APPLICATIONS NOTE

A novel signal processing approach for the detection of copy number variations in the human genome
C.Stamoulis and R.A.Betensky

Structured bioinformatics

A Lasso regression model for the construction of transcriptional networks
D.Zhang, Y.Zhou, W.Qu, M.Deng and C.Zhang

Tissue-specific subnetworks and characteristics of publicly available human protein interaction databases

BRISK—research-oriented storage kit for biology-related data
I.Bertini, D.A.Case, L.Ferella, A.Giachetti and A.Rosato

StochKit2: software for discrete stochastic simulation of biochemical systems with events
J.Y.Semegni, M.Wanulw, R.Gaujoux, G.W.Harkins, A.Gray and D.F.Martin

Simulating systems genetics data with SysGenSIM

DBK—research-oriented annotation pipeline (SHAP)
M.Z.DeMaere, F.M.Lauro, T.Thomas, S.Yau and R.Cavicchiolo

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