

# Sub-string/Pattern Matching in Sub-linear Time Using a Sparse Fourier Transform Approach

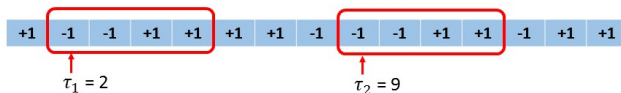
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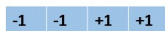


# Problem Statement

Database ( $x$ )



Query ( $y$ )



- **Database/String:**  $\underline{x} = [x[0], x[1], \dots, x[N]]$  (length  $N$ )
- **Query/Substring:**  $\underline{y} = [y[0], y[1], \dots, y[M]]$  (length  $M = N^\mu$ )
- Determine all the  **$L$  locations**  $\underline{\tau} = [\tau_1, \tau_2, \dots, \tau_L]$  with **high probability** where
  - 1 **Exact Matching:**  $\underline{y}$  appears **exactly** in  $\underline{x}$ 
    - $\underline{y} := \underline{x}[\tau : \tau + M - 1]$
  - 2 **Approximate Matching:**  $\underline{y}$  is a **noisy substring** of  $\underline{x}$ 
    - $\underline{y} := \underline{x}[\tau : \tau + M - 1] \odot \underline{b}$
    - $\underline{b}$  is a noise sequence with  $d_H(\underline{y}, \underline{x}[\tau : \tau + M - 1]) \leq K$

# Main Result

## Theorem 1

Assume that a sketch of  $\underline{x}$  of size  $O(\max(M, N/M) \log N)$  can be precomputed and stored. Then for the exact pattern matching and approximate pattern matching (with  $K = \eta M$ ,  $0 \leq \eta \leq 1/6$ ) problems, with the number of matches  $L$  scaling as  $O(N^\lambda)$ , our algorithm has

- a sketching function for  $\underline{y}$  that computes  $O(\max(M, \frac{N}{M}) \log N) = O(N^{\max(\mu, 1-\mu)} \log N)$  *samples*
- a *computational complexity* of
  - $O(\max(N^{1-\mu} \log^2 N, N^{\mu+\lambda} \log N))$  for  $\mu < 0.5$
  - $O(\max(N^\mu \log^2 N, N^{1-\mu+\lambda} \log N))$  for  $\mu > 0.5$
- a decoder that recovers all the  $L$  matching positions with a *failure probability that approaches zero asymptotically*

*Note:* Particularly when  $L = O(1)$  or  $L < \frac{N}{M}$  (i.e.  $\lambda < 1 - \mu$ ) our algorithm has a *sub-linear time* complexity.

# Some Prior Work

## Exact Matching

- [?]: First occurrence of the match (only  $\tau_1$ )
  - Average complexity -  $O(N^{1-\mu} \log N)$  (sublinear)
  - Worst case complexity -  $O(N \log N)$

## Approximate Matching

- [?]: Generalization of [?]
  - Average complexity -  $O(NK/M \log N)$  (sub-linear only when  $K \ll M$ )
- [?]:  $O(N/M^{0.359})$  (sub-linear even when  $K = O(M)$ )
  - Combinatorial in nature

## Sparse Fourier Transform Approach

- [?]: Faster GPS receiver
  - Exploited sparsity in Correlation function  $R_{XY}$
- [?]: Robust Sparse Fourier Transform
  - Sparse Graph code Approach
  - Computational complexity :  $O(N \log N)$

# Motivation

- **Cross-correlation** ( $\underline{r}$ ):

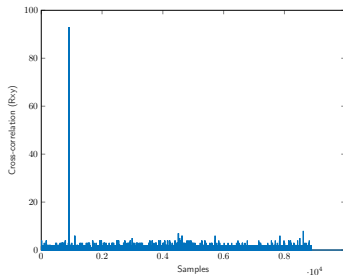
$$r[m] = (\underline{x} * \underline{y})[m] \triangleq \sum_{i=0}^{M-1} x[m+i]y[i], \quad 0 \leq m \leq N-1$$

- **Naive implementation:**  $O(MN) = O(N^{1+\mu})$  (**super-linear** complexity)
- **Fourier Transform Approach:**  $O(N \log N)$  complexity

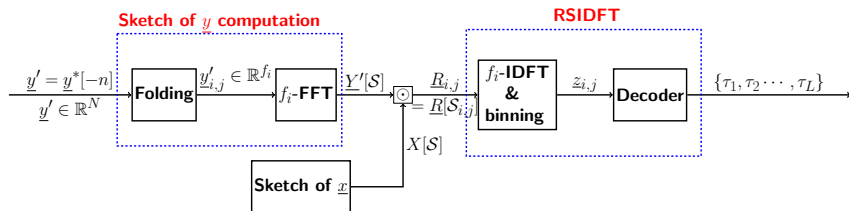
## Key Observation

- $\underline{r}$  is **Sparse** with some noise.

$$r[m] = \begin{cases} M, & \text{if } m \in \mathcal{T} \\ n_m, & m \in [N] - \mathcal{T} \end{cases}$$



# Sparse Fourier Transform Approach



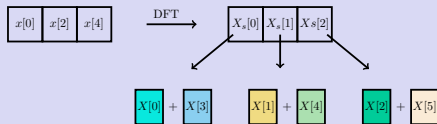
$$\underline{r} = \underset{3}{\mathcal{F}_N^{-1}} \left\{ \underset{1}{\mathcal{F}_N\{\underline{x}\}} \odot \underset{2}{\mathcal{F}_N\{\underline{y}'\}} \right\}$$

1. *Sketch of  $\underline{x}$*  : Assume  $\underline{X}[l] = \mathcal{F}\{\underline{x}\}$  is precomputed at positions  $l \in \mathcal{S}$ .
2. *Sketch of  $y$* :
  - Compute  $\underline{Y}'[l] = \mathcal{F}\{\underline{y}'\}$  for  $l \in \mathcal{S}$ .
  - Only  $M$  non-zero values in  $\underline{y}'$  - Efficient computation (folding and adding)
3. *Sparse  $\mathcal{F}^{-1}$* :
  - Robust Sparse Inverse Fourier Transform (RSIDFT)
  - Efficient Implementation- **sublinear** time and sampling complexity

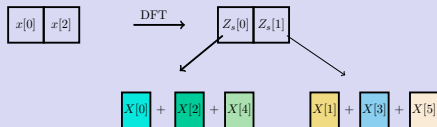
# Aliasing and Sparse Graph Codes



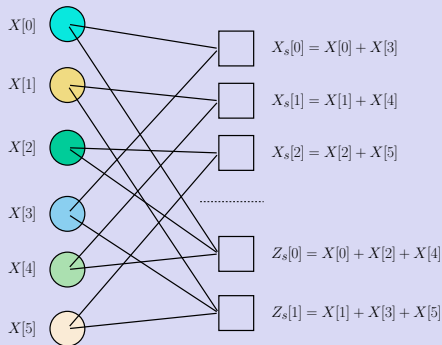
$x_s$ : Sub-sampled by  $f_1 = P_1 = 2$



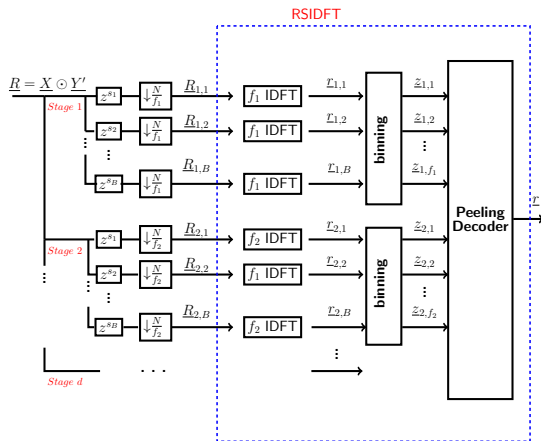
$z_s$ : Sub-sampled by  $f_2 = P_2 = 3$



Factor graph



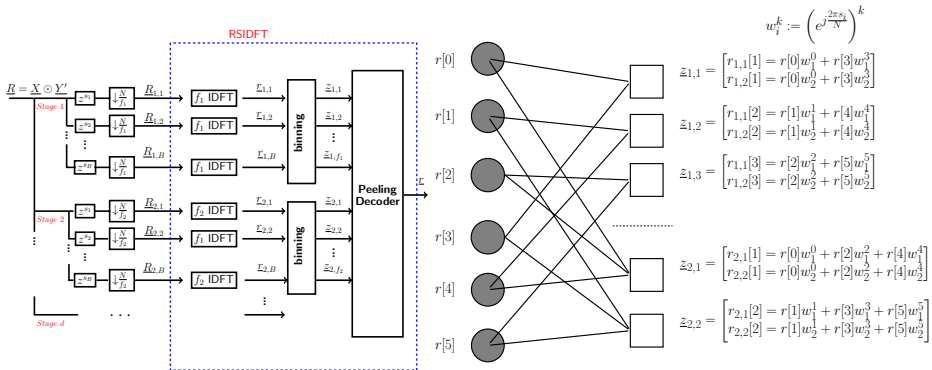
# RSIDFT Framework



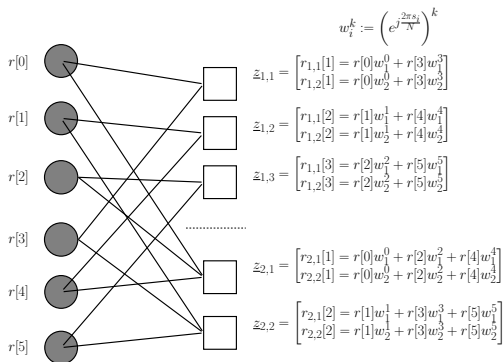
- Similar to FFAST
- Key modifications
  - Correlation peak is always **positive**
  - Take advantage in decoding algorithm - **sub-linear** time complexity



# RSIDFT Framework



# RSIDFT-Decoding (Peeling Decoder)



- **Observations:**  $\underline{z}_{i,k} = [r_{i,1}[k], r_{i,2}[k], \dots, r_{i,B}[k]]^T$
- **Decoding- 3 steps**
  - 1 Bin Classification
  - 2 Position Identification
  - 3 Peeling Process

# Decoding Process

## Bin Classification

- Classify each check-node - Zero-ton / Single-ton / Multi-ton
- **Threshold constraints** on first observation  $z_{i,k}[1] = z$
- Threshold varies with  $\eta$ 
  - different for exact( $\eta = 0$ ) and approximate matching

$$\hat{\mathcal{H}}_{i,j} = \begin{cases} \mathcal{H}_z & z/M < \gamma_1 \\ \mathcal{H}_s & \gamma_1 < z/M < \gamma_2 \\ \mathcal{H}_d & \gamma_2 < z/M < \gamma_3 \\ \mathcal{H}_m & z/M > \gamma_3 \end{cases}$$

where  $(\gamma_1, \gamma_2, \gamma_3) = (\frac{1-2\eta}{2}, \frac{3-4\eta}{2}, \frac{5-6\eta}{2})$

# Decoding Process

## Position Identification

- Observation:

$$\underline{z}_{i,k} = \mathbb{W}_{i,k} \times \begin{bmatrix} r[k + (0)f_i] \\ r[k + (1)f_i] \\ \vdots \\ r[k + (g_i - 1)f_i] \end{bmatrix}$$

- Sensing Matrix:

$$\mathbb{W}_{i,k} = \left[ \underline{w}^k, \underline{w}^{k+f_i}, \dots, \underline{w}^{k+(g_i-1)f_i} \right] \text{ where } \underline{w}^k = \begin{bmatrix} e^{\frac{j2\pi ks_1}{N}} \\ e^{\frac{j2\pi ks_2}{N}} \\ \vdots \\ e^{\frac{j2\pi ks_B}{N}} \end{bmatrix}, \quad g_i = \frac{N}{f_i}$$

- Column that gives **maximum correlation** with the observation

$$\hat{k} = \arg \max_{k \in \{j+lg_i\}} \underline{z}_{i,j}^\dagger \underline{w}^k$$

# Decoding Process

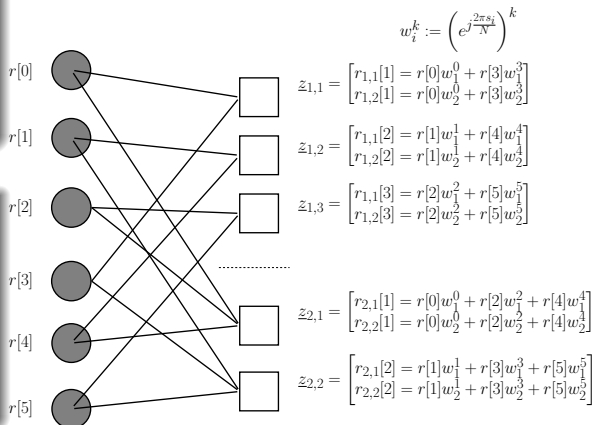
## Peeling Process:

### Exact Matching

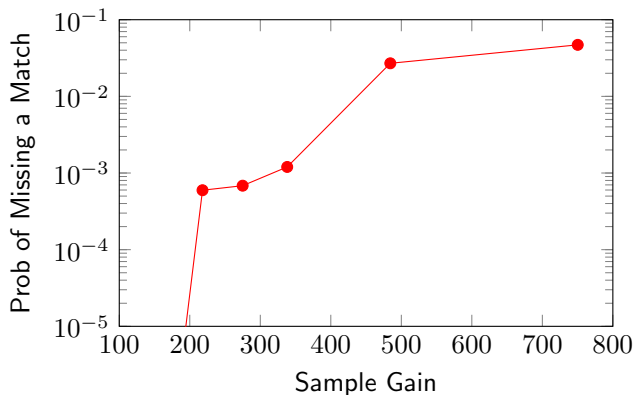
- Remove a decoded variable node's contribution from **all** participating bin nodes

### Approximate Matching

- Remove a decoded variable node's contribution only from neighboring **single-tons** and **double-tons**
- Avoid error propagation

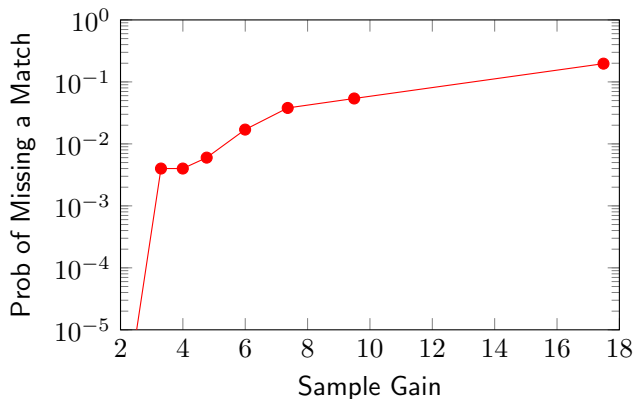


# Simulation Results



**Figure:** Plot of Probability of Missing a Match vs. Sample Gain for Exact Matching of a substring of length  $M = 10^5$  from a equiprobable binary  $\{+1, -1\}$  sequence of length  $N = 10^{12}$ , divided into  $G = 10^5$  blocks each of length  $\tilde{N} = 10^7$ . The substring was simulated to repeat in  $L = 10^6$  locations uniformly at random.

## Simulation Results



**Figure:** Plot of Probability of Missing a Match vs. Sample Gain for Exact Matching of a substring of length  $M = 10^3$  from a equiprobable binary  $\{+1, -1\}$  sequence of length  $N = 10^{12}$ , divided into  $G = 10^6$  blocks each of length  $\tilde{N} = 10^6$ . The substring was simulated to repeat in  $L = 10^6$  locations uniformly at random.

# Questions?



# Thank you!