

# Single user tracing superimposed codes

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*Abstract* — We investigate the zero-error capacity region of  $r$ -out of  $T$  user multiple access OR channel. A family  $\mathcal{F}$  subsets of  $[n] = \{1, \dots, n\}$  is an  $r$ -single-user-tracing superimposed code ( $r$ -SUT) if there exists such a single-user-tracing function  $\phi: 2^{[n]} \mapsto \mathcal{F}$  that for all  $\mathcal{F}' \subseteq \mathcal{F}$  with  $|\mathcal{F}'| \leq r$ ,  $\phi(\cup_{A \in \mathcal{F}'} A) \in \mathcal{F}'$ . In this paper we introduce the concept of these codes and investigate bounds on their rate. We also consider disjointly  $r$ -superimposed codes.

## I. SINGLE-USER-TRACING SUPERIMPOSED CODES

Suppose that  $T$  users share a common channel. To each of them a binary vector of length  $n$  is associated. The  $i^{\text{th}}$  user transmits its vector  $x_i = (x_i^1, \dots, x_i^n)$  ( $i = 1, 2, \dots, T$ ) if it is active, otherwise not. It is assumed that the transmission is bit and block synchronized. The destination of the messages is a single receiver, which gets the bitwise OR vector of the vectors associated to the active users. Moreover, suppose that at most  $r$  users are active simultaneously. In the classical framework of superimposed coding, the receiver has to be able to identify the set of all active users from the output vector  $y$  of the channel. That is, the code must satisfy the property that for all choices of  $x_1, \dots, x_k$  and  $z_1, \dots, z_\ell$  of codewords with  $1 \leq k, \ell \leq r$  and  $\{x_1, \dots, x_k\} \neq \{z_1, \dots, z_\ell\}$ , we have  $\vee_{i=1}^k x_i \neq \vee_{j=1}^\ell z_j$ . The rate of these codes have been studied extensively in e.g., [2, 3, 4].

Here we investigate the case when the receiver has to be able to identify just one user out of at most  $r$  active ones. Clearly, if a code is superimposed in the classical sense then it satisfies this requirement. A practical motivation for our study of  $r$ -SUT families rises from a novel application of combinatorial designs in genomics [1].

As the question is rather of a combinatorial nature, we switch to that terminology. Codewords are characteristic vectors of subsets of the set  $[n] = \{1, \dots, n\}$  with  $n > 0$ .

**Definition I.1** A family  $\mathcal{F} \subseteq 2^{[n]}$  is  $r$ -superimposed if  $\cup_{i=1}^k A_i \neq \cup_{j=1}^\ell B_j$  for any  $\{A_1, \dots, A_k\} \neq \{B_1, \dots, B_\ell\}$ ,  $1 \leq k, \ell \leq r$ ;  $A_1, \dots, A_k, B_1, \dots, B_\ell \in \mathcal{F}$ .

We are interested in  $r$ -single-user-tracing families ( $r$ -SUT), defined as follows.

**Definition I.2** A family  $\mathcal{F}$  is  $r$ -SUT if for all choices of  $\mathcal{F}_1, \dots, \mathcal{F}_k \subseteq \mathcal{F}$  with  $|\mathcal{F}_i| \leq r$ ,  $\cup_{A \in \mathcal{F}_1} A = \cup_{A \in \mathcal{F}_2} A = \dots =$

$\cup_{A \in \mathcal{F}_k} A$  implies  $\cap_{i=1}^k \mathcal{F}_i \neq \emptyset$ . Equivalently, there exists such a single-user-tracing function  $\phi: 2^{[n]} \mapsto \mathcal{F}$  that for all  $\mathcal{F}' \subseteq \mathcal{F}$  with  $|\mathcal{F}'| \leq r$ ,  $\phi(\cup_{A \in \mathcal{F}'} A) \in \mathcal{F}'$ .

For every base set size  $n$  and  $r$ , let  $f(n, r)$  denote the maximum size of an  $r$ -superimposed family, and  $g(n, r)$  denote the maximum size of an  $r$ -SUT family. We give bounds on the rate of  $r$ -SUT families, which is  $R_g(r) = \limsup_{n \rightarrow \infty} \frac{\log g(n, r)}{n}$ .

**Theorem I.3**  $\exists c_1, c_2 > 0$  such that  $\frac{c_1}{r^2} \leq R_g(r) \leq \frac{c_2}{r}$ .

## II. DISJOINTLY $r$ -SUPERIMPOSED CODES

Another important case implicated in the multiple access model of Section I is when the receiver must distinguish only between disjoint sets of active users. The following definition captures this notion.

**Definition II.1**  $\mathcal{F} \subseteq 2^{[n]}$  is disjointly  $r$ -superimposed if  $\cup_{i=1}^k A_i \neq \cup_{j=1}^\ell B_j$  for  $\{A_1, A_2, \dots, A_k\} \cap \{B_1, B_2, \dots, B_\ell\} = \emptyset$ ;  $1 \leq k, \ell \leq r$ ;  $A_1, \dots, A_k, B_1, \dots, B_\ell \in \mathcal{F}$ .

Let  $h(n, r)$  be the maximum size of disjointly  $r$ -superimposed families.

**Lemma II.2** If  $\mathcal{F}$  is  $r$ -superimposed then it is  $r$ -SUT. If  $\mathcal{F}$  is  $r$ -SUT, then it is disjointly  $r$ -superimposed. Hence,  $f(n, r) \leq g(n, r) \leq h(n, r)$ .

Despite the seemingly slight difference between Definitions II.1 and I.1, the extremal properties of disjointly  $r$ -superimposed codes (families) and  $r$ -superimposed ones differ substantially.

**Theorem II.3**  $\exists c_1, c_2 > 0$  such that  $\frac{c_1}{r} \leq R_h(r) \leq \frac{c_2 \log r}{r}$ .

## ACKNOWLEDGMENTS

We would like to thank Noga Alon and László Györfi for fruitful discussions.

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<sup>1</sup>Research supported by NSERC Grant 250391-02.

<sup>2</sup>Research supported in part by OTKA Grants T038198, T046234. This work relates to Department of the Navy Grant N00014-04-1-4034 issued by the Office of Naval Research International Field Office. The United States Government has a royalty-free license throughout the world in all copyrightable material contained herein.