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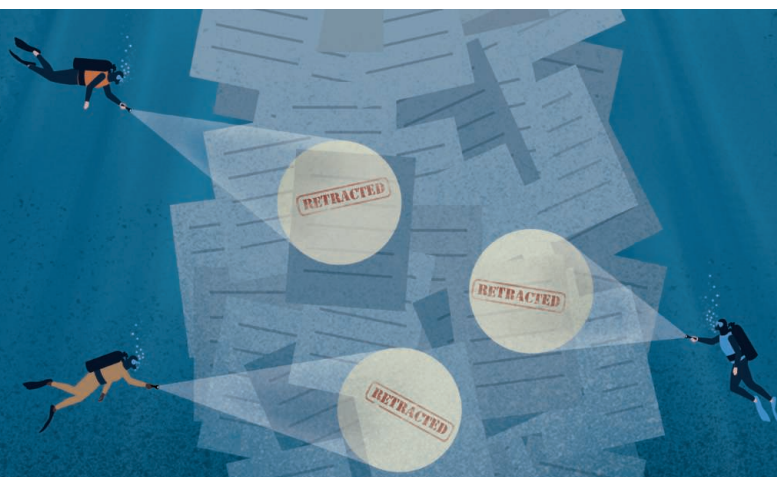
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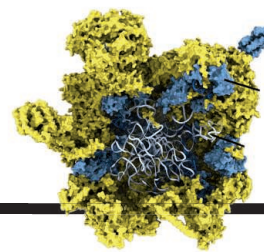
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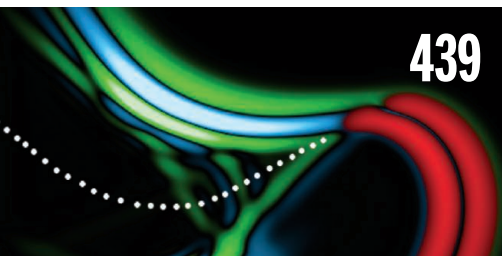
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Artist's rendering of chromatin (coiled structures) and DNA regulatory elements (yellow clusters) inside a cell's nucleus. The Cancer Genome Atlas (TCGA) Analysis Network profiled the

complex accessible chromatin landscape in 410 cancer samples from 23 cancer types, substantially expanding the catalog of known DNA regulatory elements in cancer genomes and shedding light on how genetic mutations and epigenetic regulators work together to promote tumor development. See pages 401 and 420. *Image: V. Altounian/Science*

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