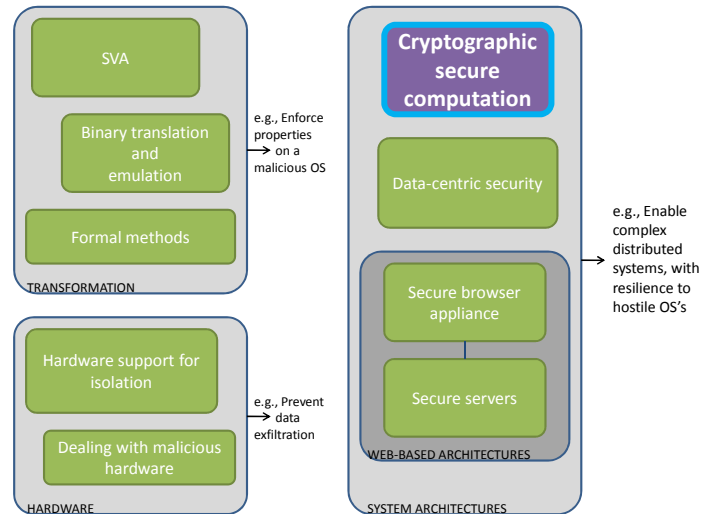


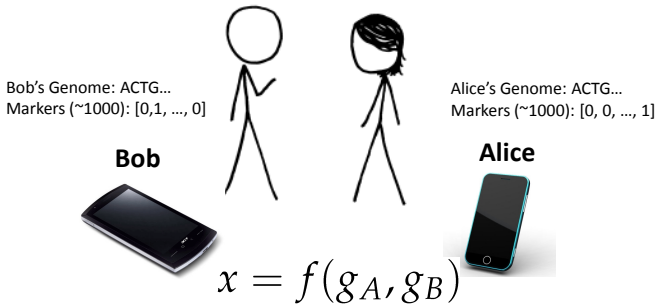
# Practical Cryptographic Secure Computation

DHOSA MURI  
Pls Meeting  
Berkeley, CA  
28 April 2011

David Evans  
University of Virginia  
<http://www.cs.virginia.edu/evans>  
<http://www.MightBeEvil.com>



## Secure Two-Party Computation



Can Alice and Bob compute a function of their private data, without exposing anything about their data besides the result?

## Secure Function Evaluation

**Alice (circuit generator)** **Bob (circuit evaluator)**  
Agree on  
Picks  $a \in \{0,1\}^s$   $f(a,b) \rightarrow x$  Picks  $b \in \{0,1\}^t$

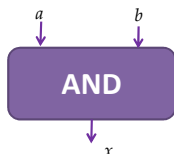
Garbled Circuit Protocol

Outputs  $x = f(a,b)$   
without revealing  $a$   
to Bob or  $b$  to Alice.

Andrew Yao, 1982/1986

## Yao's Garbled Circuits

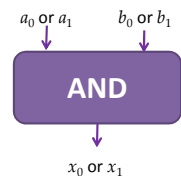
Inputs		Output
$a$	$b$	$x$
0	0	0
0	1	0
1	0	0
1	1	1



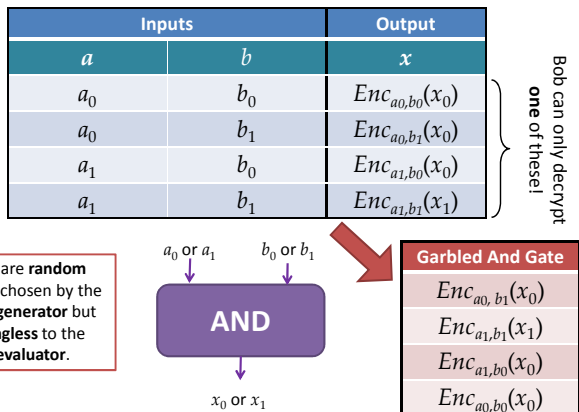
## Computing with Meaningless Values?

Inputs		Output
$a$	$b$	$x$
$a_0$	$b_0$	$x_0$
$a_0$	$b_1$	$x_0$
$a_1$	$b_0$	$x_0$
$a_1$	$b_1$	$x_1$

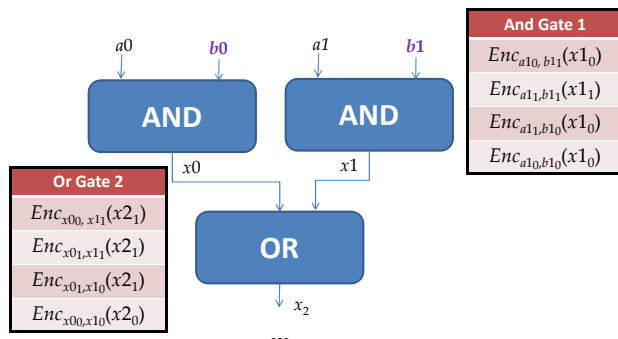
$a_i, b_i, x_i$  are random values, chosen by the circuit generator but meaningless to the circuit evaluator.



## Computing with Garbled Tables

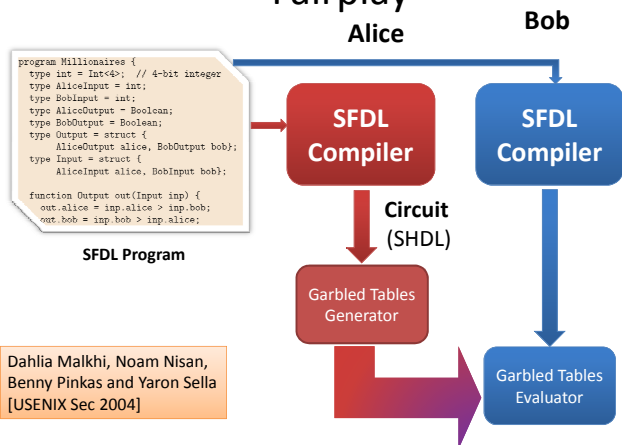


## Chaining Garbled Circuits

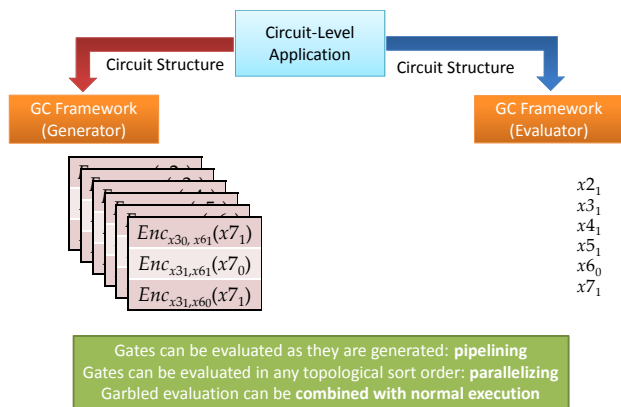


We can do *any* computation privately this way!

## Fairplay

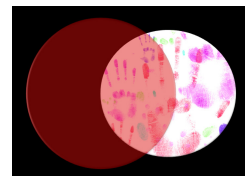


## Our Approach: Faster Garbled Circuits



## Private Set Intersection

- Do Alice and Bob have any contacts in common?
- Two countries want to compare their miscreant lists
- Identify common medical records across hospitals
- Two companies want to do joint marketing to common customers



Privacy-Preserving Biometric Matching

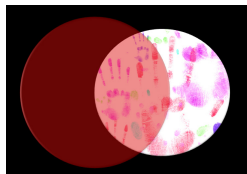
Private Personal Genomics



### Applications

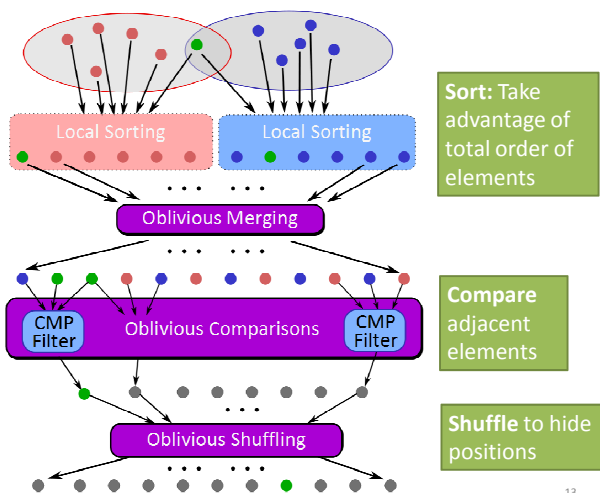


Private AES Encryption



Private Set Intersection

Sort-Compare-Shuffle

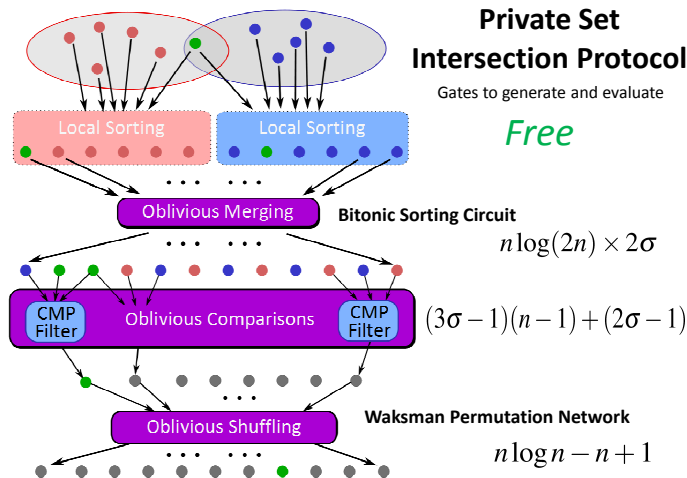


13

### Private Set Intersection Protocol

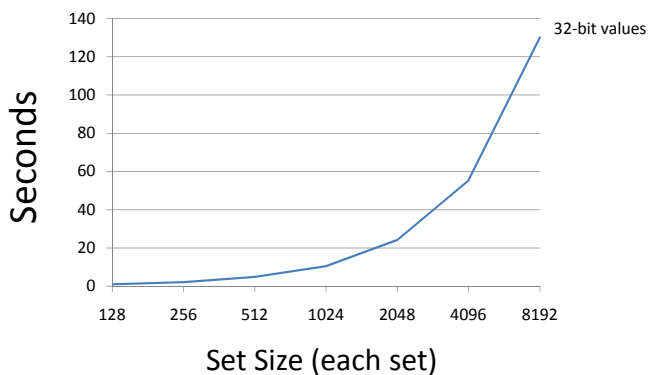
Gates to generate and evaluate

Free



14

### Private Set Intersection Results



15

### Some Other Results

	Problem	Best Previous Result	Our Result	Speedup
USENIX Security 2011	Hamming Distance (Face Recognition) – 900-bit vectors	213s [SCIFI, 2010]	<b>0.051s</b>	<b>4176</b>
	Levenshtein Distance (genome, text comparison) – two 200-character inputs	534s [Jha+, 2008]	<b>18.4s</b>	<b>29</b>
	Smith-Waterman (genome alignment) – two 60-nucleotide sequences	[Not Implementable]	<b>447s</b>	-
	AES Encryption	3.3s [Henecka, 2010]	<b>0.2s</b>	<b>16.5</b>
NDSS 2011	Fingerprint Matching (1024-entry database, 640x8bit vectors)	~83s [Barni, 2010]	<b>18s</b>	<b>4.6</b>

Scalable: 1 Billion gates evaluated at ~100,000 gates/second on laptop

16



#### Collaborators

**Yan Huang** (UVa PhD Student),  
 Yikan Chen (UVa PhD Student),  
 Samee Zahur (UVa MS Student),  
 Peter Chapman (UVa BACS Student)  
**Jonathan Katz** (University of Maryland)  
 Aaron Mackey (UVa Public Health Genomics)

**David Evans**

evans@cs.virginia.edu  
<http://www.cs.virginia.edu/evans>

17