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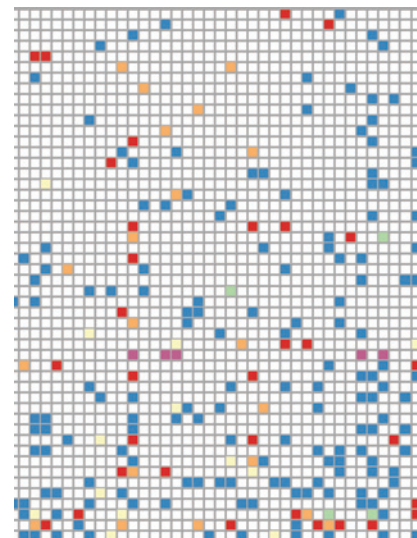
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ABOUT THE COVER

The cover image shows the recurrence of mutations, represented by colored squares, affecting known lymphoma-related genes among a cohort of relapsed and treatment-refractory DLBCL tumors subjected to exome sequencing. Each column corresponds to an individual patient with transformed lymphomas positioned on the left and *de novo* DLBCL toward the right. Each row is a gene, with the most frequently mutated genes (TP53, MLL2, CREBBP, MLL3, and STAT6) towards the top. Protein-altering mutations of different categories are indicated by color. Noncoding mutations are omitted, with the exception of mutations affecting the microRNA gene MIR142 (purple). For details see the article by Morin and colleagues on page 2290 of this issue.



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