

Motif-based method for genome-wide prediction of eukaryotic gene clusters

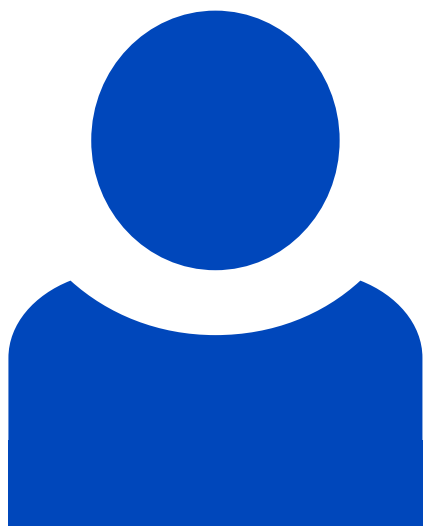
Wolf T, Shelest V, Shelest E (2013) Motif-based method for genome-wide prediction of eukaryotic gene clusters In: A. Petrosino, L. Maddalena, P. Pala (eds.) Lecture Notes in Computer Science (LNCS) 2nd International Workshop on Pattern Recognition in Proteomics, Structural Biology and Bioinformatics, Naples, Italien, 09/09/2013-09/10/2013, 8158, pp. 389-398. Springer-Verlag, Berlin, Heidelberg.

[Details](#)

Involved units

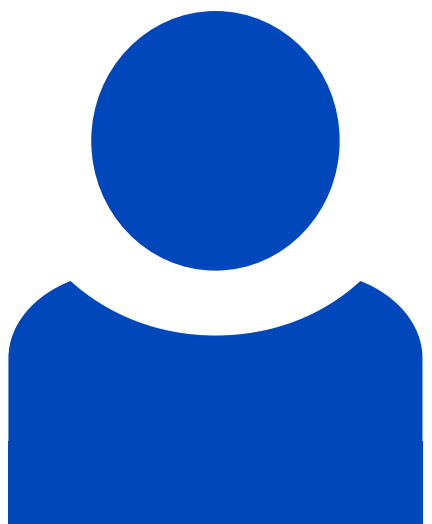
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