# Improving HIV treatment choice with multi-party computation

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- Treating HIV is not straight-forward: multiple possible treatments, many different viruses
- Virus mutates as it replicates. Bad treatment leads to more replication, which means:
  - Treatment failure
  - Accumulation of drug resistances
  - Faster progression to AIDS
- Even with optimal treatment, virus will eventually mutate

Doctors take decisions based on

- Yearly published guidelines based on current research
- Knowledge
- Experience

Every time a patient needs new treatment, they get feedback on the results of the old treatment.

Can we use this data?

Two problems:

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- 2. Even if there was a database somewhere, patient's HIV genotype is privacy-sensitive

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Solution: multi-party computation!

## Multi-party computation

Given a patient's HIV genotype, per treatment: what was the average time to failure for patients with similar HIV virus?



## **Multi-party computation**

## Secret sharing

To compute on other parties' data, we need a way to distribute secret data.  $\rightarrow$  Secret sharing



## Shamir's secret sharing scheme

Suppose we have a secret value s in some finite field  $\mathbb{F}$ , and we want to disperse it into n shares.

Fix an integer 0 < t < n, the *threshold*.

Pick a uniformly random secret polynomial  $f = s + c_1 X + c_2 X^2 + \cdots + c_t X^t$  with  $c_1, \ldots, c_t \in \mathbb{F}$ .



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 $s_i := f(\alpha_i)$  is the *i*-th share.

#### Theorem (Lagrange interpolation)

Given  $y_1, \ldots, y_{t+1} \in \mathbb{F}$  there is a unique polynomial  $h = a_0 + a_1 X + \cdots + a_t X^t$  with  $h(\alpha_i) = y_i$  for  $i = 1, \ldots, t+1$ .  $s_2$   $s_1$  $s_4$ 

#### Reconstructing the secret

#### Theorem (Lagrange interpolation)

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So: given t + 1 shares we can reconstruct f.

Privacy

Given  $\leq t$  shares, we get no information about the secret *s*.



## **MPC: Step one**



Every party shares their secret inputs to the computation. Each party has one share of every secret value.



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## Adding shares



## Adding shares



Each party can calculate a share of s + s'.

For  $c \in \mathbb{F}$  a publicly known scalar, parties can also calculate a share of  $c \cdot s$  by multiplying their own share.

So, now we can evaluate linear forms  $\mathbb{F}^n \to \mathbb{F}$ , e.g.

$$(x_1,\ldots,x_n)\mapsto a_1x_1+\cdots+a_nx_n$$

for constants  $a_1, \ldots, a_n \in \mathbb{F}$ .

What about arbitrary functions  $\mathbb{F}^n \to \mathbb{F}$ ? We need multiplication.

## Multiplication

Suppose we have secrets s, s' corresponding to polynomials f, g. deg  $fg = (\deg f) \cdot (\deg g)$ . Hence we need 2t + 1 shares to reconstruct ss'.



We can only do this a limited number of times before we run out of shares.

Suppose we have secrets s, s' corresponding to polynomials f, g. Set  $x_i := s_i \cdot s'_i$ . Party *i* can calculate this value. Now they secret-share  $x_i$  with each other party. Write  $[x_i]_j$  for the share of party *j* of the value  $x_i$ .

Note that the function mapping  $L : (s_1, s_2, \dots, s_n) \mapsto s$  is a linear form!

So every party can calculate  $L([x_1]_i, \ldots, [x_n]_i) = [s \cdot s]_i$ .

## **Evaluating arbitrary functions**

We can represent any computable function  $\mathbb{F}^m \to \mathbb{F}^k$  as a circuit of linear gates and multiplication gates.



E.g. a Boolean AND-gate is just multiplication  $b \cdot b'$  if inputs are b, b' encoded as  $\{0, 1\}$  in  $\mathbb{F}$ .

In general: circuit can be very big!

Assuming we have the desired function to be computed f as a circuit:

- 1. Every party secret-shares their input
- 2. Parties locally process linear gates, and communicate for every multiplication gate
- 3. Every party sends their shares of the output values to the parties who should receive the output, who then reconstruct the value.

Note: we assume parties strictly follow the protocol.

We have seen the basics of how MPC works.

Linear operations: cheap. Multiplications require interaction, which is the dominating cost.

Big circuits  $\rightarrow$  slow computation.

## **HIV** treatment

Given a patient's HIV genotype, per treatment: what was the average time to failure for patients with similar HIV virus?

• Every doctor has a list of

(HIV genotype, time to failure). We need to secret share each of those values.

- HIV genotype gets encoded as a vector  $\mathbf{m} \in \mathbb{F}^{\ell}$ .
- Want to compute

$$\frac{\sum_{i=1}^{N} \mathbf{1}_{\mathbf{m}_i \approx \mathbf{m}} t_i}{\sum_{i=1}^{N} \mathbf{1}_{\mathbf{m}_i \approx \mathbf{m}}}$$

• Need to check against *every* prior patient: computation scales linearly in  $\ell$ , N

Doctor has 5 minutes per patient to make a treatment decision. Work-in-progress: can we make computation fast enough?

- Can we do MPC over finite rings instead of fields, e.g.  $\mathbb{Z}/2^m\mathbb{Z}$ ?
- How can we do often used functionalities efficiently, e.g. integer division?
- Can we do machine learning tasks over MPC, e.g. decision trees and random forests?