

Identifying a piecewise affine signal from its nonlinear observation - application to DNA replication analysis

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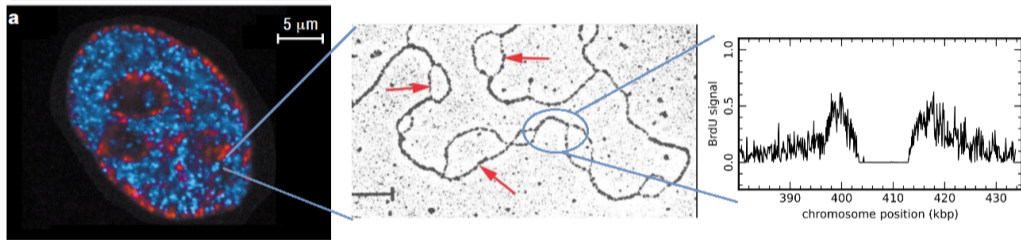
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Toulouse, October 2024

DNA replication analysis

General framework

- Microscope images are widely used to study the structure and function of cells
- Some characteristics of the replication can be studied by images



late replication
early replication

- This work aims to understand parameters of replication such as: position of origins, local speed, and replication direction

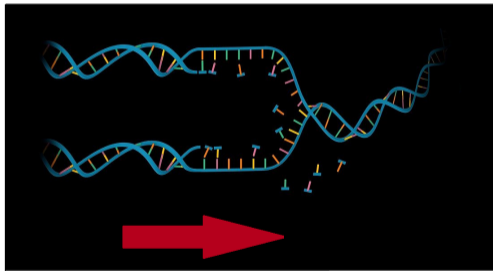
Important parameters

Main Objective: Better characterize the replication of DNA.

- Position of origins of replication
- Speed and direction of replication

Related application:

- Characterization of replication stress in cancer cells



DNA synthesis at a replication fork moving to the right

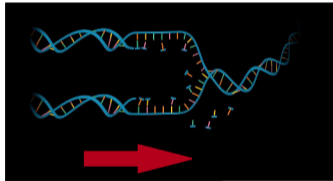
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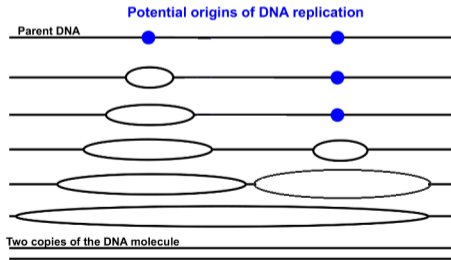
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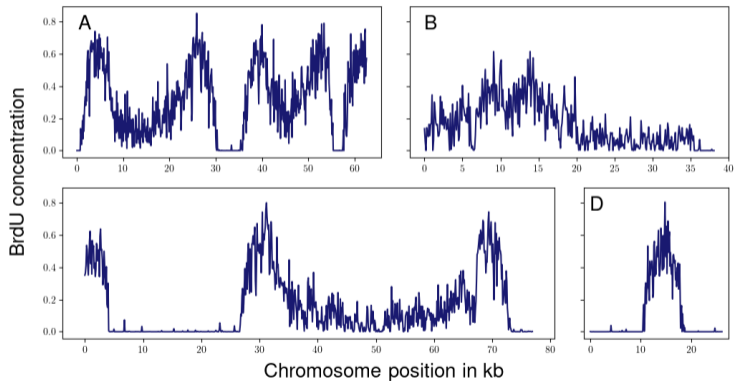
DNA synthesis at a replication fork moving to the right



DNA replication signal

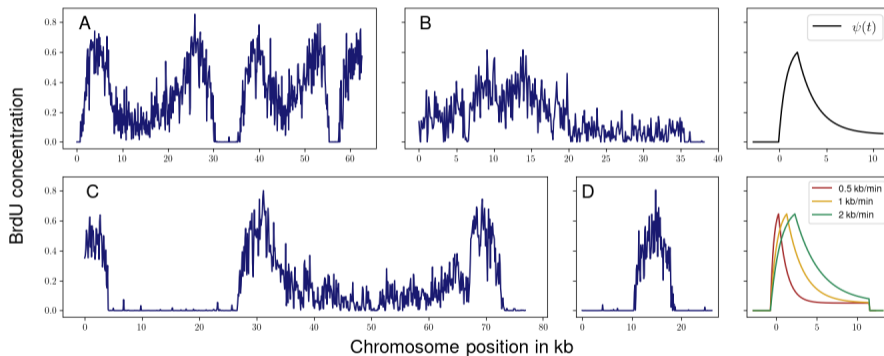
DNA Replication signal

- 1d signals generated during the replication process
- Signals combine DNA sequencing and the concentration of a chemical: BrdU



Dictionary approach

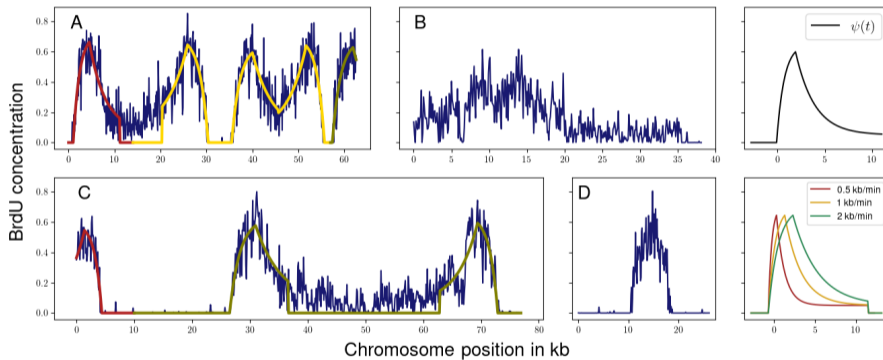
→ ψ can be used as an atom that can be translated and dilated



$$\begin{aligned} \min_x \quad & \|z - Dx\|_2^2 \\ \text{s.t.} \quad & \{x : \|x\|_0 \leq C\} \end{aligned}$$

Dictionary approach

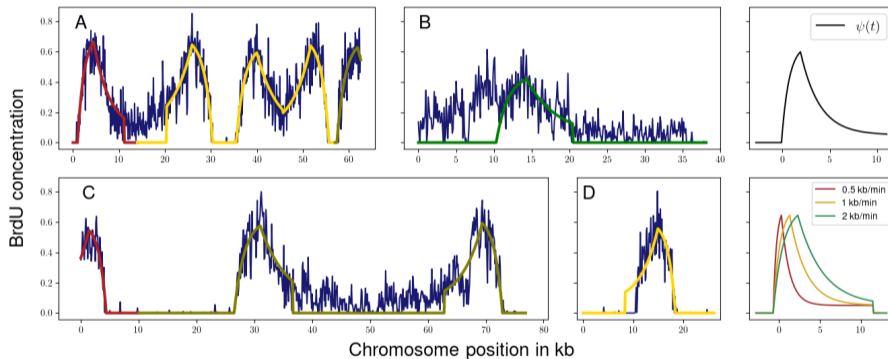
→ The dilatation is able to detect local speed for some signals



$$\begin{aligned} \min_x \quad & \|z - Dx\|_2^2 \\ \text{s.t.} \quad & \{x : \|x\|_0 \leq C\} \end{aligned}$$

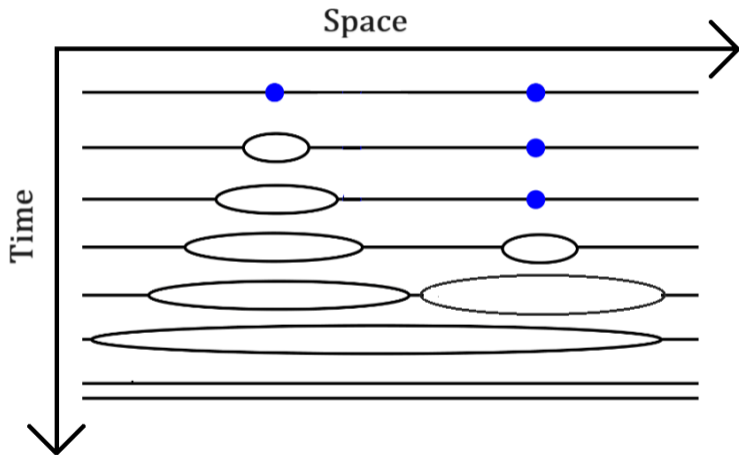
Dictionary approach

→ The method does not cover all signals

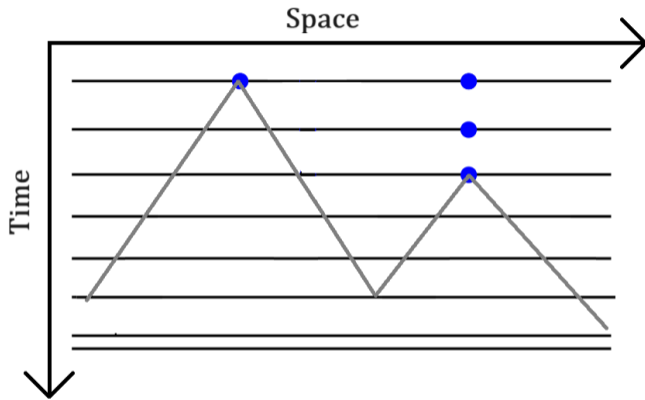


$$\begin{aligned} \min_x \quad & \|z - Dx\|_2^2 \\ \text{s.t.} \quad & \{x : \|x\|_0 \leq C\} \end{aligned}$$

Back to the basis

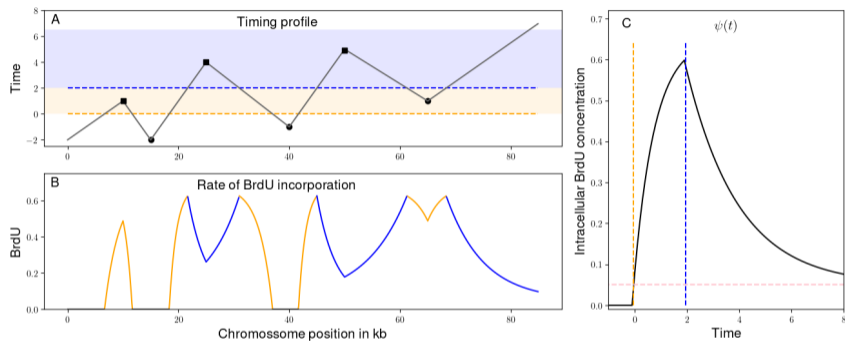


Timing profile



Assumption: Speed of replication is constant between an origin and a terminus

Change of perspective



τ - Space \times Time, Piecewise linear function

ψ - Time \times BrdU, concentration function

z - Space \times BrdU, Signal: $z(x) = \psi(\tau(x))$

The DNA replication as a nonlinear inverse problem

- Set of equidistant points $\mathcal{X} = \{x_1, \dots, x_n\}$
- Timing profile: $\boldsymbol{\tau} = (\tau_1, \dots, \tau_n) := (\tau(x_1), \dots, \tau(x_n))$

→ Measurement operator:

$$\begin{aligned}\Psi : \mathbb{R}^n &\longrightarrow \mathbb{R}_+^n \\ \boldsymbol{\tau} &\mapsto (\psi(\tau_1), \dots, \psi(\tau_n))\end{aligned}$$

Consider the inverse problem:

$$\min_{\boldsymbol{\tau} \in \mathcal{P}_C} \|\mathbf{z} - \Psi(\boldsymbol{\tau})\|_2^2, \quad (\mathbf{P1})$$

where:

$$\mathcal{P}_C := \{\boldsymbol{\tau} : \|\mathbf{L}\boldsymbol{\tau}\|_0 \leq C\}, \quad \mathbf{L}\boldsymbol{\tau} = \ell * \boldsymbol{\tau}, \quad \text{with } \ell = [1, -2, 1].$$

\mathcal{P}_C is the set of piecewise linear vectors with at most C breaks.

Nonlinear inverse problem

Challenges

$$\hat{\boldsymbol{\tau}} := \arg \min_{\boldsymbol{\tau} \in \mathcal{P}_C} \|\mathbf{z} - \Psi(\boldsymbol{\tau})\|_2^2 \quad \textbf{(P.1)}$$

- Ψ is nonlinear
- Ψ is not injective

Objective: Provide a global solutions to problem **(P1)**

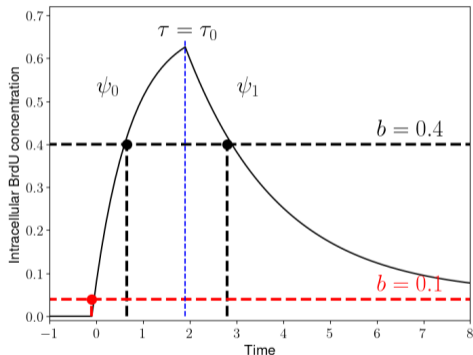
Assumption

→ The form of the operator Ψ is important

→ $\Psi(\tau) = (\psi(\tau_1), \dots, \psi(\tau_n))$

A.1 : $\exists \tau_0 > 0$ such that:

- $\psi_0 := \psi|_{[0, \tau_0]}$ is concave
- $\psi_1 := \psi|_{[\tau_0, \infty)}$ is convex
- The convexity or concavity of ψ_0 or ψ_1 is strict.
- both ψ_0, ψ_1 are injective.



As a consequence: $\#\psi^{-1}(b) = \#\{\tau : \psi(\tau) = b\} \leq 2, \forall b \in \mathbb{R}_+$

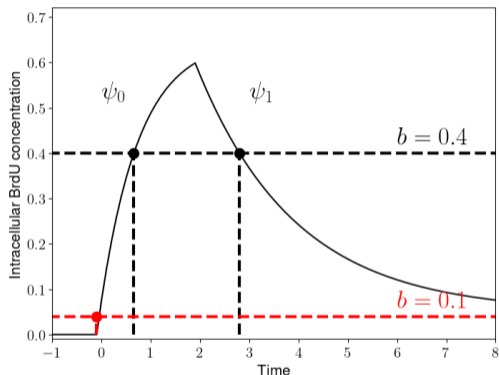
Alternative formulation

If Ψ is not injective, the definition of **P.2** is not intuitive.

$$\boxed{\min_{\tau \in \mathcal{P}_C} \|\mathbf{z} - \Psi(\tau)\|_2^2 \quad (\text{P.1})} \longleftrightarrow \boxed{? \quad (\text{P.2})}$$

→ Use the property: $\#\psi^{-1}(b) \leq 2, \forall b \in \mathbb{R}_+$ to enumerate the inverse set

Inverse image



→ Inverse image of ψ for $b \in \mathbb{R}_+$:

$$\psi^{-1}(b) = \psi_0^{-1}(b) \cup \psi_1^{-1}(b)$$

→ Inverse image of Ψ for $\mathbf{d} \in \{0, 1\}^n$:

$$\Psi_{\mathbf{d}}^{-1}(\mathbf{z}) := \psi_{d_0}^{-1}(z_0) \times \dots \times \psi_{d_n}^{-1}(z_n)$$

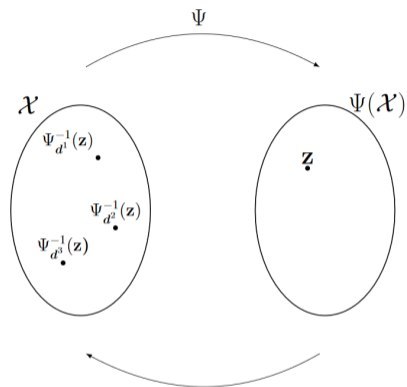
→ Define:

$$\mathcal{K}(\mathbf{z}) = \left\{ \mathbf{d} \in \{0, 1\}^n : \psi_{d_i}^{-1}(z_i) \neq \emptyset \right\}$$

→ Then: $\Psi_{\mathbf{d}}^{-1}(\mathbf{z}) \in \mathbb{R}^n$, for $\mathbf{d} \in \mathcal{K}(\mathbf{z})$

$$\Psi(\boldsymbol{\tau}) = (\psi(\tau_1), \psi(\tau_2), \dots, \psi(\tau_n))$$

Taylor approximation



Because of Taylor approximation theorem, for each $\mathbf{d} \in \mathcal{K}(\mathbf{z})$

$$\|\mathbf{z} - \Psi(\boldsymbol{\tau})\|_2^2 \approx \sum_{i=1}^n w_{\mathbf{d},i}^2 (\Psi_{\mathbf{d}}^{-1}(\mathbf{z}) - \boldsymbol{\tau})_i^2,$$

Taylor approximation

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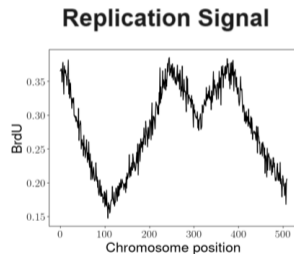
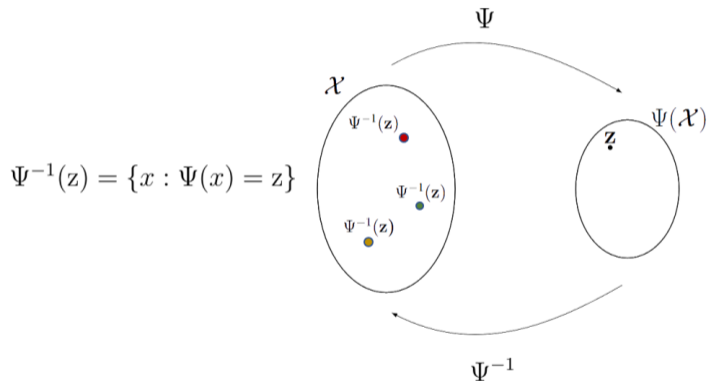
where $w_{\mathbf{d},i} := \psi'(\psi_{d_i}^{-1}(z_i)) \in \mathbb{R}$. We define: $\|\mathbf{v}\|_{\mathbf{w}} := \sqrt{\sum_{i=1}^n w_i^2 v_i^2}$. Then:

$$\text{For each } \mathbf{d} \in \{0, 1\}^n : \quad \|\mathbf{z} - \Psi(\boldsymbol{\tau})\|_2^2 \approx \|\Psi_{\mathbf{d}}^{-1}(\mathbf{z}) - \boldsymbol{\tau}\|_{\mathbf{w}_{\mathbf{d}}}^2$$

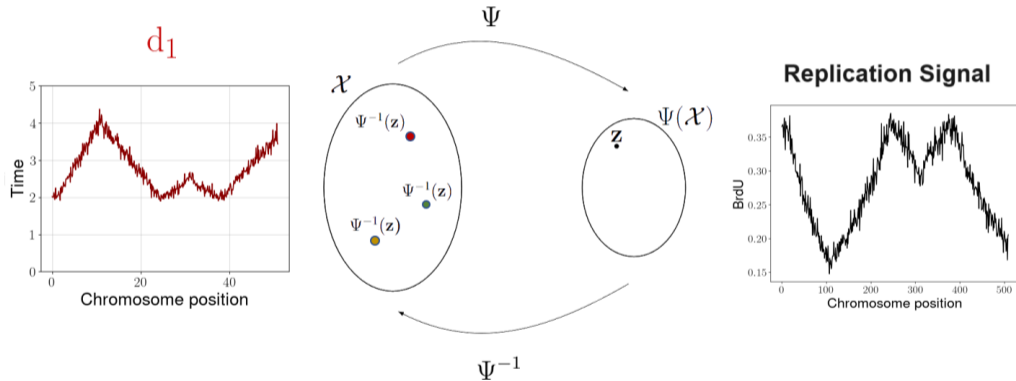
Resulting in:

$$\min_{\{(\boldsymbol{\tau}, \mathbf{d}) \in \mathcal{P}_C \times \{0, 1\}^n\}} \|\Psi_{\mathbf{d}}^{-1}(\mathbf{z}) - \boldsymbol{\tau}\|_{\mathbf{w}_{\mathbf{d}}}^2 \quad (\mathbf{P.2})$$

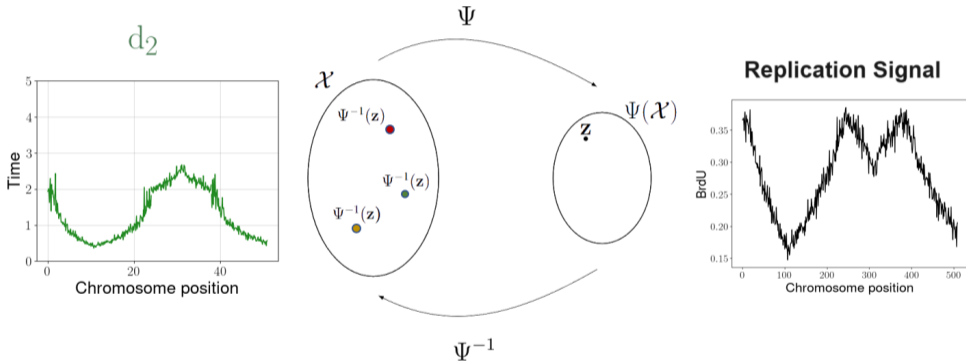
Intuition of (P.2) with a simulated signal



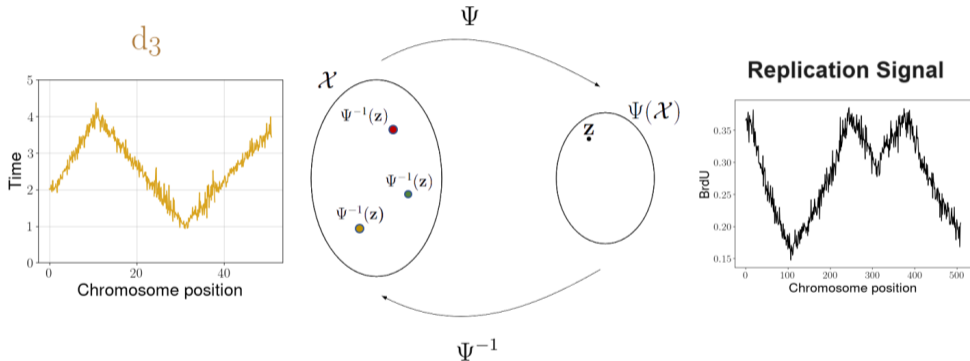
Intuition of (P.2) with a simulated signal



Intuition of (P.2) with a simulated signal



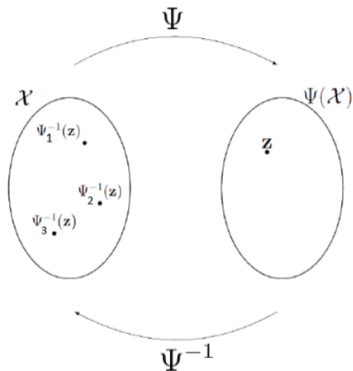
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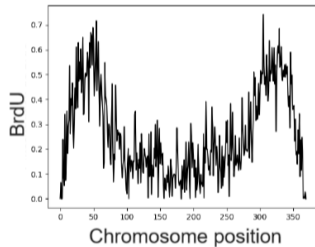
→ One of the inverse images is close to a piecewise linear behavior

Intuition of weights w_d in a real signal

$$\Psi^{-1}(z) = \{x : \Psi(x) = z\}$$

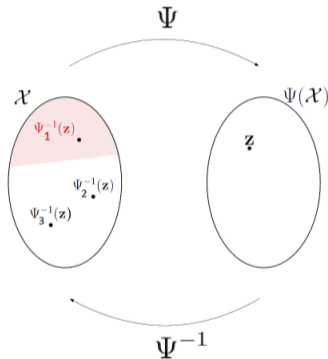


Replication Signal

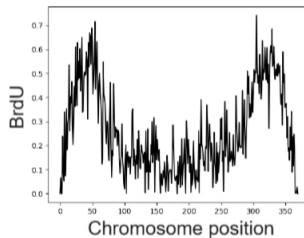


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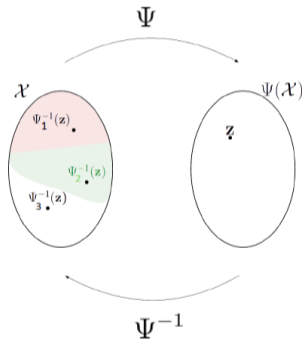
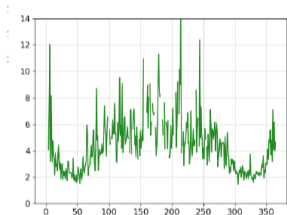


Replication Signal

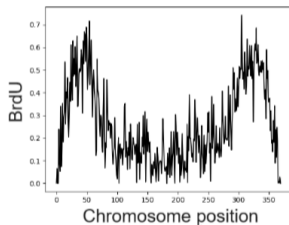


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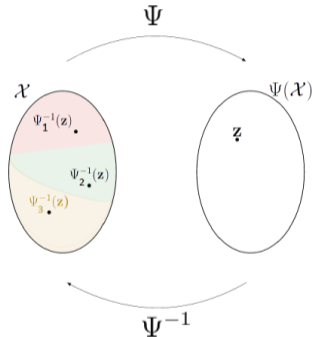
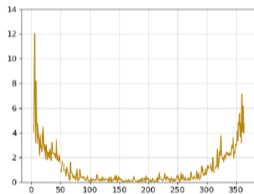


Replication Signal

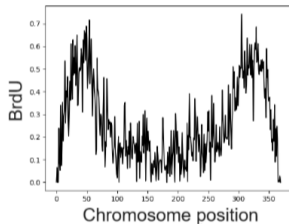


Intuition of weights w_d in a real signal

$$\Psi^{-1}(z) = \{x : \Psi(x) = z\}$$



Replication Signal



→ One of the inverse images is close to a piecewise linear behavior

Equivalence in noiseless case

$$\hat{\tau} := \arg \min_{\{\tau \in \mathcal{P}_C\}} \|z - \Psi(\tau)\|_2^2 \quad (\text{P.1}) \quad \longleftrightarrow \quad (\tau^*, \mathbf{d}^*) := \arg \min_{\{(\tau, \mathbf{d}) \in \mathcal{P}_C \times \mathcal{K}(z)\}} \|\tau - \Psi_{\mathbf{d}}^{-1}(z)\|_{w_{\mathbf{d}}}^2 \quad (\text{P.2})$$

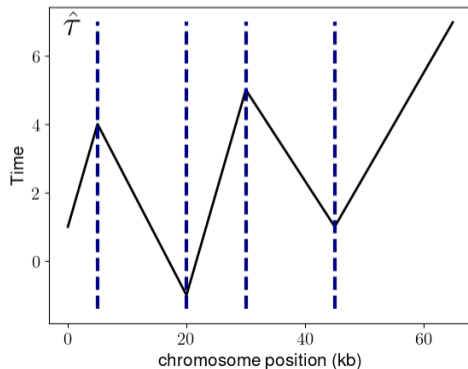
Theorem. Suppose $\hat{\tau}$ or τ^* do not have constant parts and its breakpoints are sufficiently spaced (more than 1.2 kb). Then $\hat{\tau} = \tau^*$, and \mathbf{d}^* is such that $\Psi_{\mathbf{d}^*}^{-1}(z) = \tau^* = \hat{\tau}$

Proof.

Based on the injectivity of Ψ restricted to the correspondent set in \mathcal{P}_C . □

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Numerical method

Rewriting (P.2')

$$\min_{\{(\boldsymbol{\tau}, \mathbf{d}) \in \mathcal{P}_C \times \{0,1\}^n\}} \|\Psi_{\mathbf{d}}^{-1}(\mathbf{z}) - \boldsymbol{\tau}\|_{\mathbf{w}_{\mathbf{d}}}^2 \quad (\text{P.2})$$

$$\begin{aligned} &= (\odot \text{ coordinatewise multiplication}) \\ &\quad \mathbf{w}_{\mathbf{d}} = \mathbf{d} \odot \mathbf{w}_1 + (\mathbf{1} - \mathbf{d}) \odot \mathbf{w}_0 \end{aligned}$$

$$\begin{aligned} \min_{\boldsymbol{\tau}, \mathbf{d}} \quad & \frac{1}{2} \|\mathbf{d} \odot (\boldsymbol{\tau} - \Psi_1^{-1}(\mathbf{z}))\|_{\mathbf{w}_1}^2 + \frac{1}{2} \|(\mathbf{1} - \mathbf{d}) \odot (\boldsymbol{\tau} - \Psi_0^{-1}(\mathbf{z}))\|_{\mathbf{w}_0}^2 \\ \text{s.t.} \quad & \boldsymbol{\tau} \in \mathbb{R}^n, \|\mathbf{L}\boldsymbol{\tau}\|_0 \leq C \\ & \mathbf{d} \in \{0,1\}^n \end{aligned}$$

$$\approx (\ell_1 \text{ regularization}) \quad \lambda > 0$$

$$\begin{aligned} \min_{\boldsymbol{\tau}, \mathbf{d}} \quad & \frac{1}{2} \|\mathbf{d} \odot (\boldsymbol{\tau} - \Psi_1^{-1}(\mathbf{z}))\|_{\mathbf{w}_1}^2 + \frac{1}{2} \|(\mathbf{1} - \mathbf{d}) \odot (\boldsymbol{\tau} - \Psi_0^{-1}(\mathbf{z}))\|_{\mathbf{w}_0}^2 + \lambda \|\mathbf{L}\boldsymbol{\tau}\|_1 \quad (\text{P.3}) \\ \text{s.t.} \quad & \boldsymbol{\tau} \in \mathbb{R}^n, \\ & \mathbf{d} \in \{0,1\}^n \end{aligned}$$

Solving the regularized problem

$$\begin{aligned} \min_{\boldsymbol{\tau}, \mathbf{d}} \quad & \frac{1}{2} \|\mathbf{d} \odot (\boldsymbol{\tau} - \Psi_1^{-1}(\mathbf{z}))\|_{\mathbf{w}_1}^2 + \frac{1}{2} \|(\mathbf{1} - \mathbf{d}) \odot (\boldsymbol{\tau} - \Psi_0^{-1}(\mathbf{z}))\|_{\mathbf{w}_0}^2 + \lambda \|\mathbf{L}\boldsymbol{\tau}\|_1 \quad (\mathbf{P.3}) \\ \text{s.t.} \quad & \boldsymbol{\tau} \in \mathbb{R}^n, \\ & \mathbf{d} \in \{0, 1\}^n \end{aligned}$$

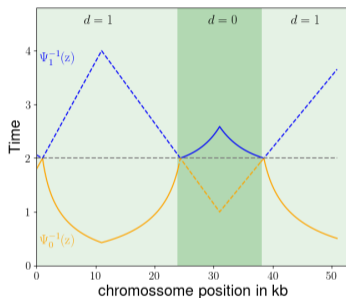
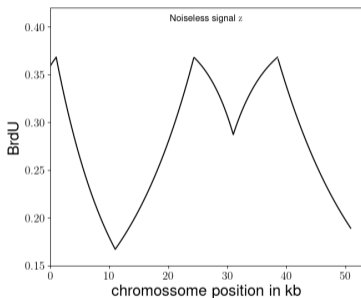
→ For each $\mathbf{d} \in \{0, 1\}^n$, **(P.3)** is convex

→ For each $\mathbf{d} \in \{0, 1\}^n$, **(P.3)** is similar to *generalized lasso*

Proposition. For each $\mathbf{d} \in \{0, 1\}^n$, problem **(P.3)** is equivalent to its dual formulation, which is a *quadratic optimization problem*.

Constraining the set $\{0, 1\}^n$

- The set $\{0, 1\}^n$ is excessively large to be tractable
- We can constraint this set without changing its optimal solution



$$\mathcal{D} = \{\mathbf{d} : \mathbf{d} \text{ can be the optimal solution}\} = \{\mathbf{d} : d_i = d_{i+1}, \forall i \in I_{\mathcal{A}}\}$$

$$I_{\mathcal{A}} = \{i : |\psi_0^{-1}(z_i) - \psi_1^{-1}(z_i)| > \epsilon\}$$

Algorithm

Algorithm: DNA-Inverse

Data: Input data: \mathbf{z} . Parameters: $\epsilon > 0$.

Initialization:

Compute weights w_d . Compute set \mathcal{D} . Set: $\mathcal{D}_{\text{past}} = \emptyset$;

Main Loop:

for $d \in \mathcal{D}$ do

Step 0: $\mathcal{D}_{\text{past}} \leftarrow \mathcal{D}_{\text{past}} \cup \{d\}$;

Step 1: Solve (P.3), obtaining a solution τ_d^* ;

Step 2:

$$d^* := \arg \min_{d \in \mathcal{D}_{\text{past}}} F(\tau_d^*), \quad \tau^* := \tau_{d^*}^*,$$

Output: τ^*, d^*

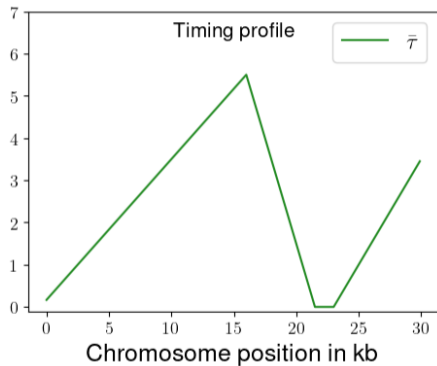
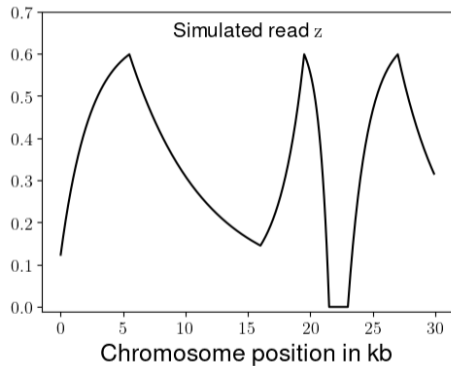
where:

$$F(\tau) := \frac{1}{2} \|\mathbf{d} \odot (\tau - \Psi_1^{-1}(\mathbf{z}))\|_{\mathbf{w}_1}^2 + \frac{1}{2} \|(\mathbf{1} - \mathbf{d}) \odot (\tau - \Psi_0^{-1}(\mathbf{z}))\|_{\mathbf{w}_0}^2$$

Numerical results

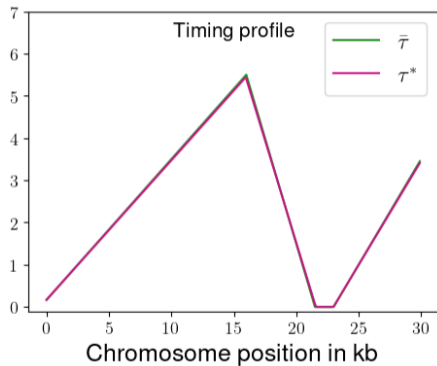
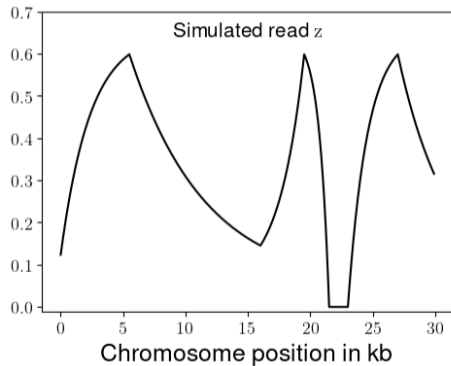
Methods for nonlinear inverse problem

→ Example of a noiseless signal



Methods for nonlinear inverse problem

→ DNA Inverse solution (pink)



Methods for nonlinear inverse problem

→ Other optimization methods can solve the ℓ_1 -regularized (P.1)

$$\min_{\boldsymbol{\tau} \in \mathbb{R}^n} \|\mathbf{z} - \Psi(\boldsymbol{\tau})\|_2^2 + \gamma \|\mathbf{L}\boldsymbol{\tau}\|_1,$$

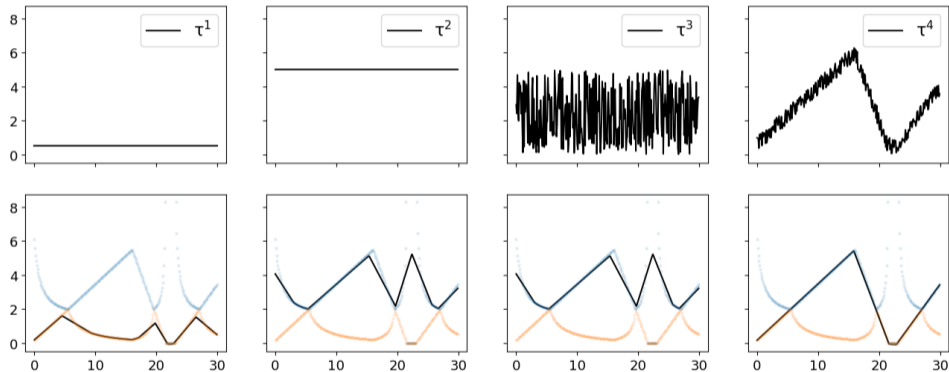
→ One of these methods is the Primal-Dual method (Valkonen, 2019):

$$G(\mathbf{u}) = \|\mathbf{u} - \mathbf{z}\|_2^2, \quad G^*(\mathbf{y}) = \sup_{\mathbf{u} \in \mathbb{R}^n} \langle \mathbf{u}, \mathbf{y} \rangle - G(\mathbf{u}).$$

$$\begin{cases} \boldsymbol{\tau}^{k+1} = \text{prox}_{\sigma_1 \gamma \|\cdot\|_1}(\boldsymbol{\tau}^k - \sigma_1 \Psi'(\boldsymbol{\tau}^k) \mathbf{y}^k) \\ \mathbf{y}^{k+1} = \text{prox}_{\sigma_2 (G^* - 2\langle \Psi(\boldsymbol{\tau}^k), \cdot \rangle)}(\mathbf{y}^k - \sigma_2 \Psi(\boldsymbol{\tau}^k)), \end{cases}$$

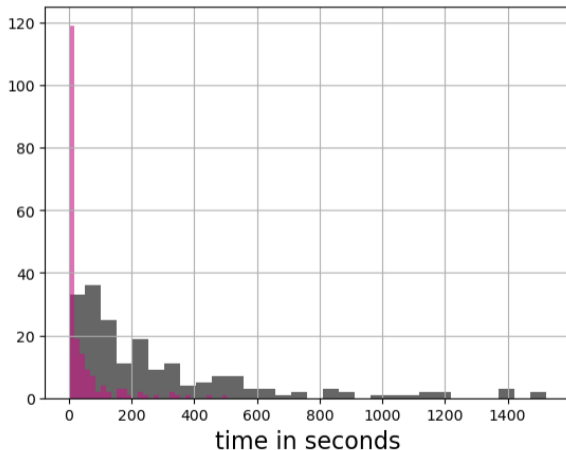
Methods for nonlinear inverse problem

→ The PDPS provide local solutions that depend on the initial point: $\tau^i, i \in \{1, 2, 3, 4\}$



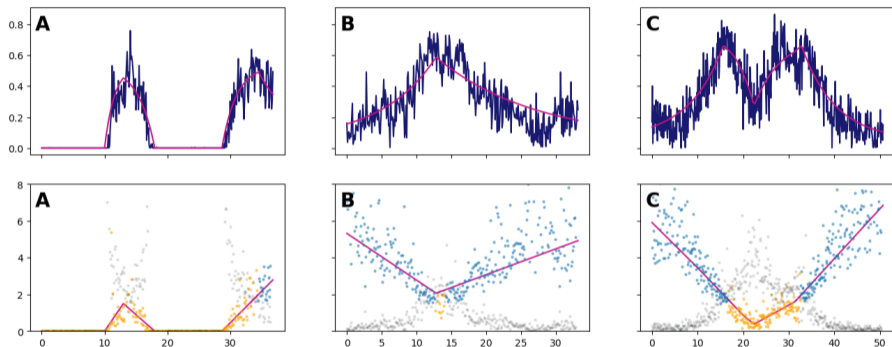
Execution time

→ When compared to PDPS with smart initialization (grey), DNA inverse (pink) exhibits faster performance



Results for real data

→ DNA inverse identifies the biological reality behind the signal




→ Timing profile τ^* (below, pink), and a signal approximation (above, pink)

→ The colored dots are selected by the integer variable d^*

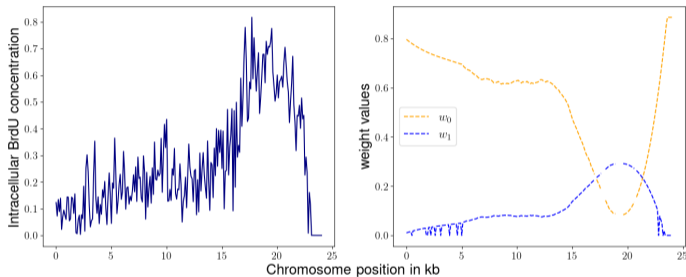
Thanks for your attention!!

References

-  Valkonen, T. et al (2019). “Acceleration and Global Convergence of a First-Order Primal-Dual Method for Nonconvex Problems”. In: *SIAM Journal on Optimization* 29.1, pp. 933–963. DOI: [10.1137/18M1170194](https://doi.org/10.1137/18M1170194). URL: <https://doi.org/10.1137/18M1170194>.

Extension to the noisy case

- We extend \mathbf{w}_d to be zero in coordinates $i \in \{1, \dots, n\}$ such that $\psi_{d_i}(\mathbf{z}) = \emptyset$
- These coordinates contain less information about the signal position



In the noisy case, consider the optimization problem:

$$(\boldsymbol{\tau}^*, \mathbf{d}^*) := \arg \min_{\{(\boldsymbol{\tau}, \mathbf{d}) \in \mathcal{P}_C \times \{0,1\}^n\}} \|\Psi_{\mathbf{d}}^{-1}(\mathbf{z}) - \boldsymbol{\tau}\|_{\mathbf{w}_d}^2 \quad (\mathbf{P.2}')$$