Getting Polymers to Tell a Story: Scaling up DNA Data Storage and Functionality

Luis Ceze and Karin Strauss

Molecular Information Systems Lab

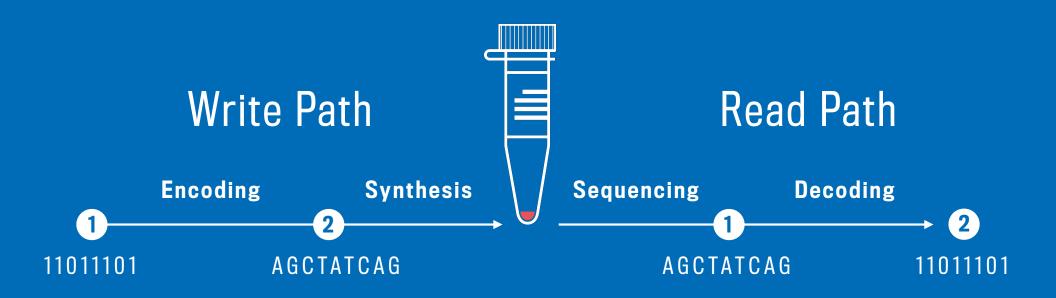


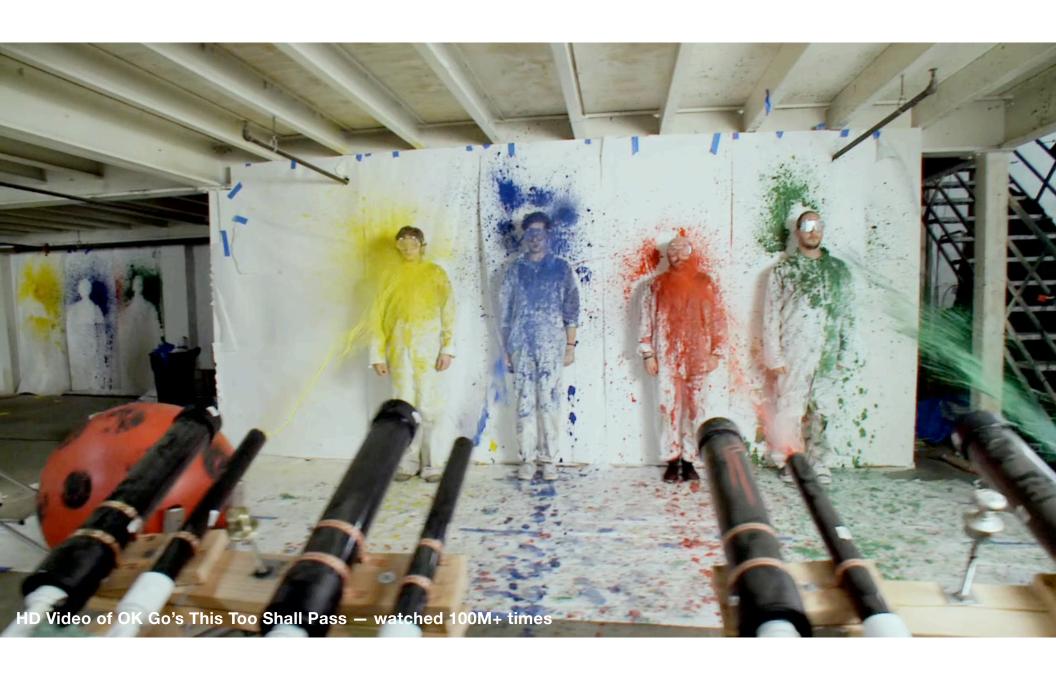


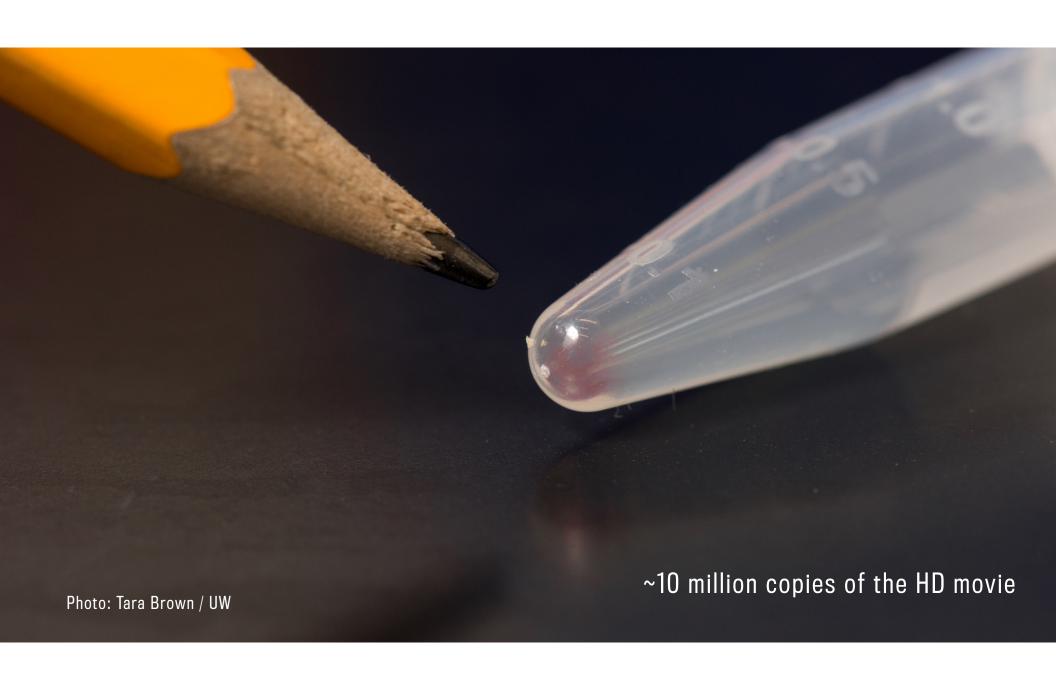
joint work with Georg Seelig, Doug Carmean, Sergey Yekhanin, Lee Organick, Yuan-Jyue Chen, Bichlien Nguyen, Chris Takahashi, Ashley Stephenson, Pranav Vaid, Sharon Newmann, Cyrus Rashtchian, Miklos Racz, Siena Ang, David Ward, Randolph Lopez, Max Willsey, Kendall Stewart, James Bornholt, Rob Carlson, Hsing-Yeh Parker.

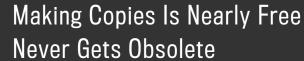
LoC DSA, September 2018

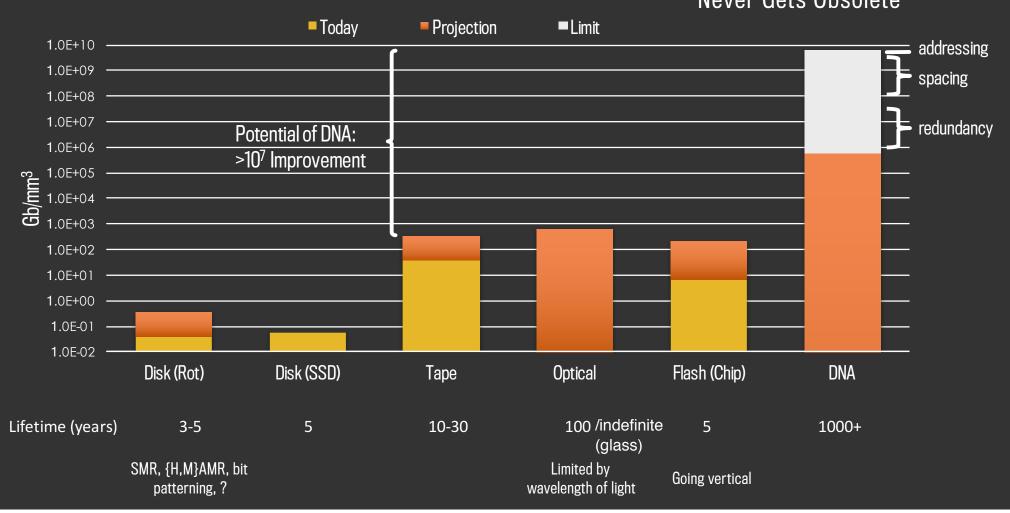


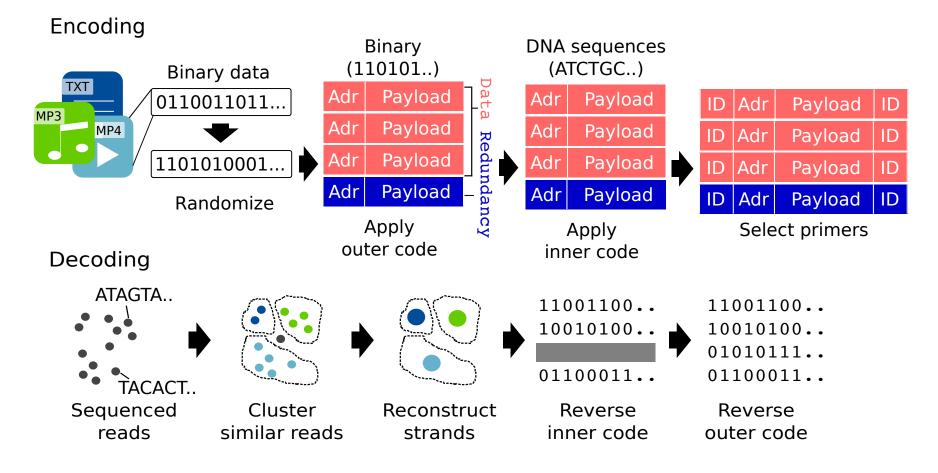




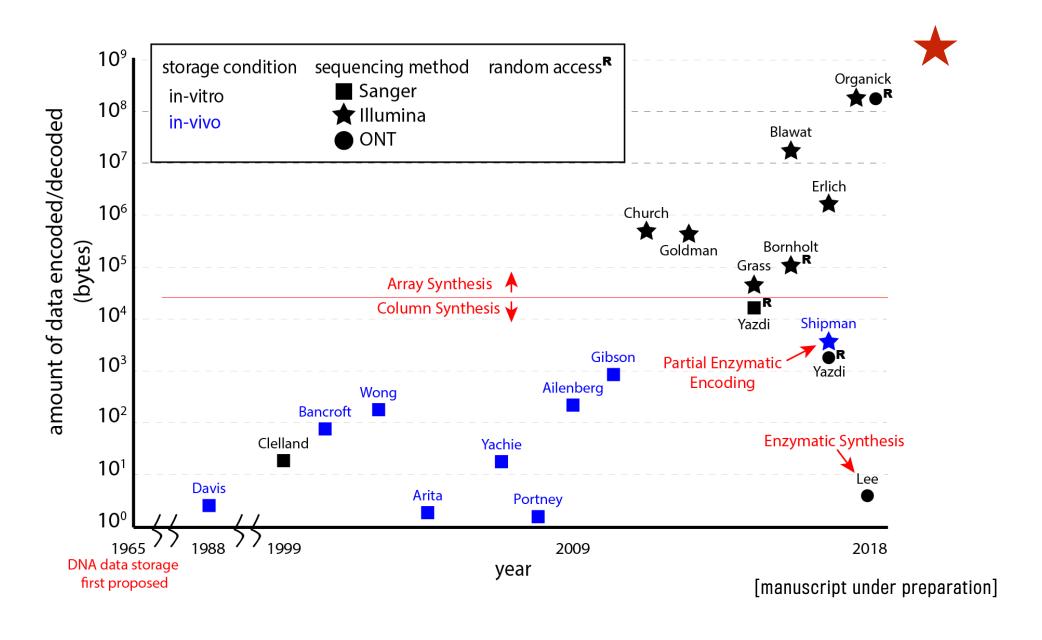








[Nature Biotechnology'18, NIPS'17]



Challenges

to mainstream availability

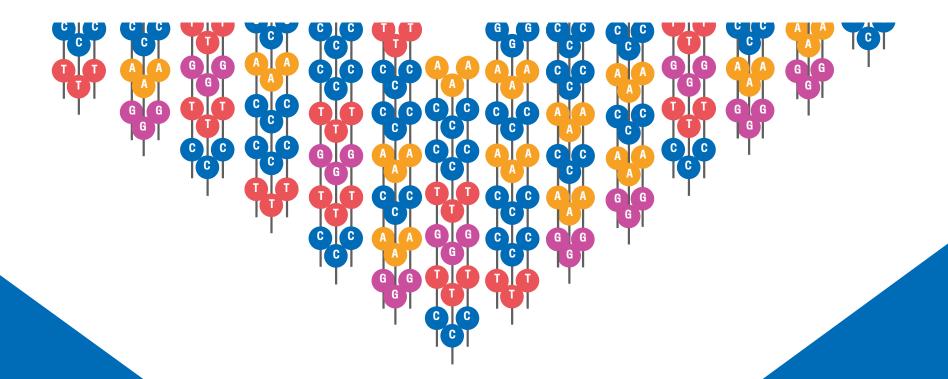
Bandwidth goal is O(TB/s), today writes are at O(KB/s)

Very high throughput and low cost writing and reading

Large-scale fluidics automation

Scalable DNA physical organization and retrieval

Computational costs

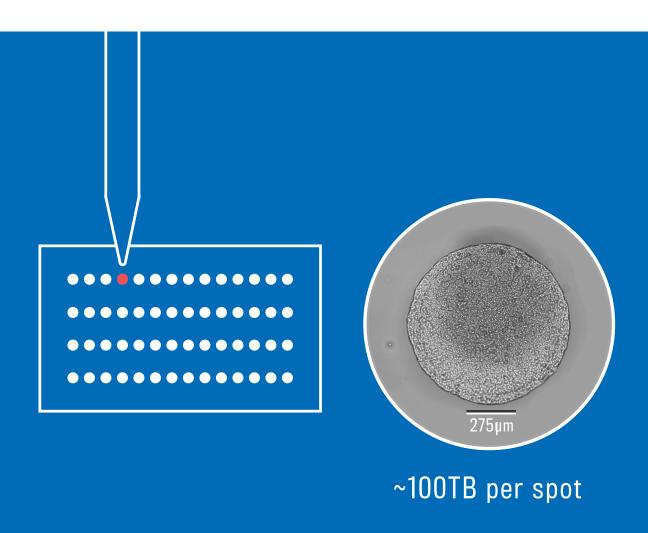


Large array DNA synthesis

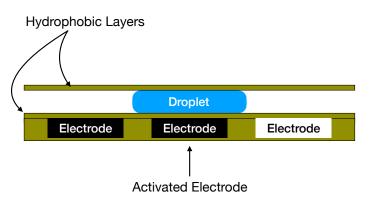
Each spot grows many copies of a given sequence.

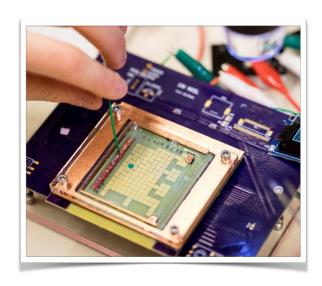
Many spots.



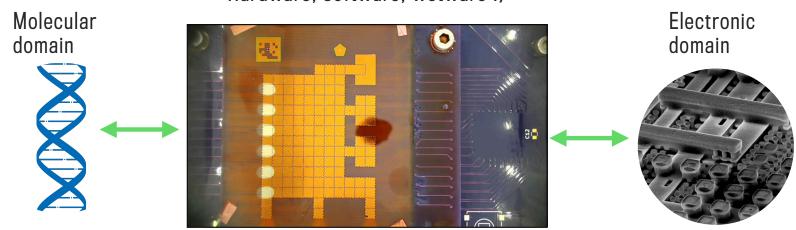


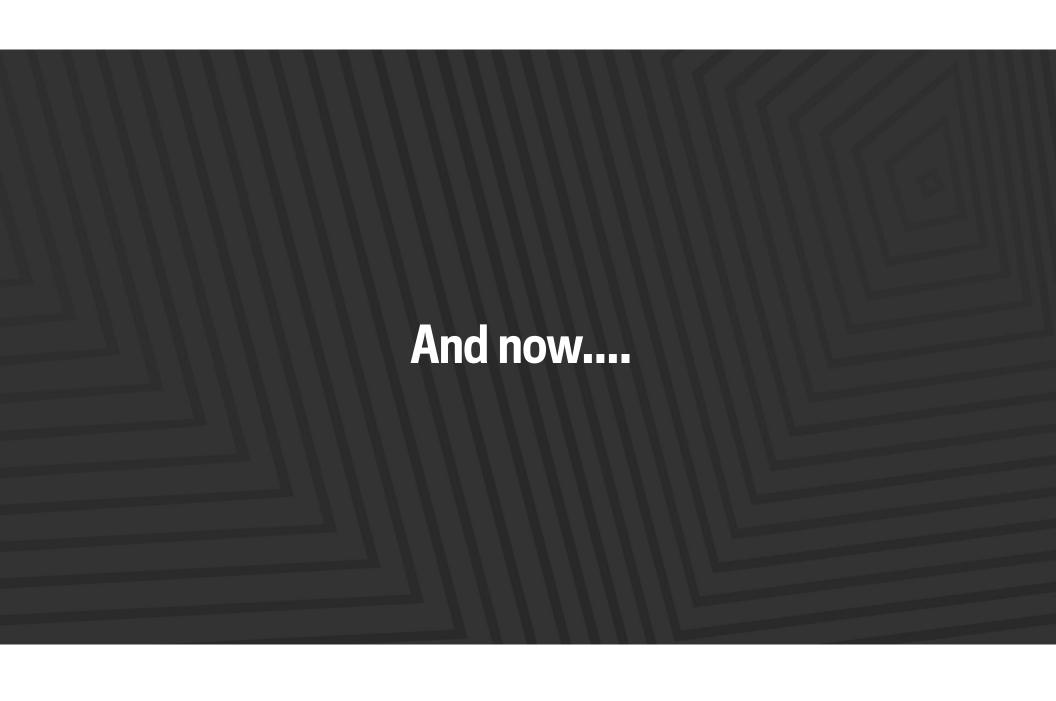
Digital microfluidics

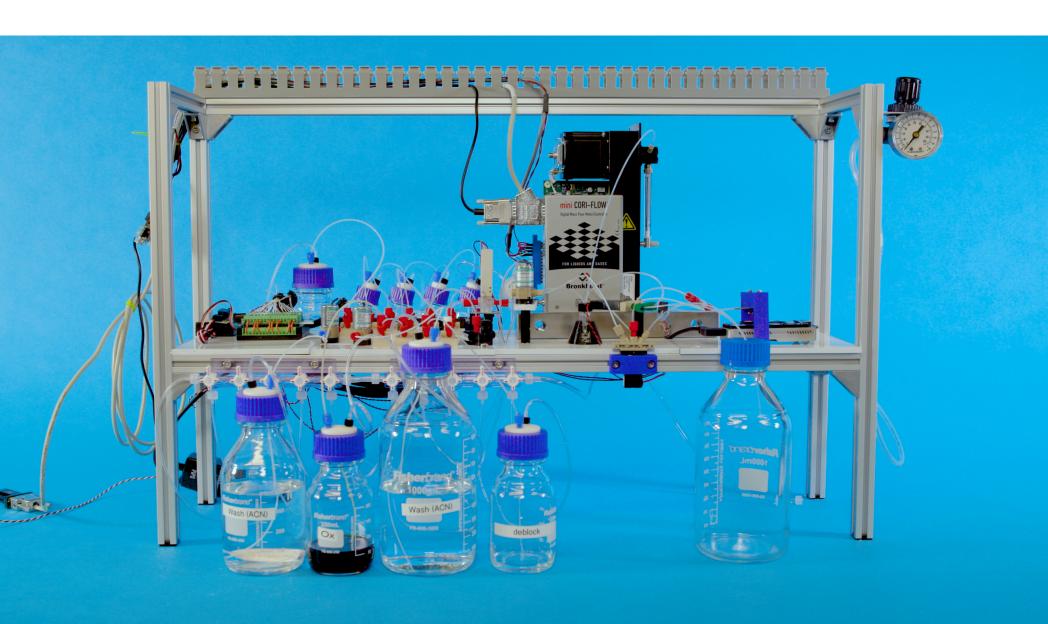




Hardware, software, wetware:)

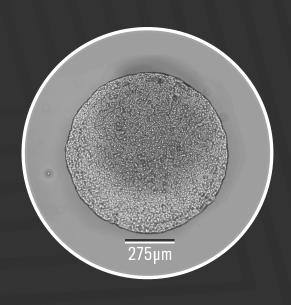








DNA "computing" in the age of big data



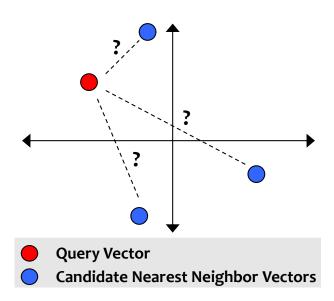
~100TB per spot

If DNA data storage succeeds, what if we could process data directly in DNA?

Extremely parallel and energy efficient

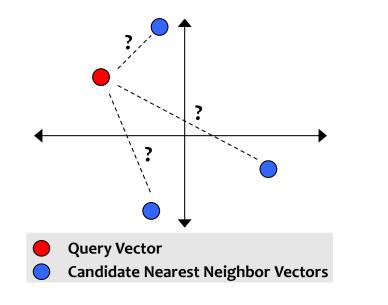
Content-based media search

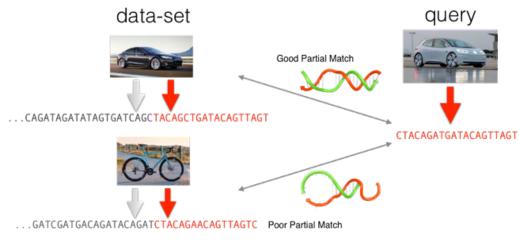
Extract feature vector, search in a high-dimensional space

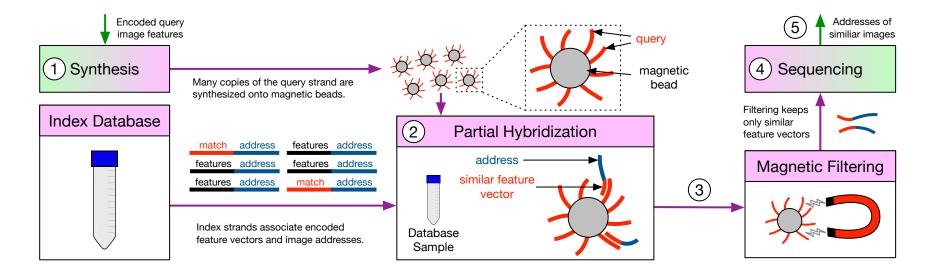


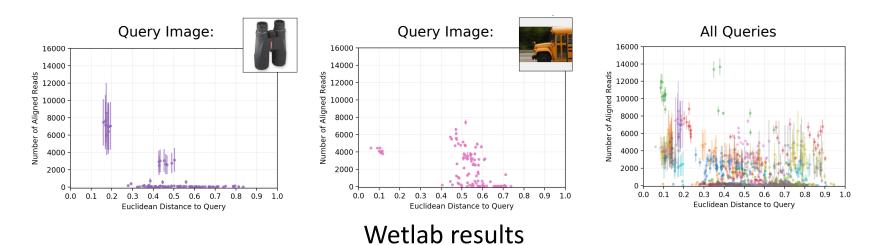
Content-based media search ... in DNA

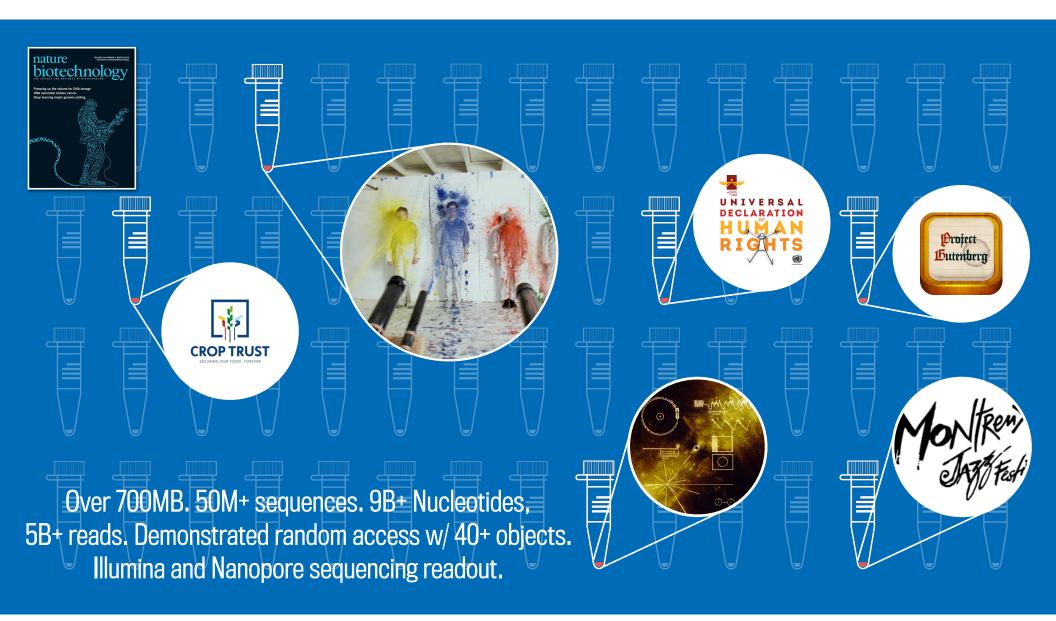
Map features vector do DNA such that molecules mapping to similar vectors "stick"

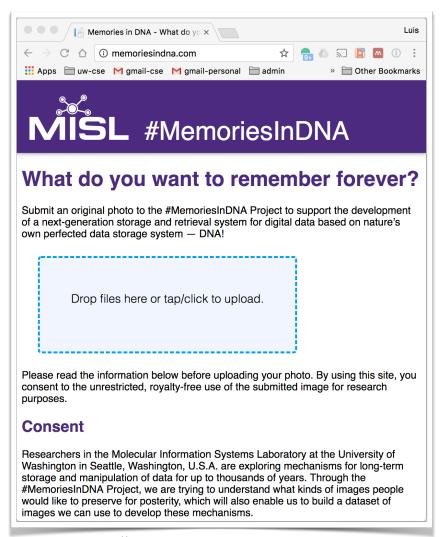












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