



DNA SECRET SHARING

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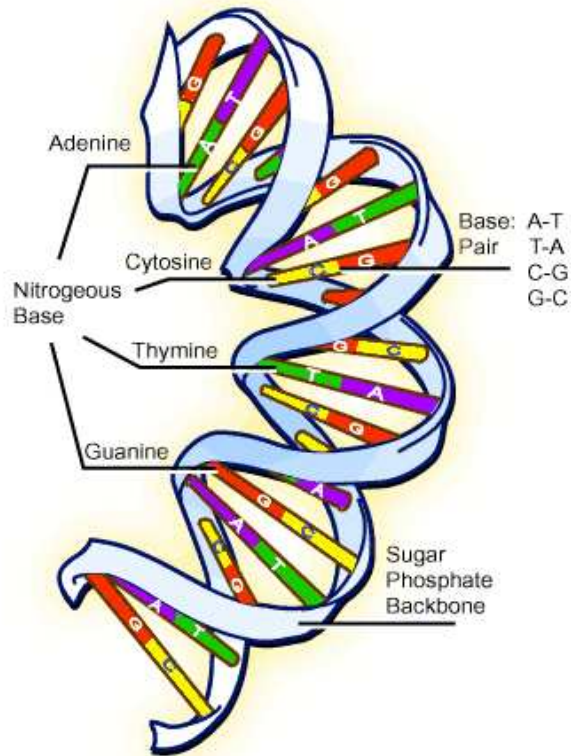
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Aim of Our Work

Our aim is to distribute a **secret binary string** **using DNA computing** to a set $\{P_1, P_2, \dots, P_n\}$ of n participants in such a way that certain designated set of participants can reveal the secret by pulling their shares, but no forbidden set of participants has **any information** about the secret binary string. In short, **our aim is to sharing a secret using DNA.**

Why DNA for Secret Sharing?

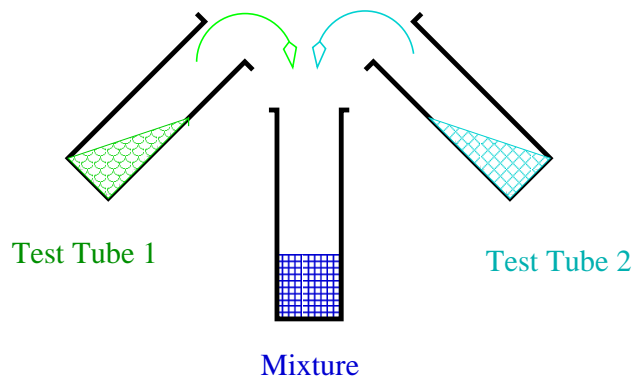


- The very small size,
- The huge storage capacity,
- Easy to carry or hide,
- Made up of A, T, G, C,
- Huge parallel computing,
- Stable as a DNA double strand,
- High longevity,
- Easy to get synthesized DNA

DNA encoding of binary strings

- a binary string can be represented as a set of integers that corresponds the positions where the bits are 1 from left to right.
- 1011 can be represented as a set $\{1, 3, 4\}$,
- each integer i can be represented $ds_i = \updownarrow S_0(GAATTGC^5)^i GAATTC S_1$, where $\updownarrow GAATTC$ is the restriction site for EcoRI and S_0 and S_1 be suitable 20 to 30 base pair long DNA strand not containing $\updownarrow GAATTC$.
- if $\alpha = 1011$, then the DNA double strand representation is $T[\alpha] = \{ds_1, ds_3, ds_4\}$.

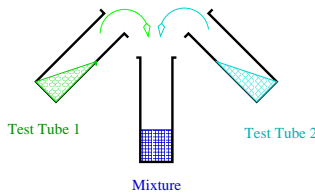
Mixing operation



- Take the content of two test tubes.
- Mixing can be done by dehydrating the tube contents (if not already in solution) and then combining the fluids together into a new tube, by pouring and pumping.

Bio-mathematical operations

- Boolean “*or*” operation between two binary strings
- if $\alpha = 1011$ and $\beta = 1001$,
- the binary “*or*” of two strings will be 1011.
- $T[\alpha]$ ($T[\beta]$) the test tube corresponding to the binary string α (β).
- pore the contents of the two test tubes to get binary “*or*”.



Example of DNA Secret Sharing :

- Consider the secret sharing scheme on $\mathcal{P} = \{1, 2, 3, 4\}$ of 4 participants, where $\Gamma_0 = \{\{1, 2\}, \{2, 3\}, \{2, 4\}, \{1, 4\}\}$.
- $\Gamma_{Qual} = \{Y \subseteq \mathcal{P} : X \subseteq Y \text{ for some } X \in \Gamma_0\}$ and $\Gamma_{Forb} = 2^{\mathcal{P}} \setminus \Gamma_{Qual}$.
- let the secret binary string be $x = x_1x_2x_3 = 011$.
- Assume that the DNA encoding, the mixing process are public.

Share Distribution :

- The dealer chooses two Boolean matrices

$$G_0 = \begin{bmatrix} 0 & 1 & 0 & 1 \\ 0 & 1 & 0 & 1 \\ 0 & 1 & 0 & 0 \\ 0 & 0 & 0 & 1 \end{bmatrix} \text{ and } G_1 = \begin{bmatrix} 1 & 0 & 1 & 0 \\ 0 & 1 & 1 & 0 \\ 1 & 0 & 0 & 0 \\ 0 & 0 & 0 & 1 \end{bmatrix} .$$

- Since $x_1 = 0$, the dealer considers the matrix G_0 and apply a random permutation to the columns of G_0 and produces a matrix M_1 .
- Similarly, for x_2 and x_3 on G_1 to produce matrices M_2 and M_3 .
- Let $M = M_1 || M_2 || M_3$.

Share Distribution :

- first row of M is $\alpha_1 = 010110101010$. Similarly $\alpha_2 = 010101100110$, $\alpha_3 = 010010001000$ and $\alpha_4 = 000100010001$.
- dealer converts them to DNA representations to get $T[\alpha_1] = \{ds_2, ds_4, ds_5, ds_7, ds_9, ds_{11}\}$, $T[\alpha_2] = \{ds_2, ds_4, ds_6, ds_7, ds_{10}, ds_{11}\}$, $T[\alpha_3] = \{ds_2, ds_4, ds_5, ds_9\}$, $T[\alpha_4] = \{ds_4, ds_8, ds_{12}\}$,
- $T[\alpha_i]$ is given to the participants P_i . Also the values $m = 4$ and $k = 3$ are given to each participants even through an insecure channel.

Dccryption by the Qualified participants

- Let P_1, P_2 come together.
- They use mixing procedure with test tubes $T[\alpha_1]$ and $T[\alpha_2]$ to get $T[\alpha_1] \cup T[\alpha_2] = \{ds_2, ds_4, ds_5, ds_6, ds_7, ds_9, ds_{10}, ds_{11}\}$.
- Execute automated DNA sequencing method to read the DNA double strands.
- With the knowledge of decoding the DNA representation to the binary string, the values of $k = 3$ and $m = 4$, the participants P_1 and P_2 can convert the DNA representation to the binary string $y = 010111101110$.

Dccryption by the Qualified participants

- Since, the value of m is known to the participants, P_1 and P_2 can break y as $y = (0101)(1110)(1110)$.
- Next they will find the value of w as 3 and then they will compute $z = 011$, as $BW(0101) < 3$, $BW(1110) = 3$. Thus P_1 and P_2 can recover the secret 011.

Forbidden set of participants

- Let $Y = \{P_3, P_4\}$ come together.
- They use mixing procedure with test tubes $T[\alpha_3]$ and $T[\alpha_4]$ to get $T[\alpha_1] \cup T[\alpha_2] = \{ds_2, ds_4, ds_5, ds_8, ds_9, ds_{12}\}$.
- they will convert the DNA representation to the binary string $y = (0101)(1001)(1001)$.
- Thus looking at those it is not possible to predict whether they correspond to 0 or 1.

Construction of Generating Matrices

Let us consider the following two associated system of linear equations over the binary field \mathbb{Z}_2 ,

$$(1) \quad A\mathbf{x} = \mathbf{0}$$

$$(2) \quad A\mathbf{x} = \mathbf{1}$$

where, A is a $2 \times n$ known Boolean matrix of rank 2; \mathbf{x} is an $n \times 1$ vector of unknowns; $\mathbf{0}$ and $\mathbf{1}$ are $r \times 1$ vectors of 0's and 1's respectively. Let G_0 (G_1) be an $n \times 2^{n-2}$ Boolean matrix whose columns are all possible solutions of the system (1) ((2)).

Generating Matrices

Lemma 1 *Let $(\Gamma'_{Qual}, \Gamma'_{Forb})$ be a strong access structure on a set $\mathcal{D} = \{1, 2, \dots, p\}$ of p participants with $\Gamma'_0 = \{B_i, B_j\}$ where $p \leq n$, $B_i, B_j \subseteq \mathcal{D}$, $|B_i \cup B_j| = p$, $|B_i| \geq 2$ and $|B_j| \geq 2$. Then there exists generating matrices G_0 and G_1 for $(\Gamma'_{Qual}, \Gamma'_{Forb})$.*

Generating Matrices

Lemma 2 *Let G_0^1 and G_1^1 (G_0^2 and G_1^2) denote the generating matrices of a given access structure $(\Gamma_{Qual}^1, \Gamma_{Forb}^1)$ ($(\Gamma_{Qual}^2, \Gamma_{Forb}^2)$) on the set of participants $X_1 = \{i_{1_1}, i_{1_2}, \dots, i_{1_k}\}$ ($X_2 = \{i_{2_1}, i_{2_2}, \dots, i_{2_s}\}$). Then there exist generating matrices G_0 and G_1 for the access structure $(\Gamma_{Qual}^1 \cup \Gamma_{Qual}^2, \Gamma_{Forb}^1 \cap \Gamma_{Forb}^2)$ on the set of participants $X = X_1 \cup X_2$.*

Main Theorem

Theorem 1 *Let $(\Gamma_{Qual}, \Gamma_{Forb})$ be a strong access structure on a set $\mathcal{P} = \{1, 2, \dots, n\}$ of n participants with $\Gamma_0 = \{B_1, B_2, \dots, B_k\}$ where $B_i \subseteq \mathcal{P}, \forall i = 1, 2, \dots, k$. Let σ be a permutation on $\{1, 2, \dots, n\}$. Then there exists generating matrices G_0 and G_1 for the access structure $(\Gamma_{Qual}, \Gamma_{Forb})$ on \mathcal{P} .*



Conclusion

- It is a perfectly secure scheme.
- It has low error rate and it is easy to implement.
- Huge amount of data can be shared secretly.
- the “or” operation can be carried out very efficiently.
- Finally, it is suitable for secret agents and spies.
- The results in DNA-cryptography are very few and demand attention from the researchers.