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You will indemnify, defend and hold harmless KULEUVEN, its directors, officers, employees and agents from and against all liability, losses, damages and expenses (including attorney's fees and costs) arising out of any claims, demands, actions or other proceedings made or instituted by any third party against any of them and arising out of or relating to any breach of this Agreement by you, or any use of the Software by you, except insofar as such claims or liability result from KULEUVEN gross negligence or willful misconduct.

#### Article 8 - Term

- 8.1 This Agreement is effective from the Effective Date until you delete the Software and any and all related files from your computing system. This Agreement will terminate immediately without notice from K.U. LEUVEN if you fail to comply with any provision of this Agreement.
- 8.2 In case of termination the provisions of Article 3, 6, and 7 shall remain in full force and effect.

#### Article 9 - Miscellaneous

- 9.1 Any notice authorised or required to be given to KULEUVEN under this Agreement shall be in writing and shall be deemed to be duly given if left at or sent by registered post.
- 9.2 The terms and conditions herein contained constitute the entire agreement between the Parties and supersede all previous agreements and understandings, whether oral or written, between the parties hereto with respect to the subject matter thereof.

## **Article 10 - Conflicts**

In the event of conflicts in the interpretation and/or performance of this Agreement, the parties shall first undertake to settle their differences amicably. If no amicable settlement can be reached concerning the execution and/or interpretation of this Agreement, such conflict shall be brought before the courts of Leuven and Belgian Law shall be applicable.

Choose one of the options below:

By downloading and or installing the Software and associated files on your computing system you agree to use the Software under the terms and condition as specified in this Agreement.

# Or

By crossing the I agree button below	you acknowledge that yo	ou agree on terms of	use specified above
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## Annex A: ModuleDigger

### Description of the Software:

The detection of *cis*-regulatory modules (CRMs) that mediate transcriptional responses in eukaryotes remains a key challenge in the postgenomic era. A CRM is characterized by a set of co-occurring transcription factor binding sites (TFBS). *In silico* methods have been developed to search for CRMs by determining the combination of TFBS that are statistically overrepresented in a certain geneset. Most of these methods solve this combinatorial problem by relying on computational intensive optimization methods. As a result their usage is limited to finding CRMs in small datasets (containing a few genes only) and using binding sites for a restricted number of transcription factors (TFs) out of which the optimal module will be selected.

ModuleDigger is a tool based on an itemset mining based strategy for computationally detecting *cis*-regulatory modules (CRMs) in a set of genes. ModuleDigger can handle larger dataset as well as considering a genome-wide view which means it is considered specific for the whole cluster of input genes if the CRM is statistically more overrepresented in this cluster of genes than in the remainder of the genome. By exploiting the computational efficiency of an itemset mining approach and combining it with a well-designed statistical scoring scheme, we were able to prioritize the biologically valid CRMs in a large set of coregulated genes using binding sites for a large number of potential TFs as input.

#### Reference:

- 1. Sun H., De Bie T., Storms V., Fu Q., Dhollander T., Lemmens K., Verstuyf A., De Moor B., Marchal K., "ModuleDigger: an itemset mining framework for the detection of cis-regulatory modules". BMC bioinformatics, Vol. 10 Suppl 1 (2009).
- 2. Sun, H., De Bie, T., Storms, V., Fu, Q., Dhollander, T., Lemmens, K., Verstuyf, A., De Moor, B., Marchal K., ModuleDigger: an itemset mining framework for the detection of cis-regulatory modules. In: Proceedings of 7th Asia-Pacific Bioinformatics Conference APBC2009.