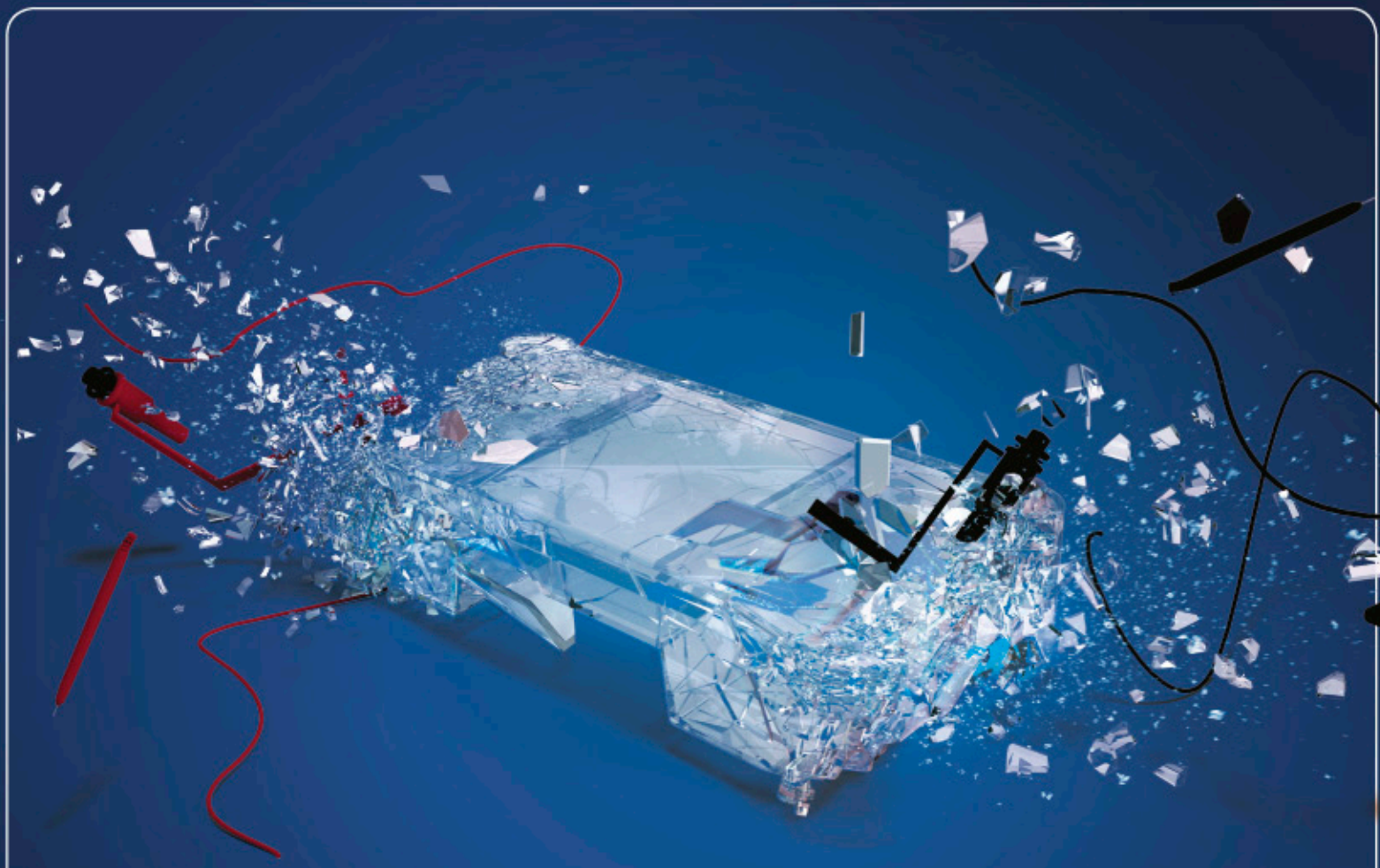

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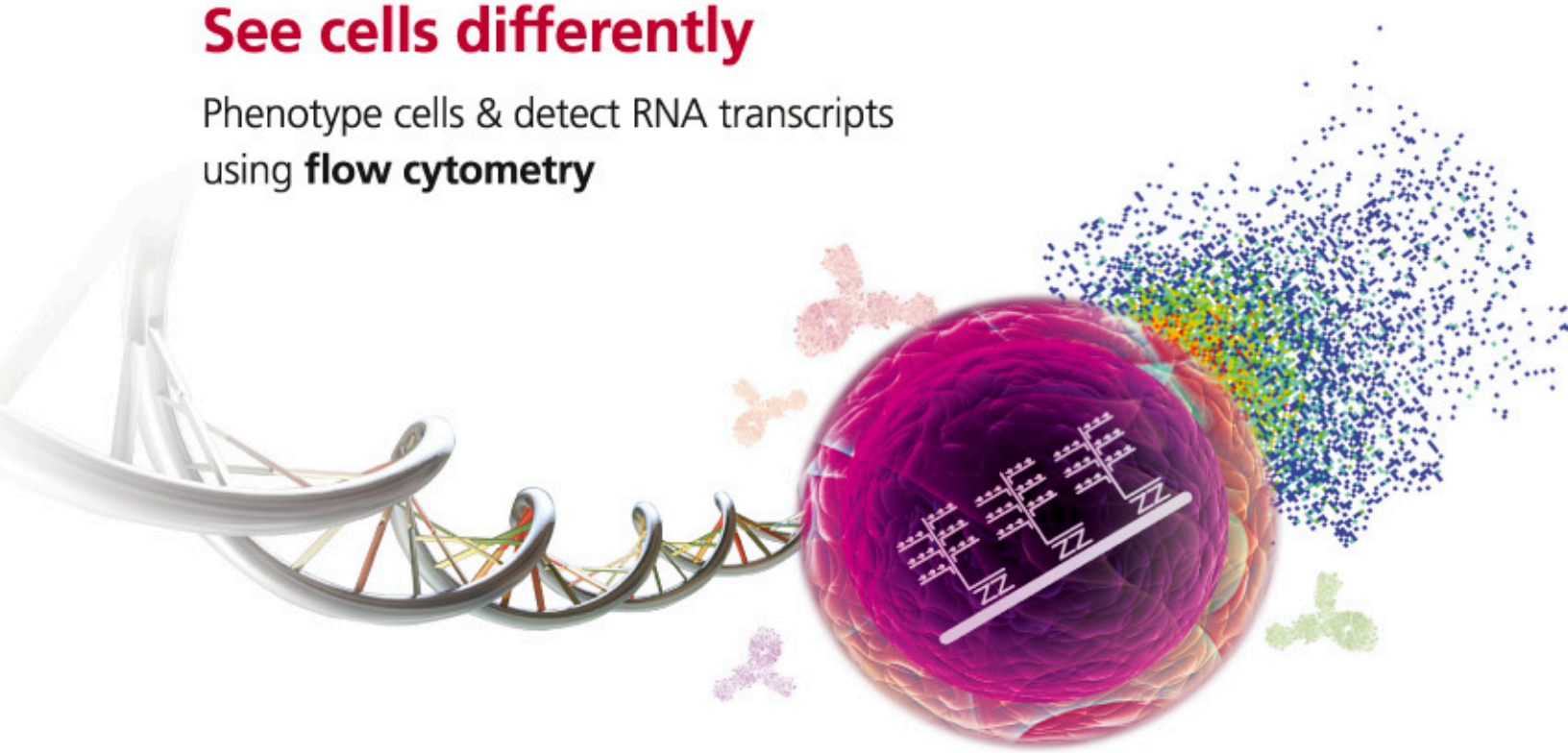

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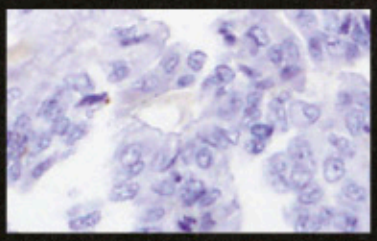
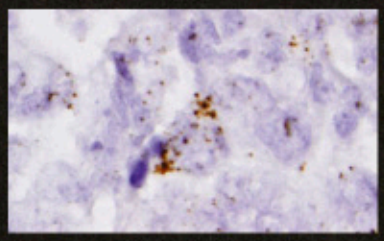


***IHC** fails to detect Napsin A expression in lung adenocarcinoma*



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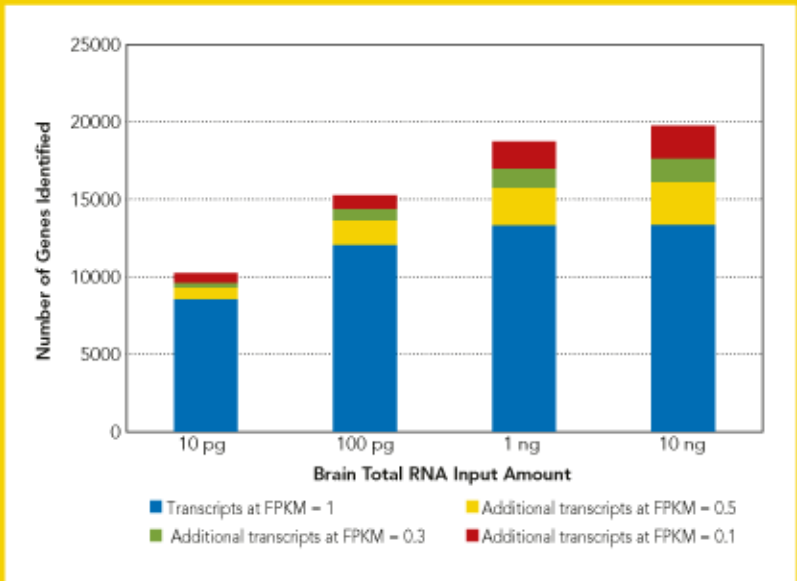

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1. Frommer M, McDonald C, Millar DS et al (1992). A genomic sequencing protocol that yields a positive display of 5-methylcytosine residues in individual DNA strands. Proc.Nat.Acad.Sci., USA 89: 1827-1831.
2. Lister R, Pelizzola M, Dowen RH, et al (2009). Human DNA methylomes at base resolution show widespread epigenomic differences. Nature. 19;462(7271): 315-22.

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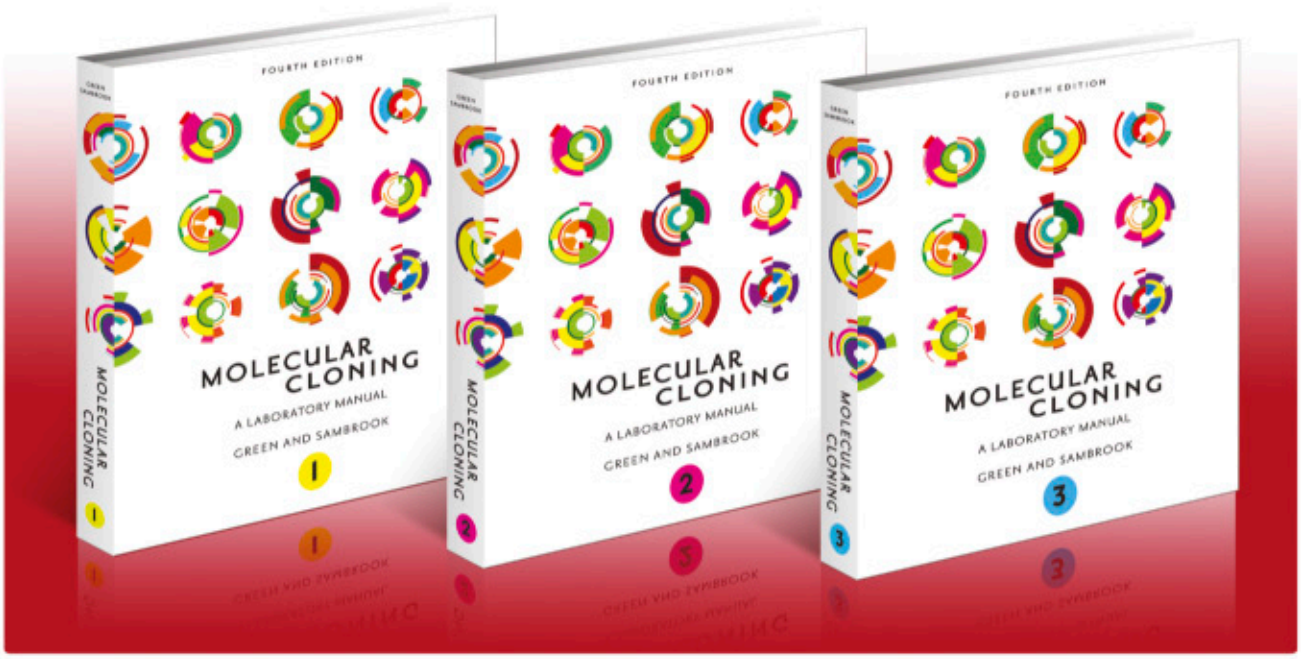

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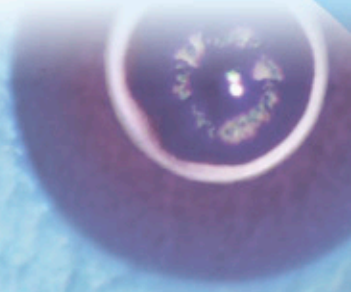

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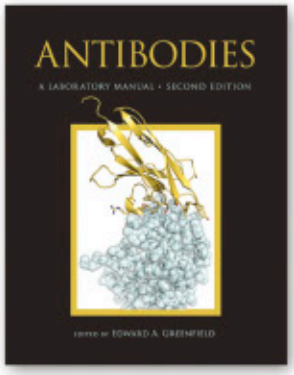

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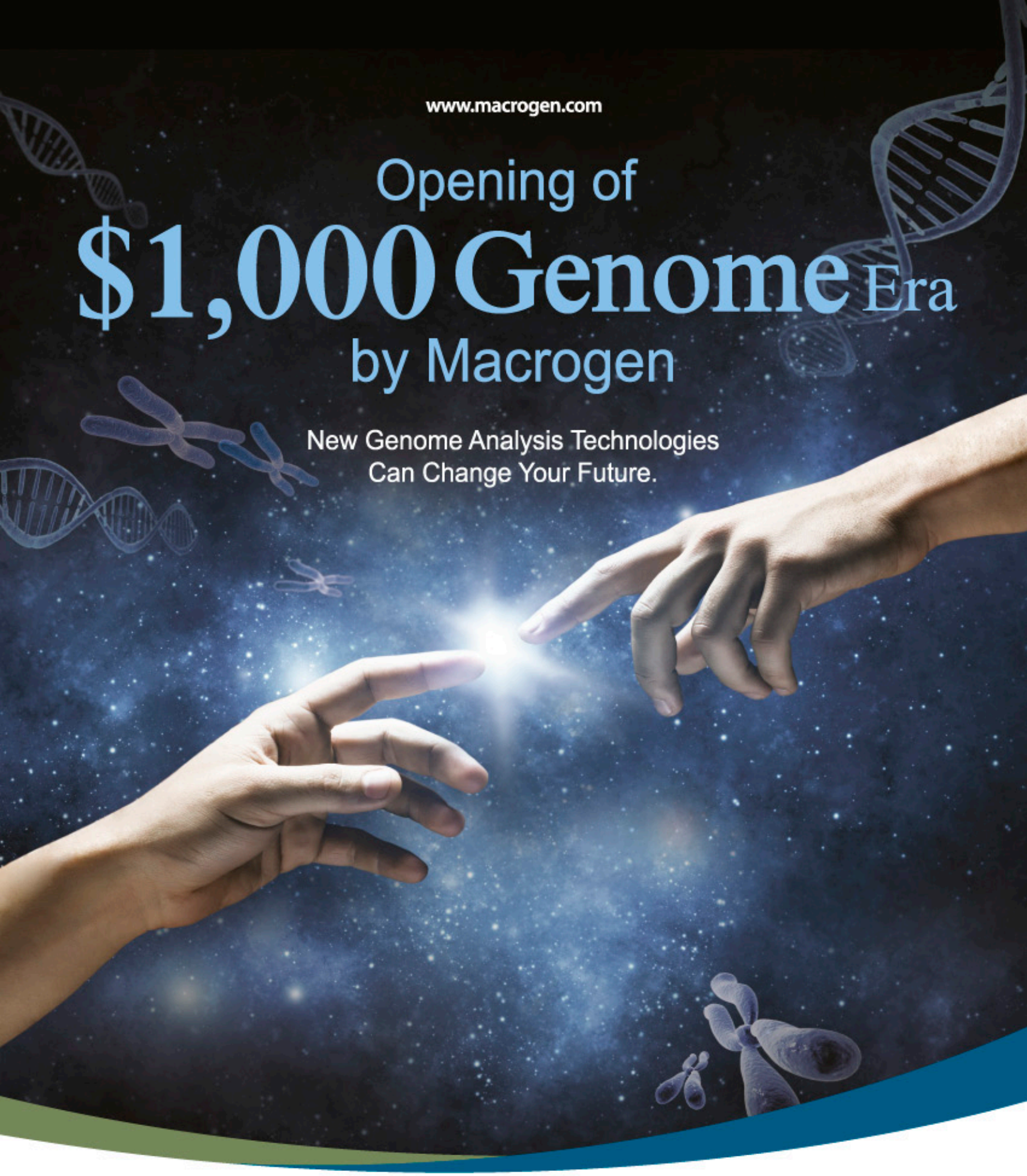

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