



## Transcription factor binding sites detection in

### Paramecium

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Keyword1 | Keyword2 | Keyword3

Abbreviations: SAM, self-assembled monolayer; OTS, octadecyltrichlorosilane

#### Introduction

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- Whole Genome Duplications background
- WGD influences on Transcription Factor Binding Sites?
- the *Paramecium* project, experimental part, etc. some work is done and random sequences?
- Here, focusing on the computational part, developing pipeline, showed that etc.

#### Results

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#### Simulations.

#### Simulation 1

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#### Simulation 2

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Real Data. Aliquam interdum pellentesque scelerisque. Sed tincidunt suscipit purus, id aliquet nulla vehicula quis. Duis sed nisl lorem. Vivamus erat ante, dignissim et aliquam vel, adipiscing vitae magna. Cras id dapibus metus. Cum sociis natoque penatibus et magnis dis parturient montes, nascetur ridiculus mus. Proin ut lectus ut nisi congue ullamcorper. Ut ac turpis ligula. Sed faucibus bibendum nunc eget gravida.

#### Discussion

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Reserved for Publication Footnotes

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#### Materials and Methods

-i developped a whole pipeline (show simple pipeline graph) From annotations and assembly files -i extracted upstream sequences of given length minimum length 15nt cutoff at 250nt. From families with specific conditions. X families identified, In each family CDSs were retrieved, aligned using protein-guided alignment (TranslatorX) and ML tree (PhyML) was computed -i used BigFoot to consider the phylogenetic signal +i Comparison with MEME.

From this, X motifs matched partially (quantify!) between MEME and BigFoot - ¿ search through the genome this motifs in each species. Using publicly available RNA expression data, search correlations between motifs and expression level.

**Definition 1.** A bounded function  $\theta$  is a weak solution of QG if for any  $\phi \in C_0^\infty(\mathbb{R}/\mathbb{Z} \times \mathbb{R} \times [0, \varepsilon])$  we have

$$\begin{split} &\int_{\mathbb{R}^{+}\times\mathbb{R}/_{\mathbb{Z}}\times\mathbb{R}}\theta(x,y,t)\,\partial_{t}\phi\left(x,y,t\right)dydxdt + \\ &+ &\int_{\mathbb{R}^{+}\times\mathbb{R}/_{\mathbb{Z}}\times\mathbb{R}}\theta\left(x,y,t\right)u(x,y,t)\cdot\nabla\phi\left(x,y,t\right)dydxdt = 0 \quad \text{[1]} \end{split}$$

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where u is determined previously.

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**Theorem 1.** If the active scalar  $\theta$  satisfies the equation [1], then  $\varphi$  satisfies the equation

$$\frac{\partial \varphi}{\partial t}(x,t) = \int_{\mathbb{R}/\mathbb{Z}} \frac{\frac{\partial \varphi}{\partial x}(x,t) - \frac{\partial \varphi}{\partial u}(u,t)}{\left[(x-u)^2 + (\varphi(x,t) - \varphi(u,t))^2\right]^{\frac{1}{2}}} \\
\chi(x-u,\varphi(x,t) - \varphi(u,t))du + \\
+ \int_{\mathbb{R}/\mathbb{Z}} \left[\frac{\partial \varphi}{\partial x}(x,t) - \frac{\partial \varphi}{\partial u}(u,t)\right] \\
\eta(x-u,\varphi(x,t) - \varphi(u,t))du + Error \quad [2]$$

with  $|Error| \leq C \, \delta |log\delta|$  where C depends only on  $\|\theta\|_{L^{\infty}}$  and  $\|\nabla \varphi\|_{L^{\infty}}$ .

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#### **Appendix**

An appendix without a title.

#### Appendix: Appendix title

An appendix with a title.

**ACKNOWLEDGMENTS.** This work was partially supported by a grant from the Spanish Ministry of Science and Technology.

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 $FIGURE \ \textbf{1.} \quad \text{Figure caption}$ 

Table 1. Table caption

Treatments	Response 1	Response 2
Treatment 1	0.0003262	0.562
Treatment 2	0.0015681	0.910
Treatment 3	0.0009271	0.296

