution outperforms previous solutions in both computation, bandwidth and storage

Secure Signal Processing

Privacy tools from SSP

Privacy Tools from SSP

Homomorphic Encryption

- Example: Paillier (additive)
  - $E_k(x) = (1 + x \cdot n) \cdot r^n \; mod \; n^2$
  - $E_k(x + y) = E_k(x) \cdot E_k(y) \; mod \; n^2, E_k(x \cdot k) = E_k(x)^k \; mod \; n^2$
- SHE and FHE (both additions and multiplications)

$$(P, +, \cdot) \longrightarrow^{E_k} (C, +, \cdot)$$

- SHE example: Lauter cryptosystem (RLWE based cryptosystem)
- Both homomorphic cyclic convolutions and additions[1]

[1] A. Pedrouzo-Ulloa, J. R. Troncoso-Pastoriza, and F. Pérez-González, "Number Theoretic Transforms for Secure Signal Processing," *in IEEE Transactions on Informations Forensics and Security*, vol.12, no. 5, pp. 1125-1140, May 2017.

Lattice-based SHE Cryptosystem

- Somewhat homomorphic cryptosystem

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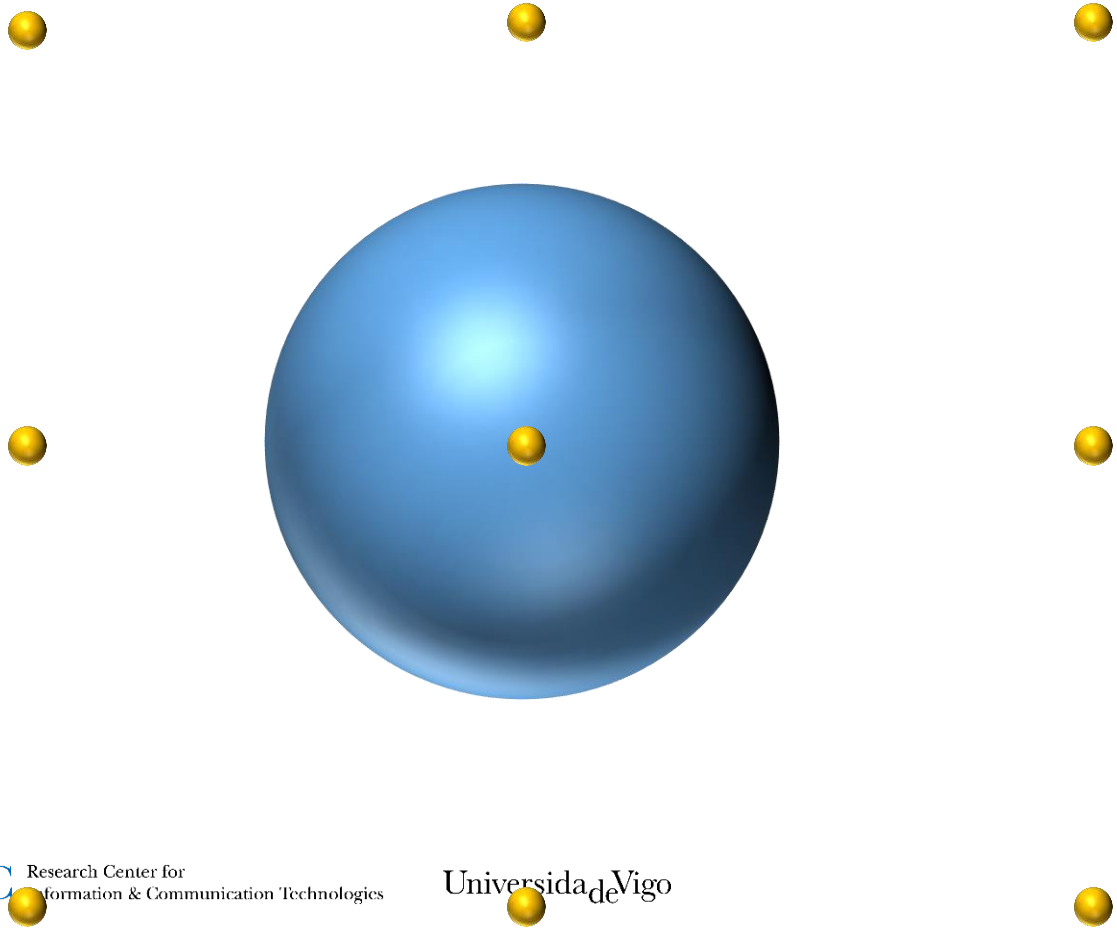

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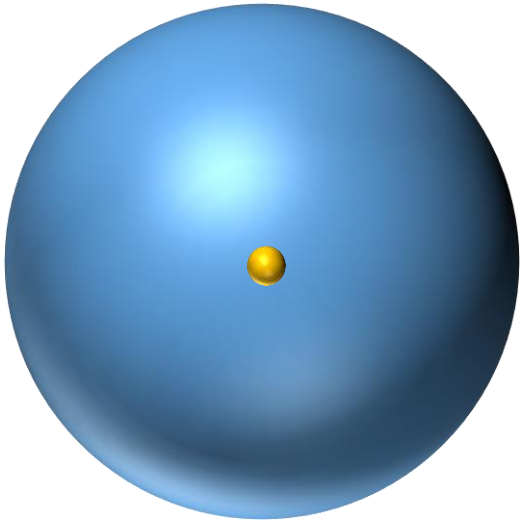

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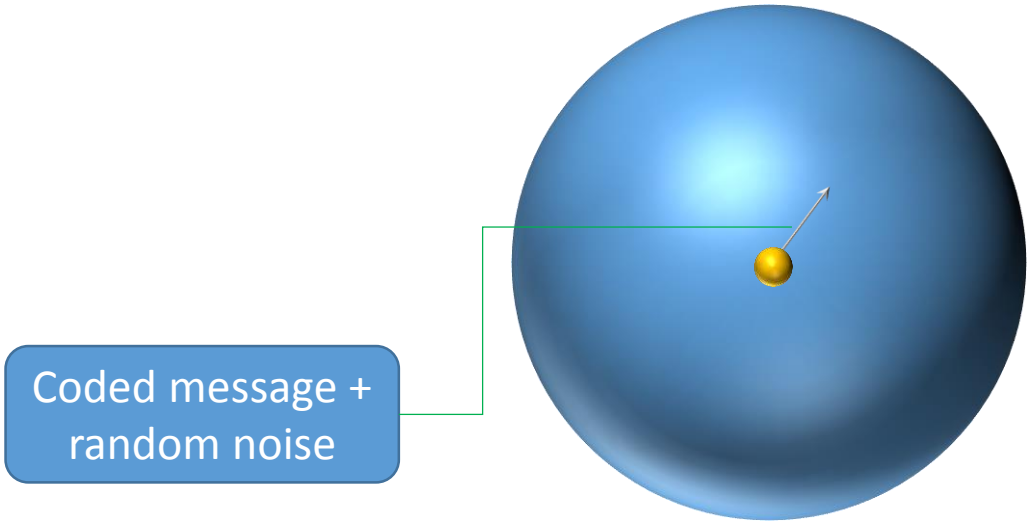

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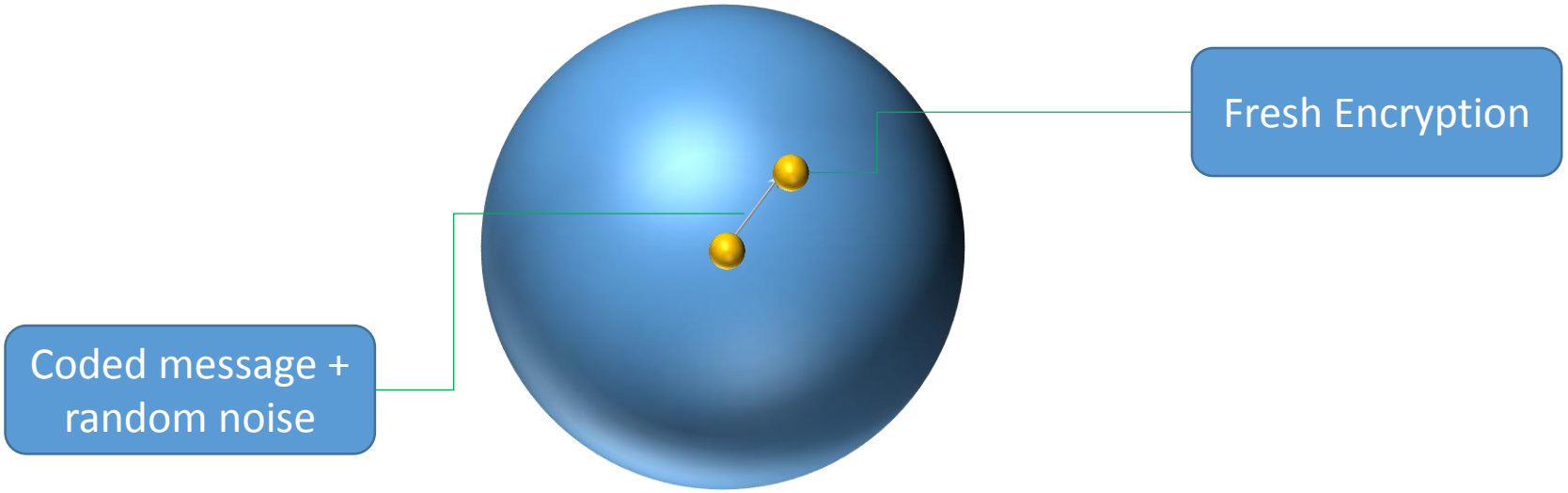

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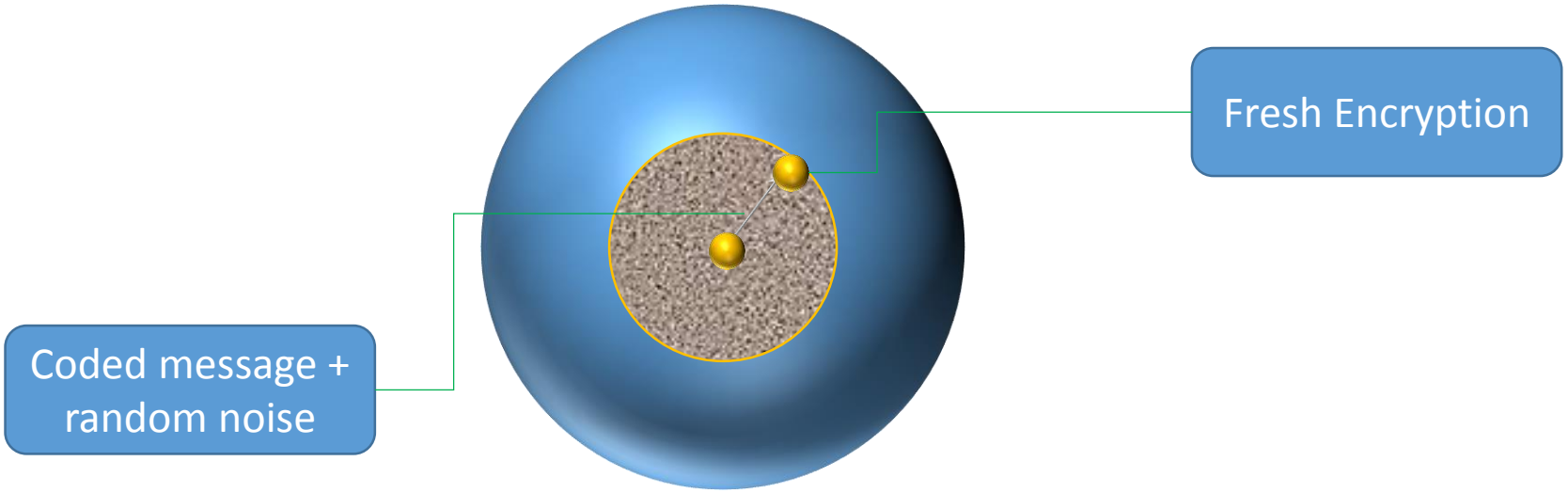

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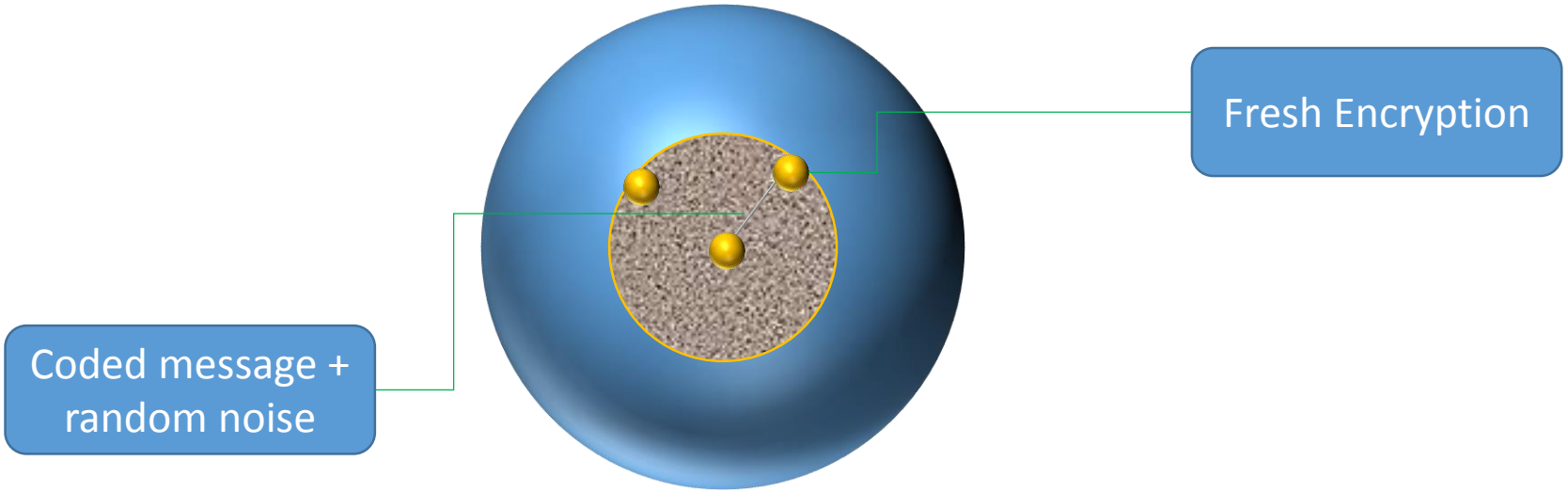

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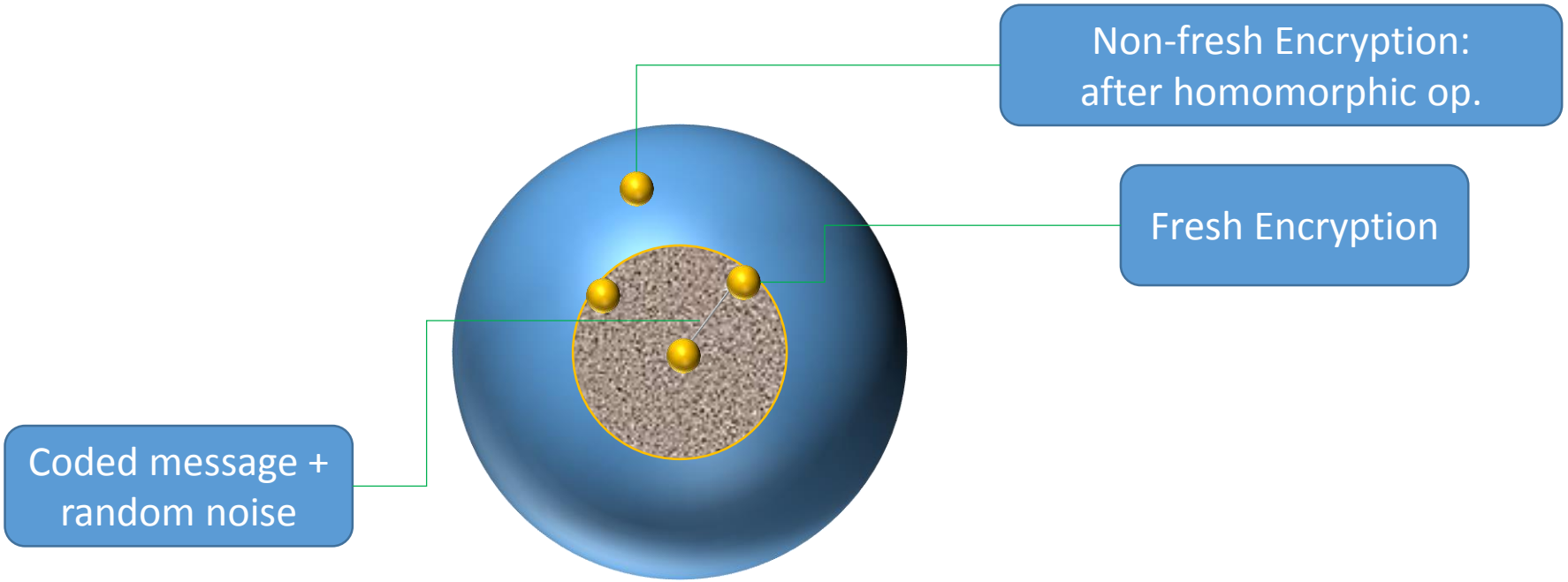

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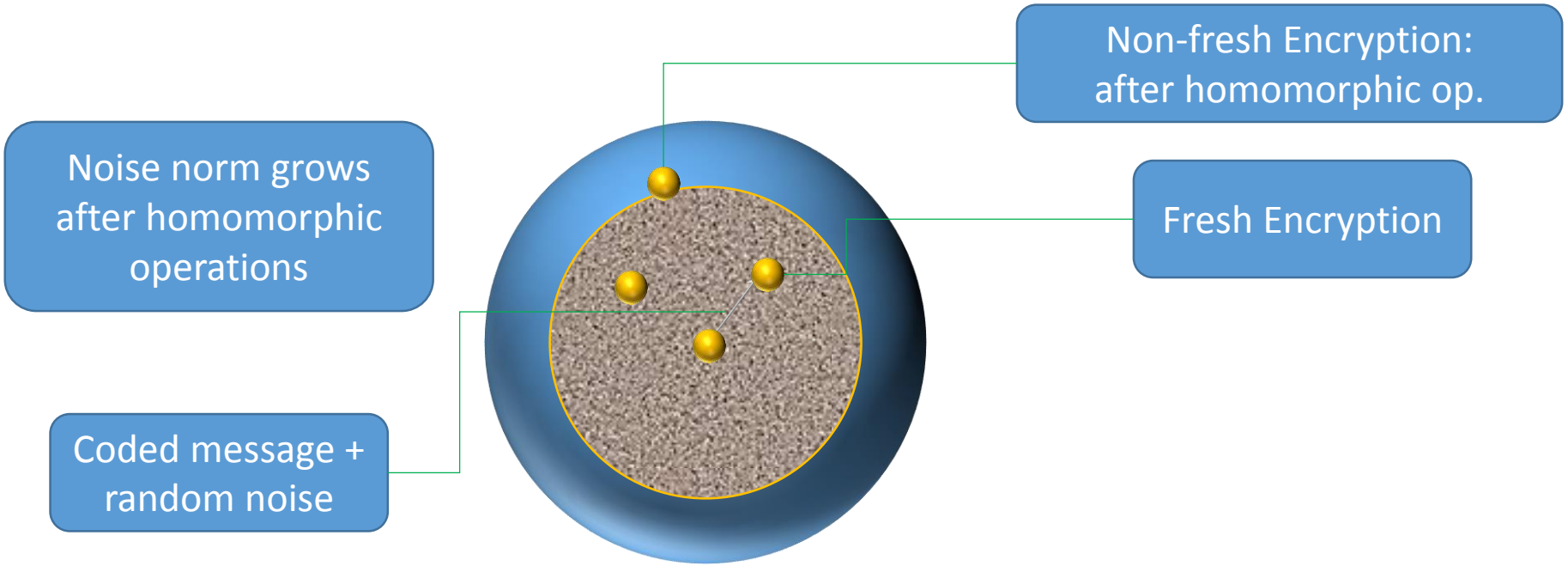

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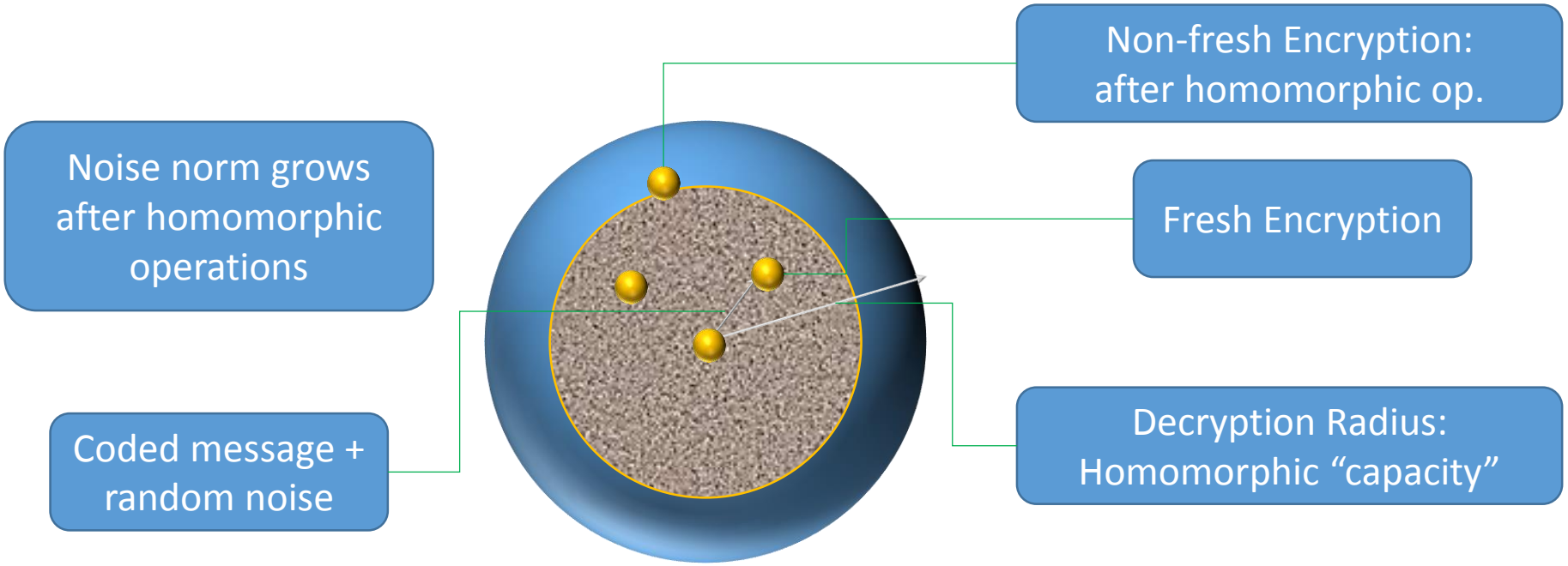

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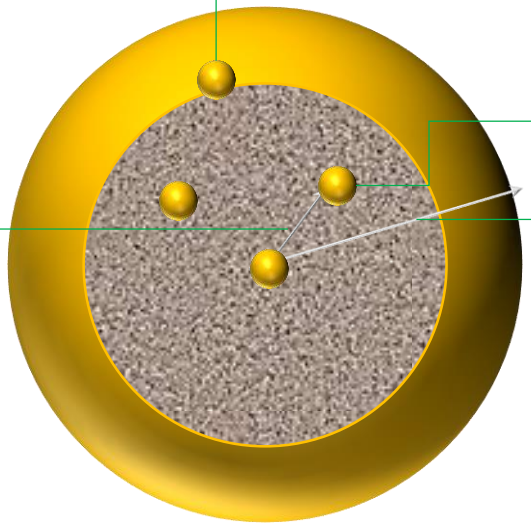

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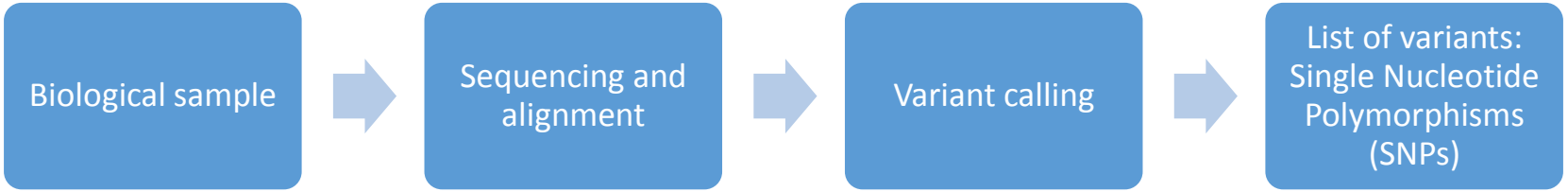
  - Can execute a bounded number of homomorphic operations
  - FHE can get unlimited homomorphic operations
    - FHE is too costly
    - As we know the number of homomorphic operations beforehand, SHE is a perfect candidate for our purposes

Private Genomic Susceptibility Testing

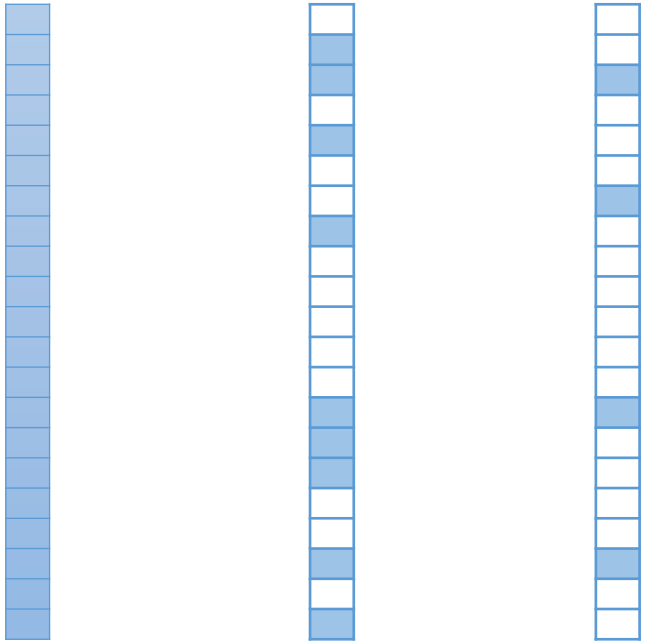
Privacy-preserving genomic susceptibility testing



Potential SNPs ~50M

Patient SNPs ~4M

Relevant SNPs (markers)



Probabilities

$pr_0^{X,i}, pr_1^{X,i}$

$pr_0^{X,j}, pr_1^{X,j}$

$pr_0^{X,k}, pr_1^{X,k}$

$pr_0^{X,l}, pr_1^{X,l}$

Contributions

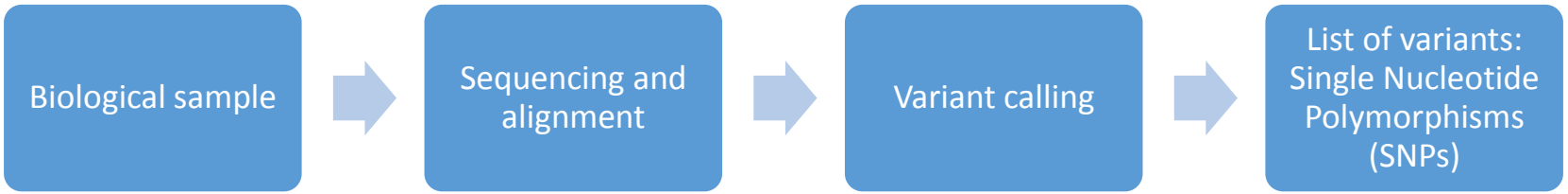
$c^{X,i}$

$c^{X,j}$

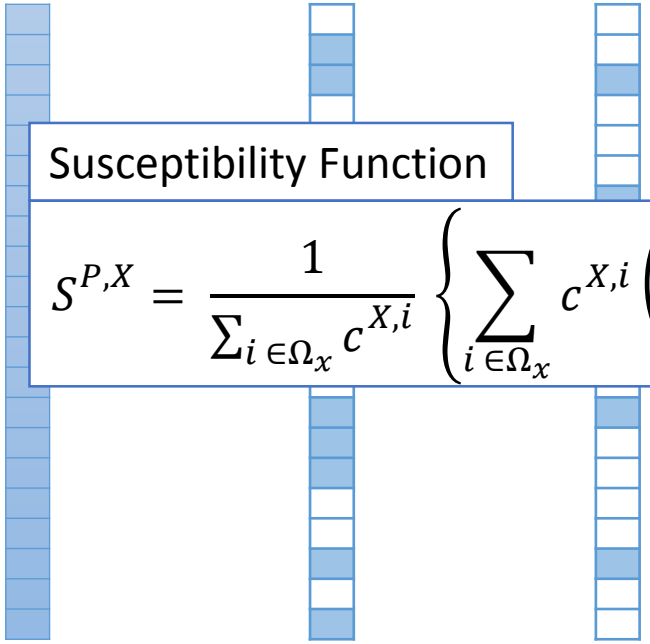
$c^{X,k}$

$c^{X,l}$

Privacy-preserving genomic susceptibility testing



Potential SNPs ~50M | Patient SNPs ~4M | Relevant SNPs (markers) | Probabilities | Contributions



$pr_0^{X,i}, pr_1^{X,i}$ $c^{X,i}$

$pr_0^{X,j}, pr_1^{X,j}$ $c^{X,j}$

$pr_0^{X,k}, pr_1^{X,k}$ $c^{X,k}$

$pr_0^{X,l}, pr_1^{X,l}$ $c^{X,l}$

Susceptibility Function

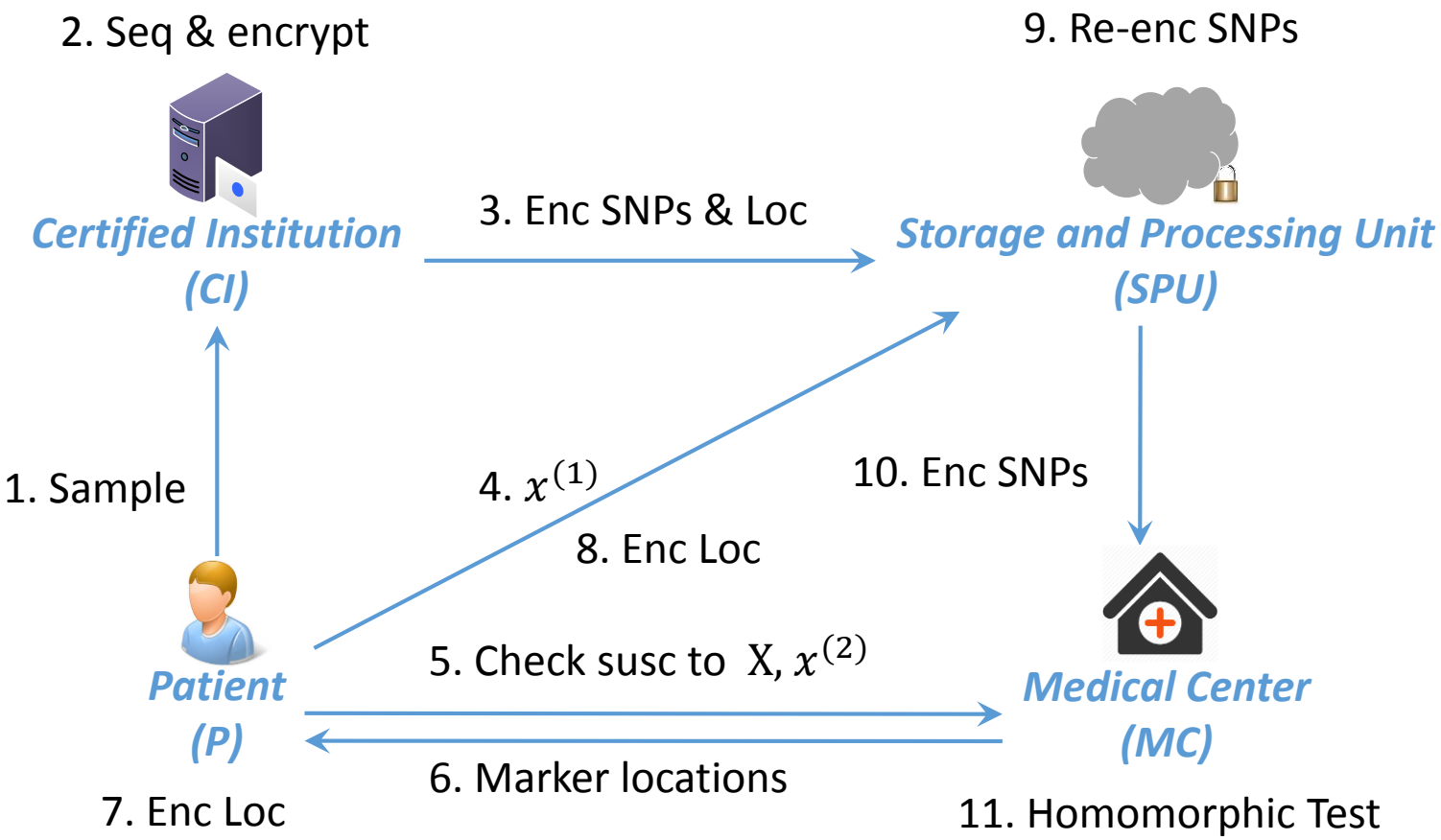
$$S^{P,X} = \frac{1}{\sum_{i \in \Omega_x} c^{X,i}} \left\{ \sum_{i \in \Omega_x} c^{X,i} \left(\frac{pr_0^{X,i}}{0-1} \left[SNP^{p,i} - 1 \right] + \frac{pr_1^{X,i}}{1-0} \left[SNP^{p,i} - 0 \right] \right) \right\}$$

Previous Schemes

Ayday et al. [2]

[2] E. Ayday, J. L. Raisaro, and J. P. Hubaux, "Privacy-Enhancing Technologies for Medical Tests Using Genomic Data," *in 20th Annual Network & Distributed System Security Symposium NDSS*, San Diego, CA, USA, Feb. 2013.

Privacy-preserving genomic susceptibility testing



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Paillier-encrypted Susceptibility

$$S_E^{P,X} = \prod_{i \in \Omega_x} \left(\left[SNP_E^{p,i} \cdot (-1)_E \right]^{\frac{-c^{X,i} \cdot pr_0^{X,i}}{\sum_{i \in \Omega_x} c^{X,i}}} \cdot \left[SNP_E^{p,i} \right]^{\frac{c^{X,i} \cdot pr_1^{X,i}}{\sum_{i \in \Omega_x} c^{X,i}}} \right)$$

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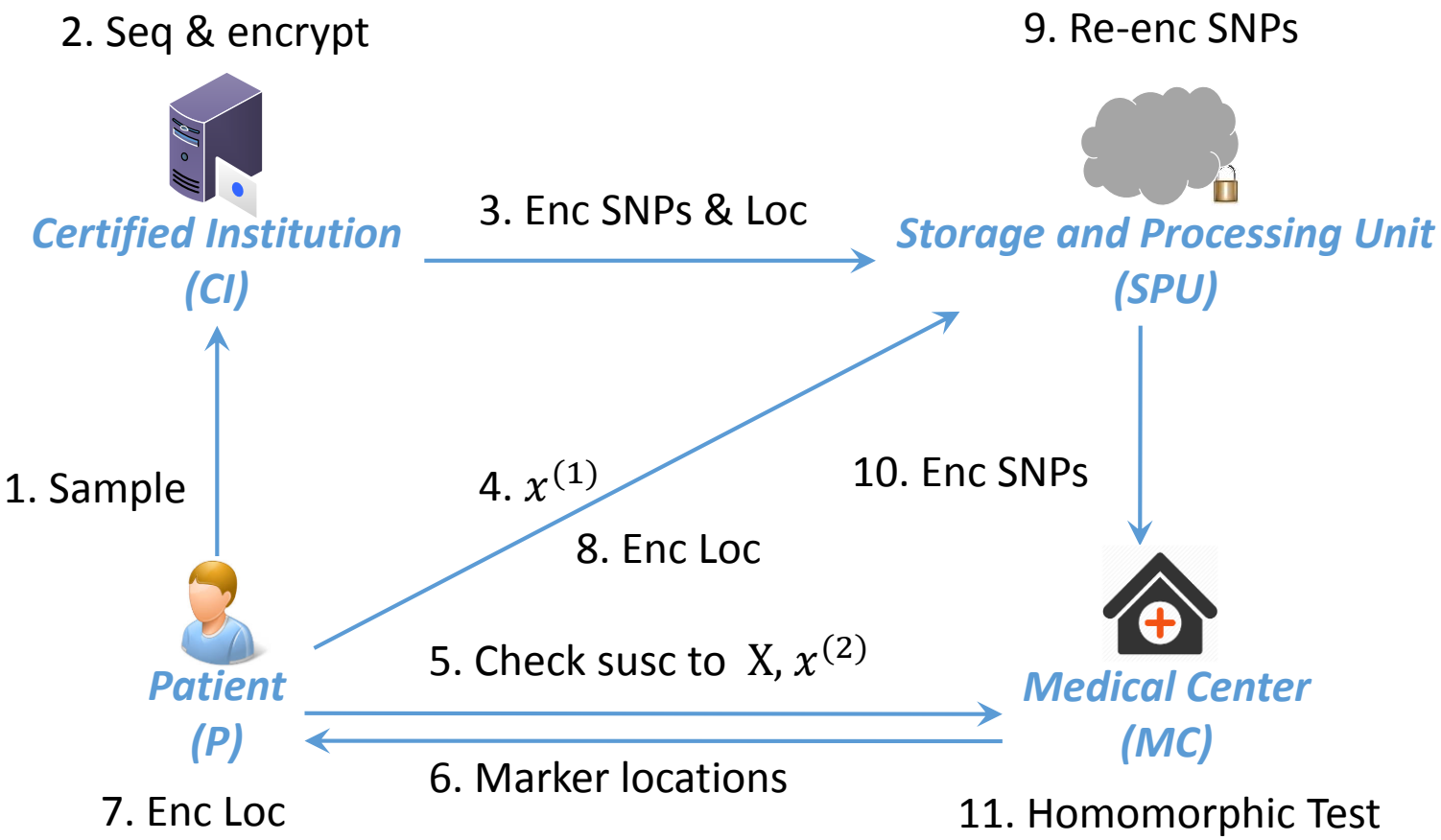

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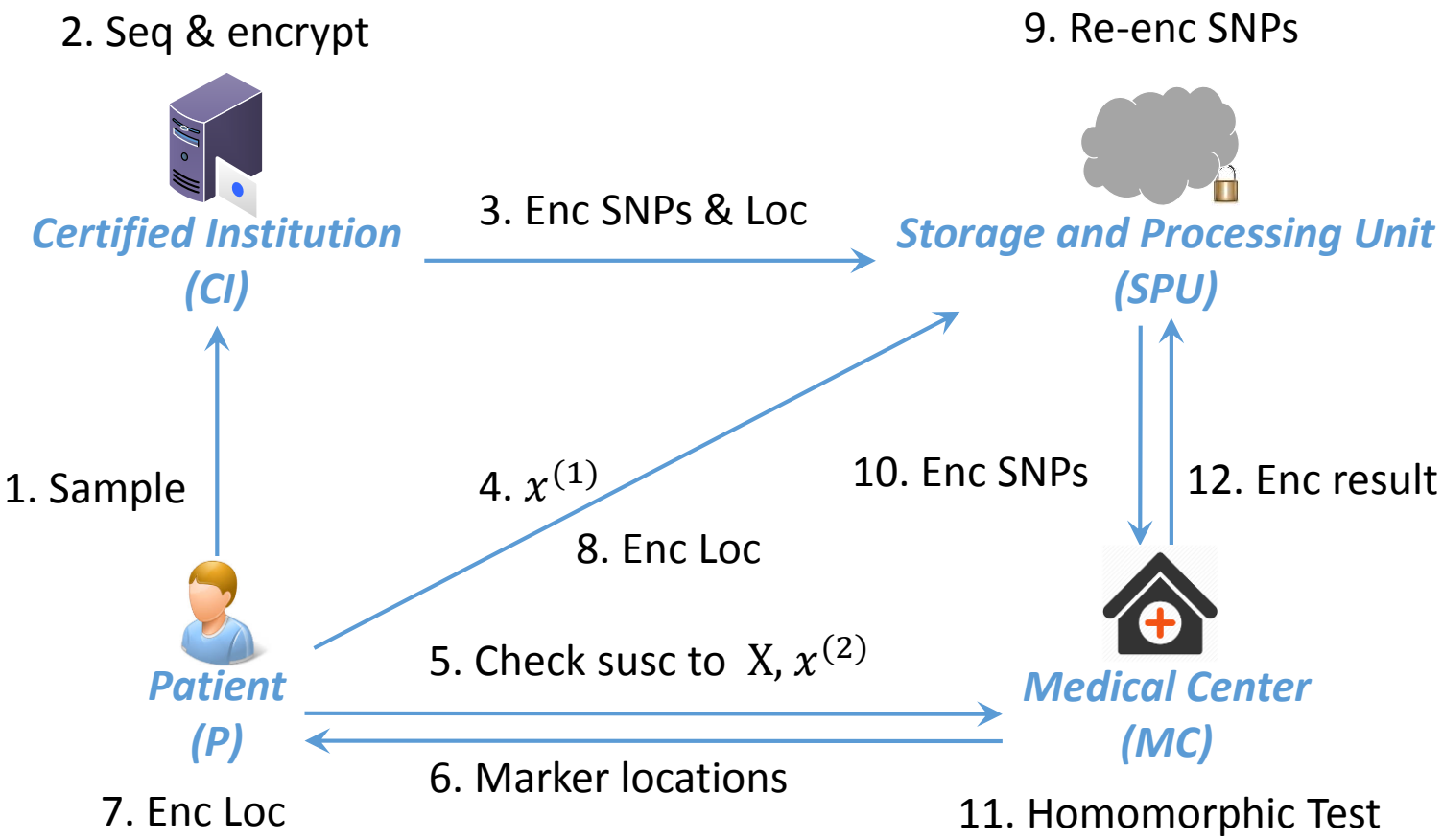

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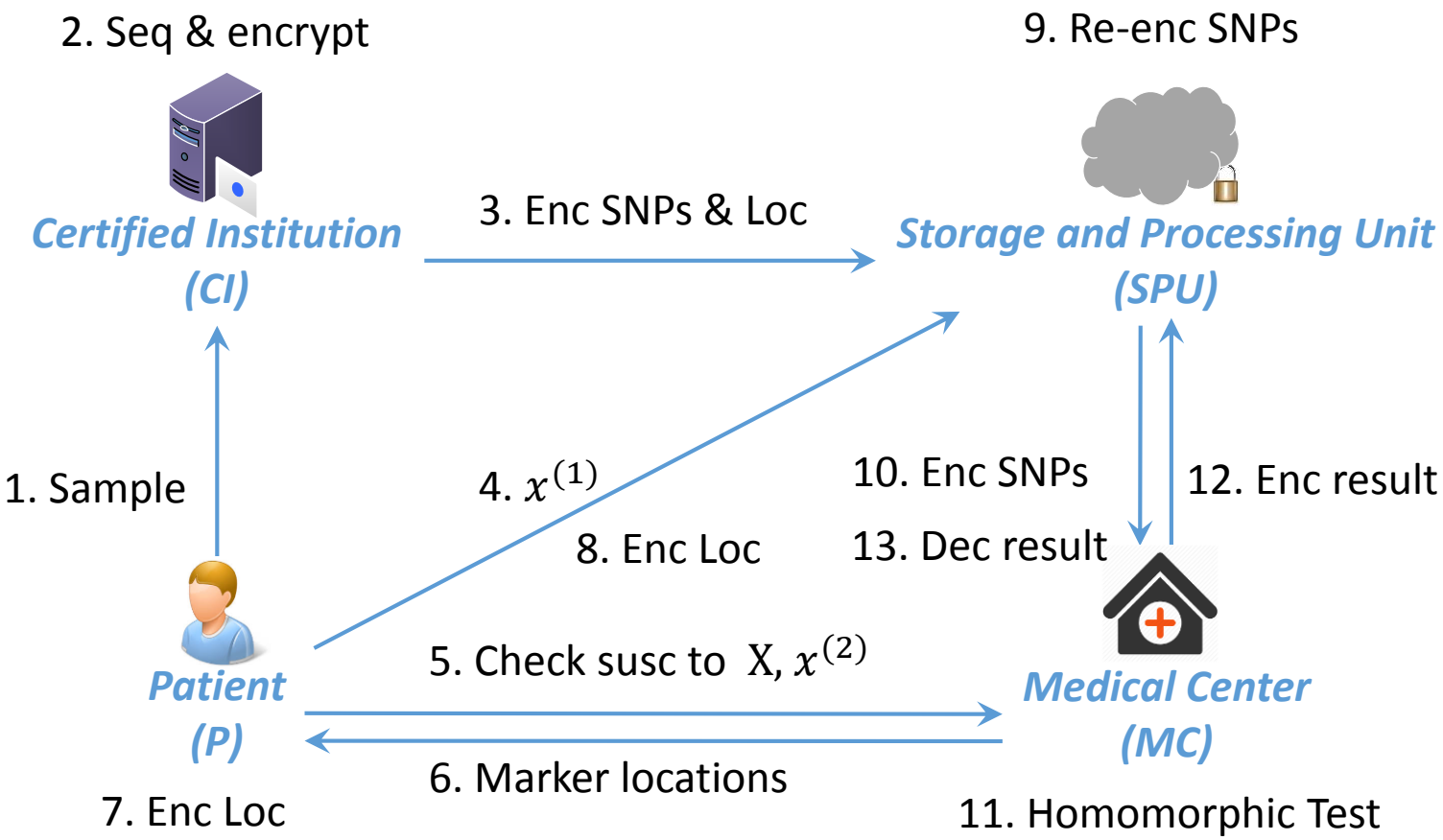


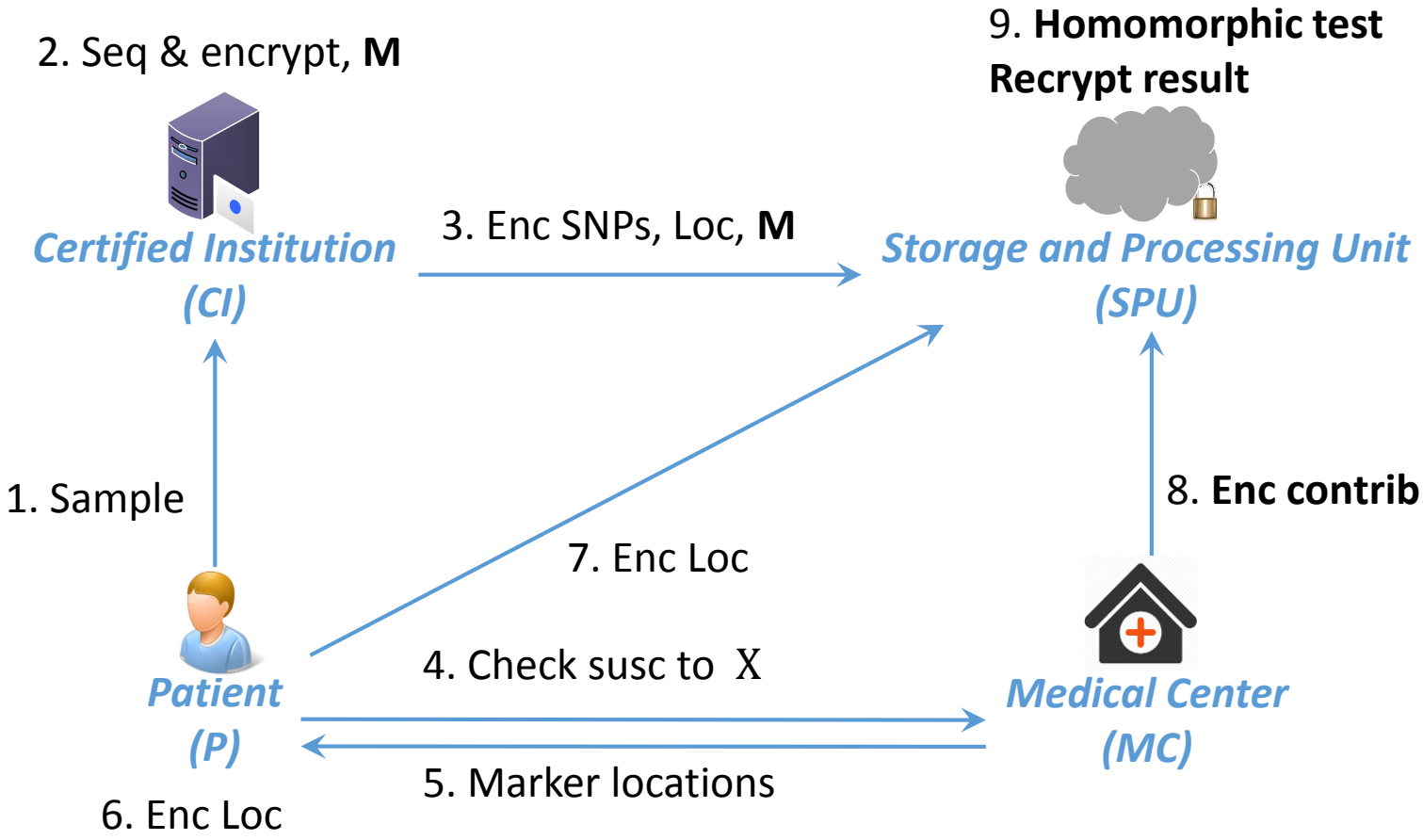
Previous Schemes

Namazi et al. [3]

[3] M. Namazi, J. R. Troncoso-Pastoriza, and F. Pérez-González, "Dynamic Privacy-Preserving Genomic Susceptibility Testing," *in Proceedings of the 4th ACM Workshop on Information Hiding and Multimedia Security*. 2016, IH&MMSec`16, pp. 45-50, ACM.

Privacy-preserving genomic susceptibility testing

- Modified scheme with lattice-based encryptions [NTP16]



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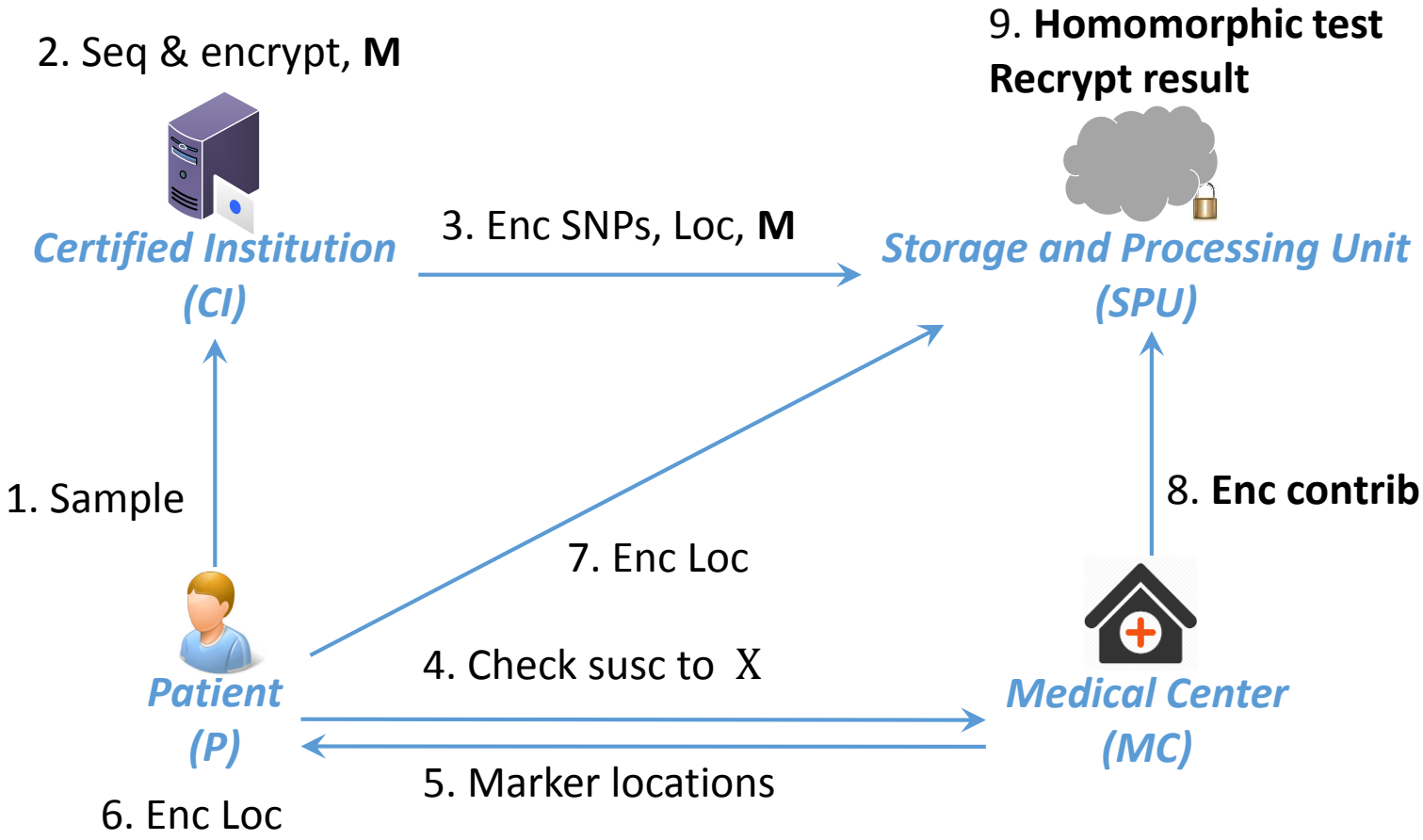

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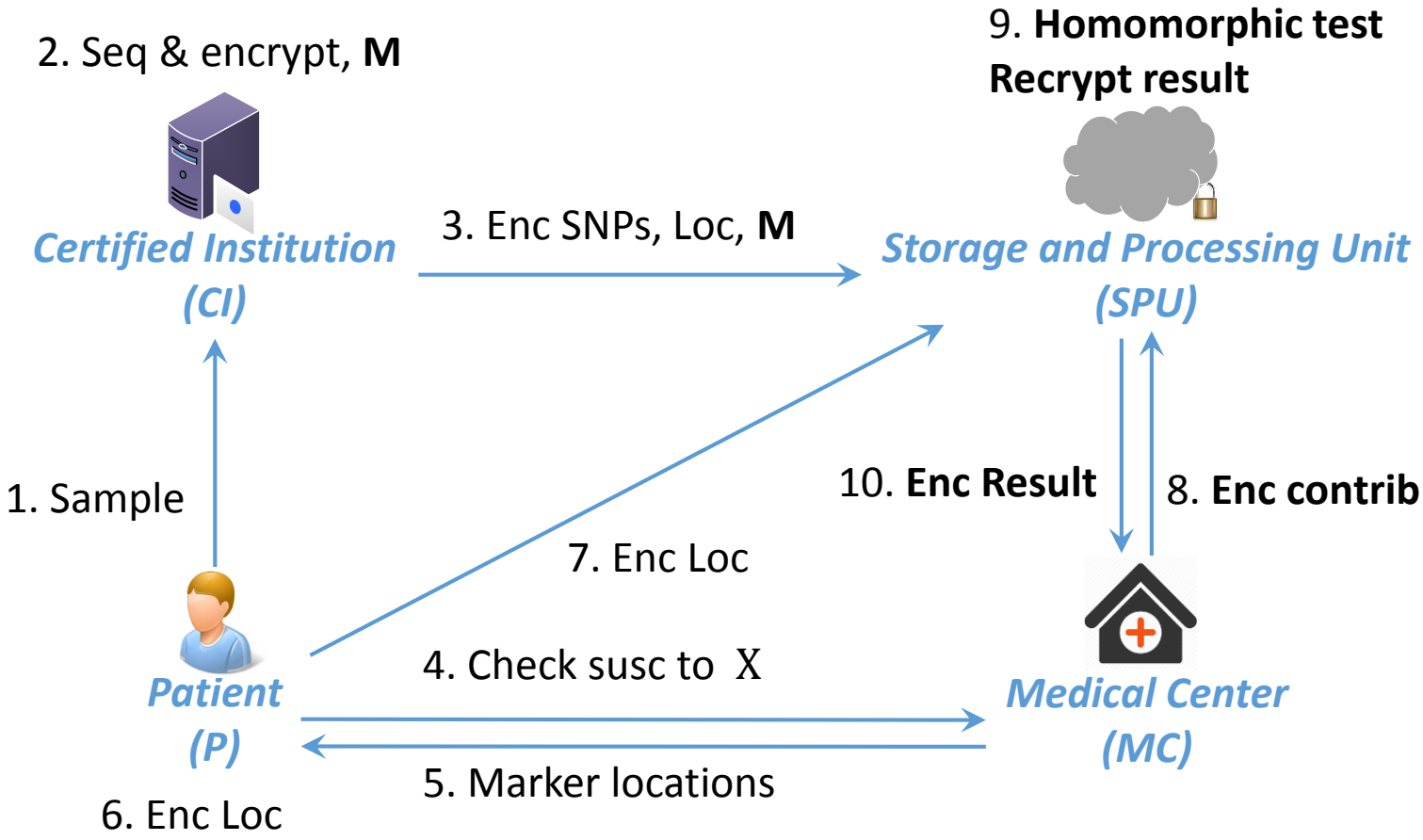

Privacy-preserving genomic susceptibility testing

- Complexity
  - 4M SNPs, 10-marker test
  - Paillier: 2048-bit modulus, Lauter: 4096-dim. lattice

Ayday et al.	CI	SPU		MC	
112 bit security	Encrypt/ SNP	Recrypt	Proxy recrypt	Homomorphic calculation	Paillier decrypt
Time	33.2 ms	304.3 ms	30.3 ms	39.3 ms	30.3 ms
Size	4,1 GB	10.2 kB	1.02 kB	1.02 kB	

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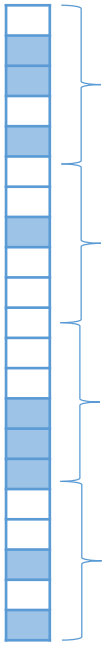
Proposed Scheme

Privacy-preserving genomic susceptibility testing

- One step further: packing
  - Reduce cipher expansion

Patient SNPs



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Privacy-preserving genomic susceptibility testing

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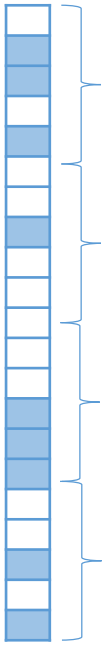

$$SN = \left(\sum_{i \in P} SNP^{p,i} \cdot z^{i} \right)$$

Markers

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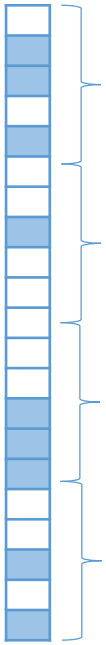
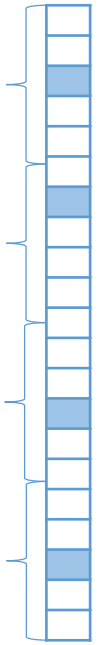


$$SN = \left(\sum_{i \in P} SNP^{p,i} \cdot z^i \right)$$

$$\beta_b = \left[\sum_{i \in \Omega_x} \frac{(-1)^{1-b} c^{X,i} pr_b^{X,i}}{\sum_{i \in \Omega_x} c^{X,i}} \cdot z^i \right]$$

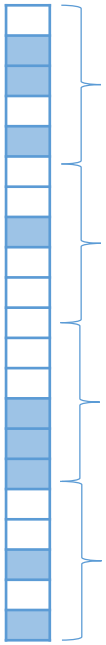
Markers



Privacy-preserving genomic susceptibility testing

- One step further: packing
  - Reduce cipher expansion
  - Needed scalar product, polynomial product available

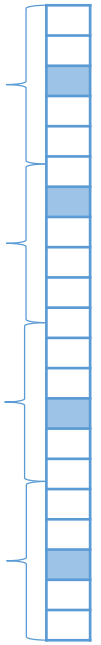
Patient SNPs



$$SN = \left(\sum_{i \in P} SNP^{p,i} \cdot z^i \right)$$

$$\beta_b = \left[\sum_{i \in \Omega_x} \frac{(-1)^{1-b} c^{X,i} pr_b^{X,i}}{\sum_{i \in \Omega_x} c^{X,i}} \cdot z^i \right]$$

Markers



Privacy-preserving genomic susceptibility testing

- One step further: packing
  - Reduce cipher expansion
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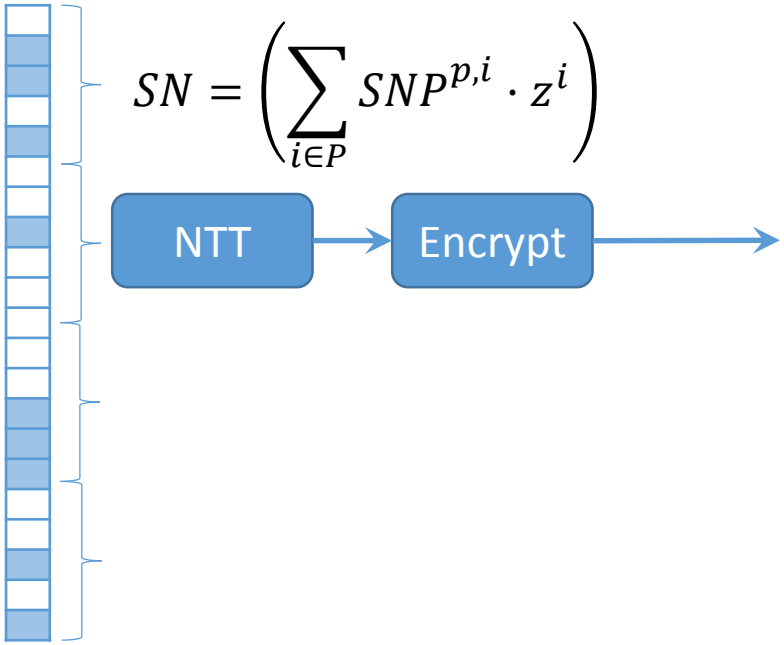

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NTT → Encrypt

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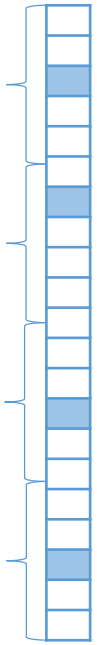

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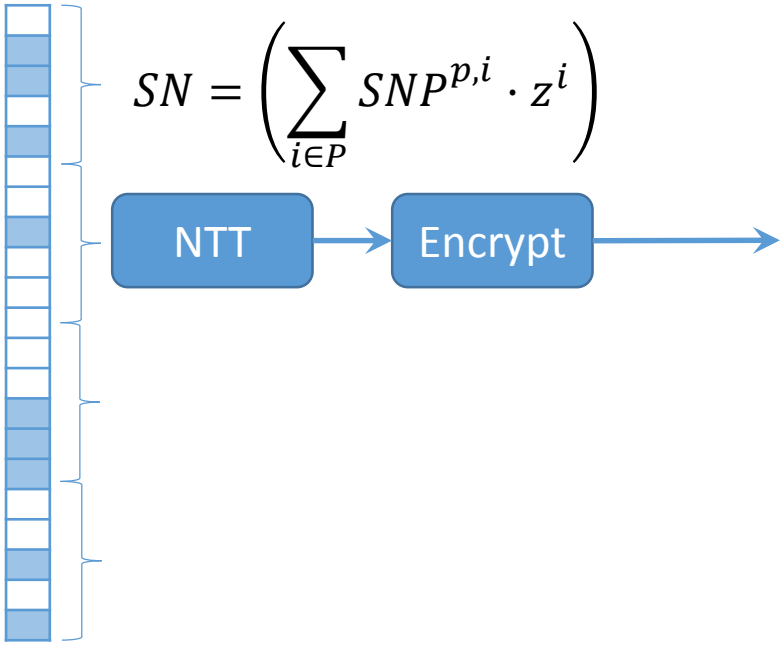

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Patient SNPs

Markers

$$SN_E = \left(NTT\left[\sum_{i\in P} SNP^{p,i}\cdot z^i\right]\right)_E$$

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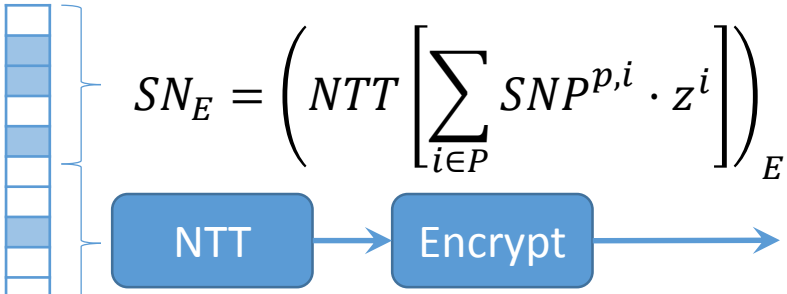

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Markers

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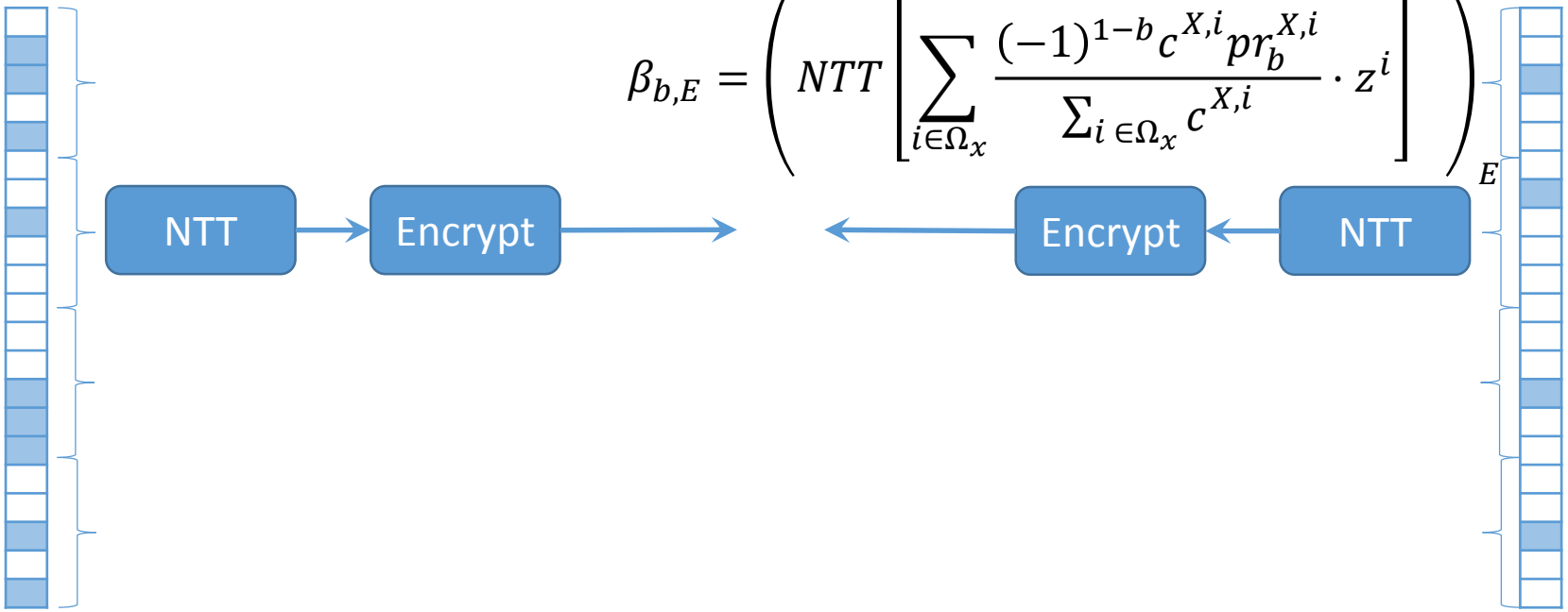

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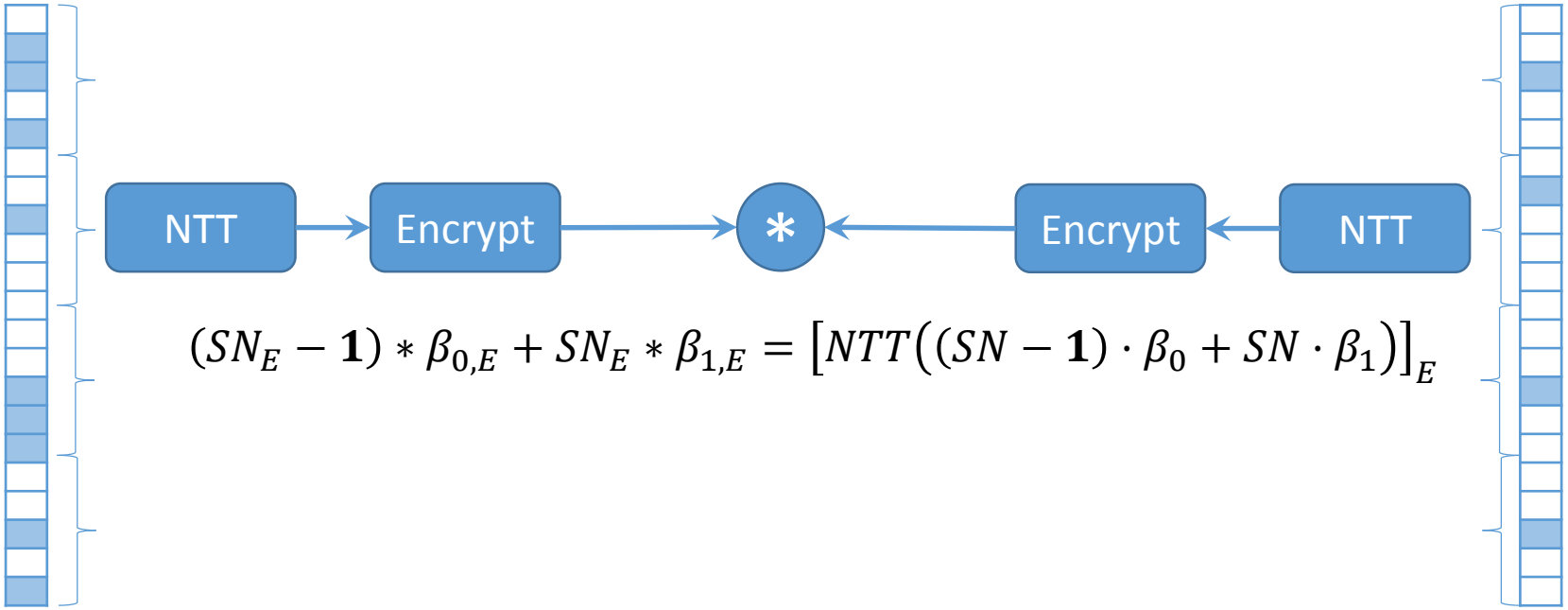

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Privacy-preserving genomic susceptibility testing

- One step further: packing
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$$NTT[\boldsymbol{x}]_k = \sum_{i\in[0,N)} x_i \alpha^{i\cdot k}$$

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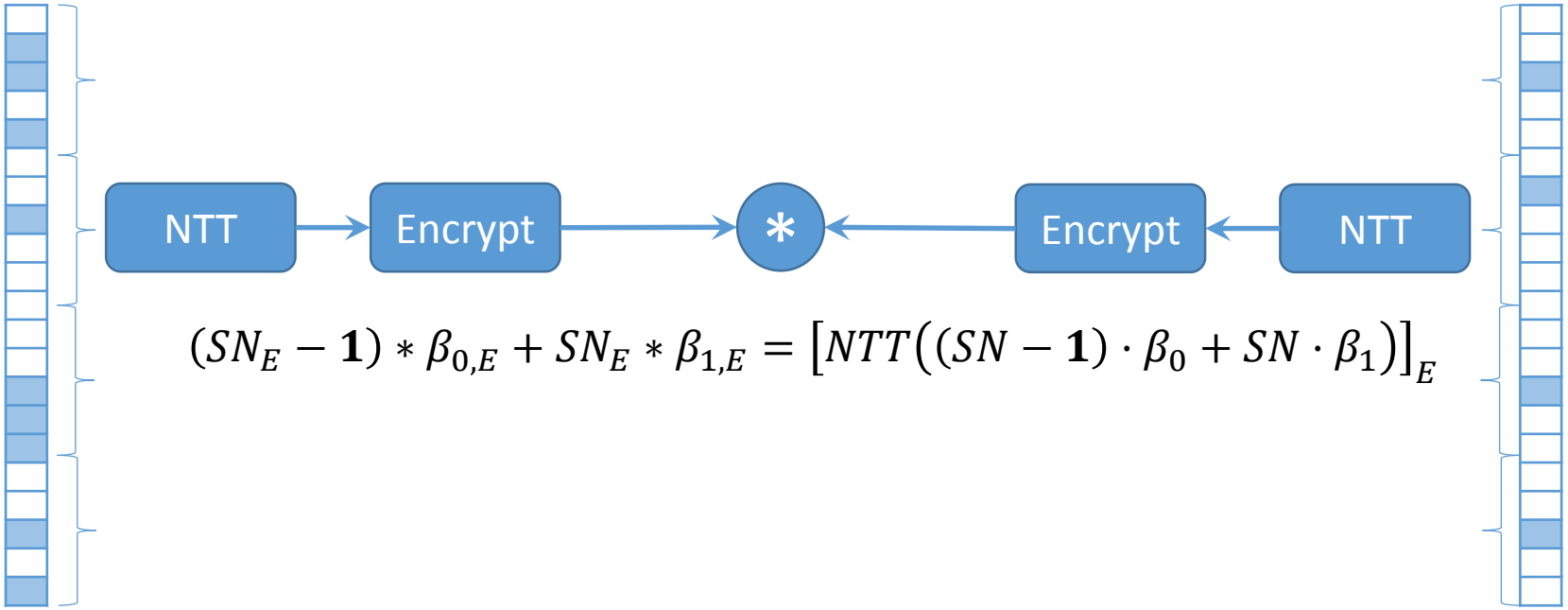

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$$NTT[\boldsymbol{x}] = \begin{pmatrix} 1 & 1 & \dots & 1 \\ 1 & \alpha & \dots & \alpha^{N-1} \\ \vdots & \vdots & \ddots & \vdots \\ 1 & \alpha^{N-1} & \dots & \alpha^{(N-1)(N-1)} \end{pmatrix} \begin{pmatrix} x_0 \\ x_1 \\ \vdots \\ x_{N-1} \end{pmatrix}$$

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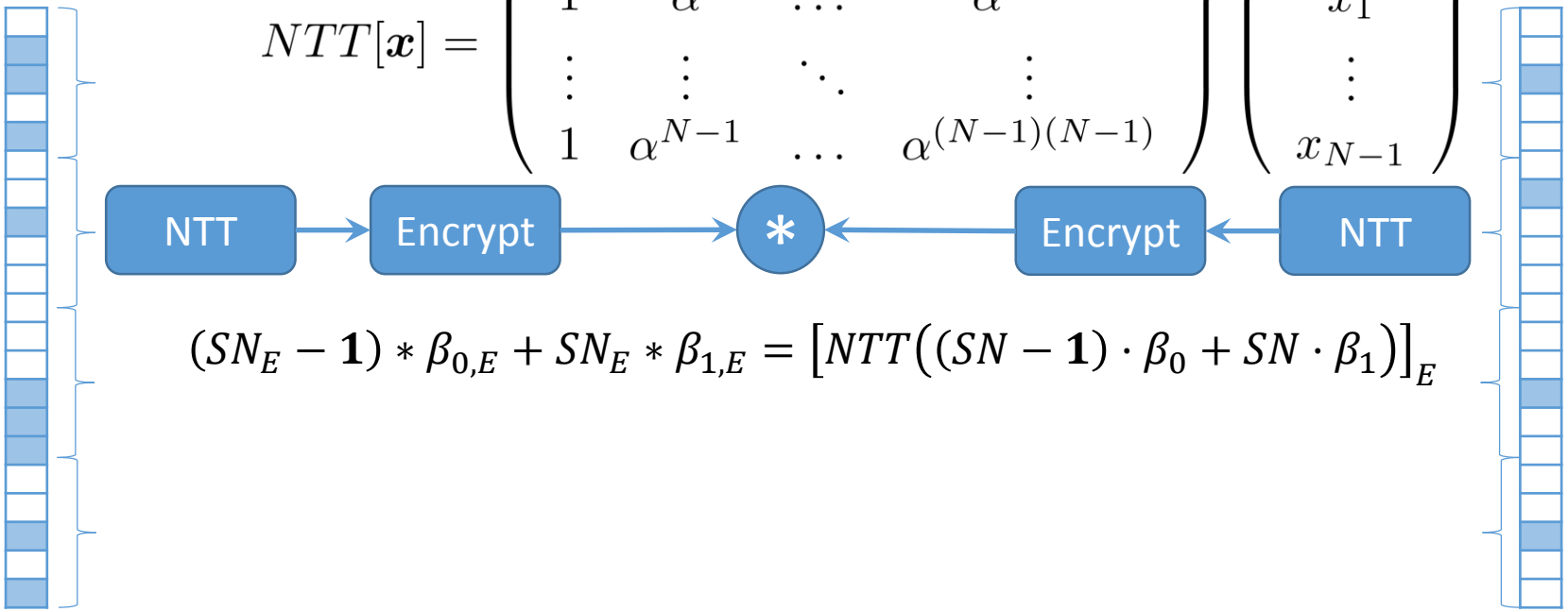

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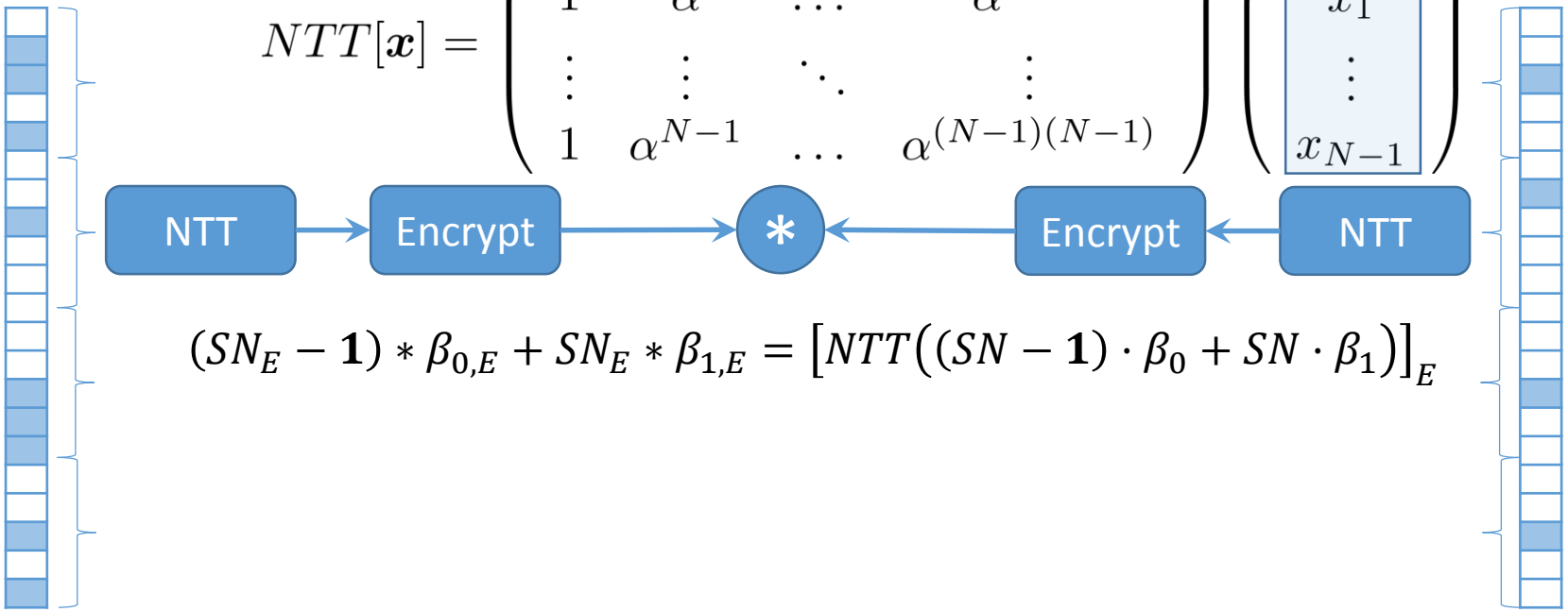

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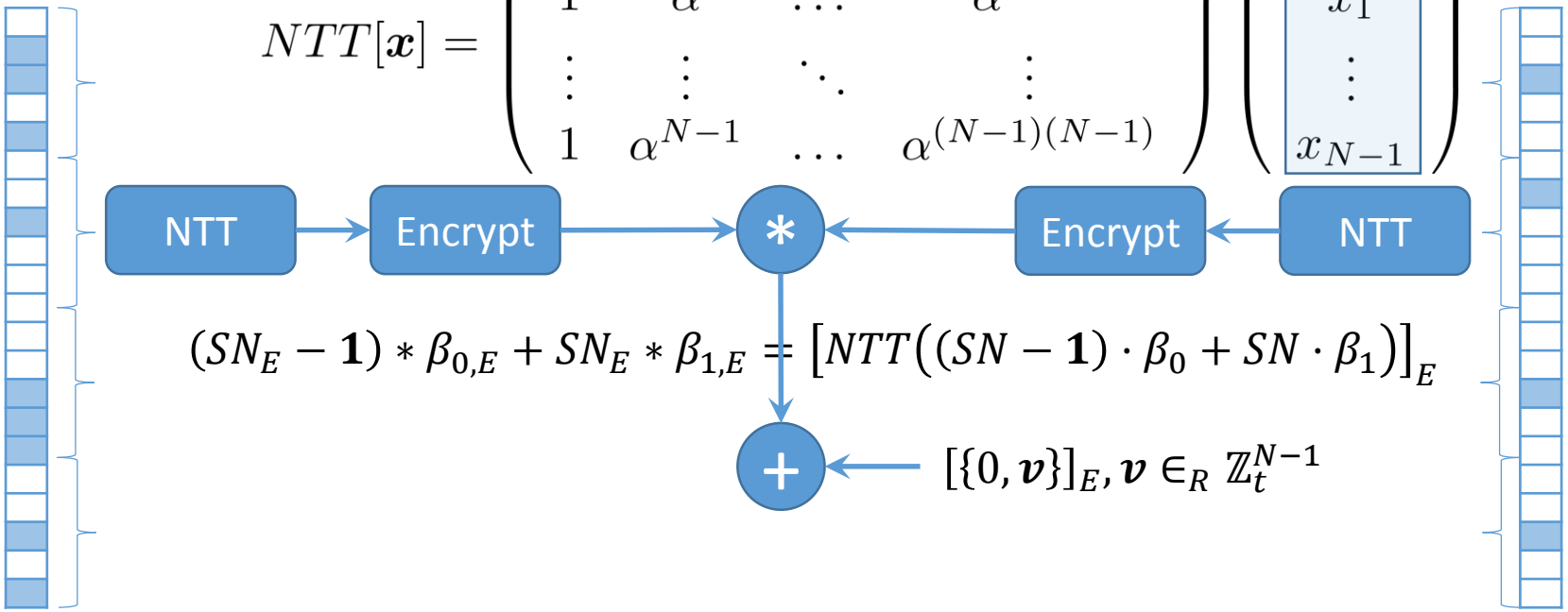

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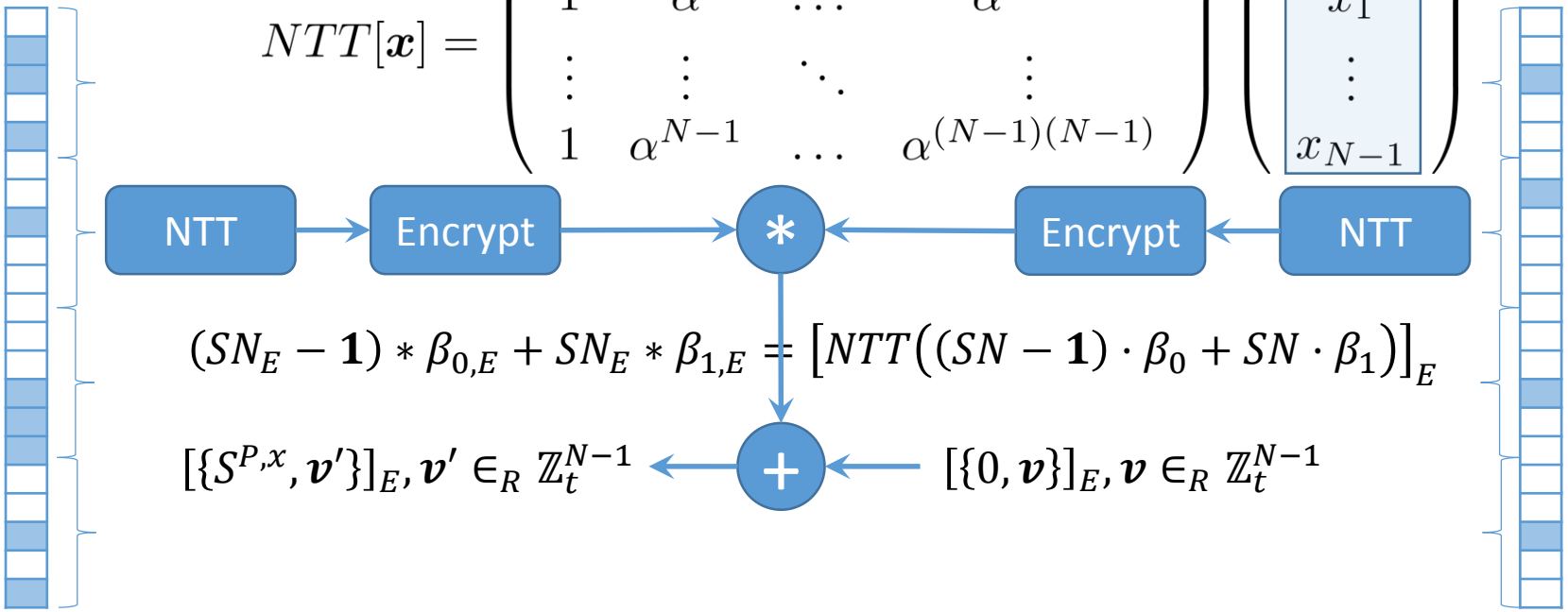

Security and Performance Evaluation

4M SNPs, 10-marker test

Paillier: 2048-bit modulus, 112 bit security
Lauter: 2048-dim. lattice, 127 bit-security (δ = 1.005)

Ayday et al.	**CI**	**SPU**		**MC**	
112 bit security	Encrypt/ SNP	Recrypt	Proxy recrypt	Homomorphic calculation	Paillier decrypt
Time	33.2 ms	304.3 ms	30.3 ms	39.3 ms	30.3 ms
Size	4.1 GB	10.2 kB	1.02 kB	1.02 kB	
Namazi et al.	**CI**	**SPU**		**MC**	
127 bit security	Encrypt/ SNP	Homomorphic calculation	Reliniariz	Enc params	Decrypt
Time	0.22 ms	1.08 ms	1.1 ms	4.5 ms	0.46 ms
Size	131.1 GB		32.8 kB	655 kB	
Proposed	**CI**	**SPU**		**MC**	
127 bit security	Encrypt/ SNP	Homomorphic calculation	Reliniariz	Enc params	Decrypt
Time	0.00011 ms	0.05 – 1.08 ms	1.1 ms	0.22 – 4.5 ms	0.46 ms
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364 bit security	Encrypt/ SNP	Homomorphic calculation	Reliniariz	Enc params	Decrypt
Time	0.45 ms	2.17 ms	2.32 ms	9.1 ms	0.96 ms
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Conclusions

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- An efficient protocol to deal with encrypted genomic susceptibility tests is proposed
- We introduce some optimizations:
  - A reasonable choice of the cryptosystem parameters (for both efficiency and security)
  - A transformed input packing strategy
  - Avoiding costly unpacking/repacking operations
- It moves the bulk of the computation to the SPU
- It outperforms previous solutions in both computational cost, bandwidth and storage

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Research Center for
Information & Communication Technologies

Thanks for your attention!