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^{OA}Open Access paper.



Cover Colored tiles are playfully suspended on vertical strings to illustrate how Spark, presented in this issue, uses clustering to group genomic regions (tiles) with similar data patterns (colored heatmaps) and facilitates genome-wide data exploration. (Cover illustration by Martin Krzywinski and Cydney Nielsen. [For details, see Nielsen et al., pp. 2262–2269.])