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TGTCTTTGATTCCTG**

**TATGTCGCAGTATCT
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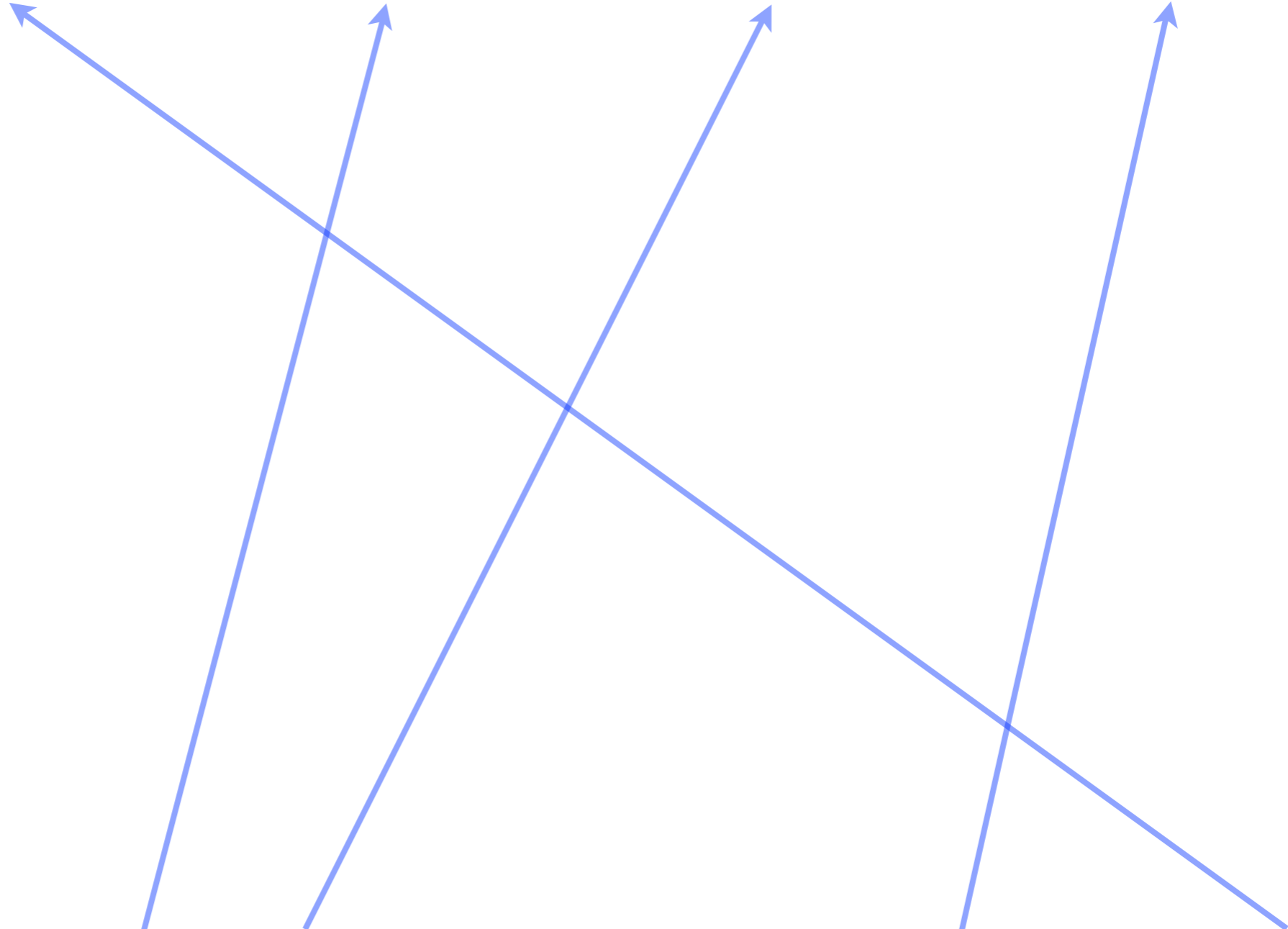
**CACCCTATGTCGCAG
TGGAGCCGGAGCACC
GCATTGCGAGACGCT**

**GAGACGCTGGAGCCG
CGCTGGAGCCGGAGC
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Reads

Your genome

CGTCTGGGGGTATGCACGCGATAGCATTGCGAGACGCTGGAGCCGGAGCACCCTATGTCGCAGTATCTGTCCTTTGATTCCTG



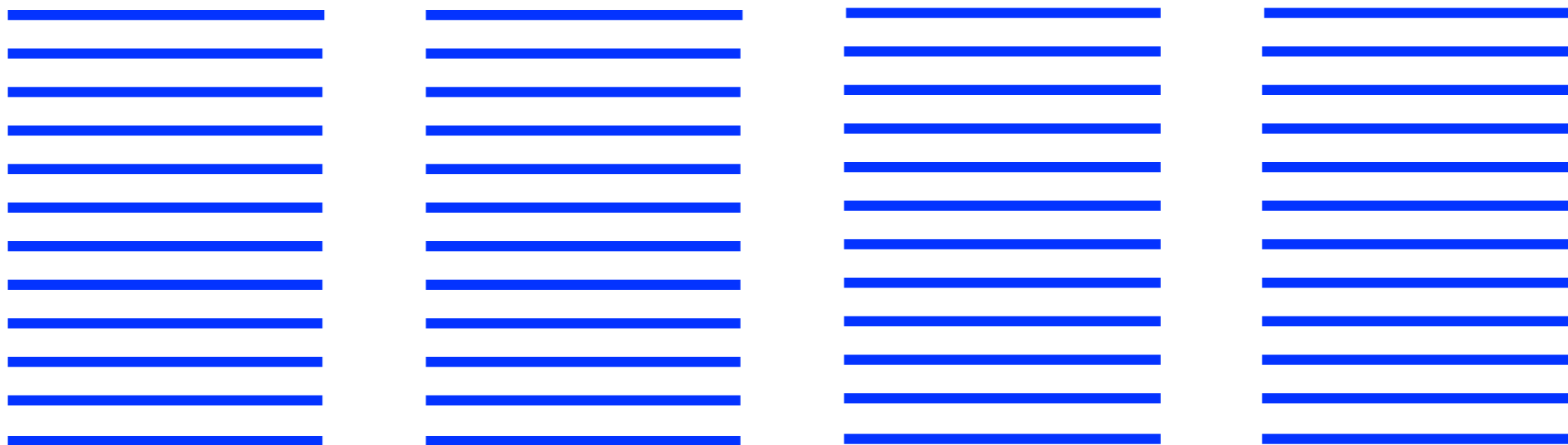
Reads

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GTATGCACGCGATAG	ACCTACGTTCAATAT	TATTTATCGCACCTA	CCACTCACGGGAGCT
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GACGCTGGAGCCGGA	GCACCCTATGTCGCA	GTATCTGTCTTTGAT	CCTCATCCTATTATT
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CACGGGAGCTCTCCA	TGCATTTGGTATTTT	CGTCTGGGGGGTATG	CACGCGATAGCATTG

Your genome

CGTCTGGGGGGTATGCACGCGATAGCATTGCGAGACGCTGGAGCCGGAGCACCCTATGTCGCAGTATCTGTCTTTGATTCCTG

Reads



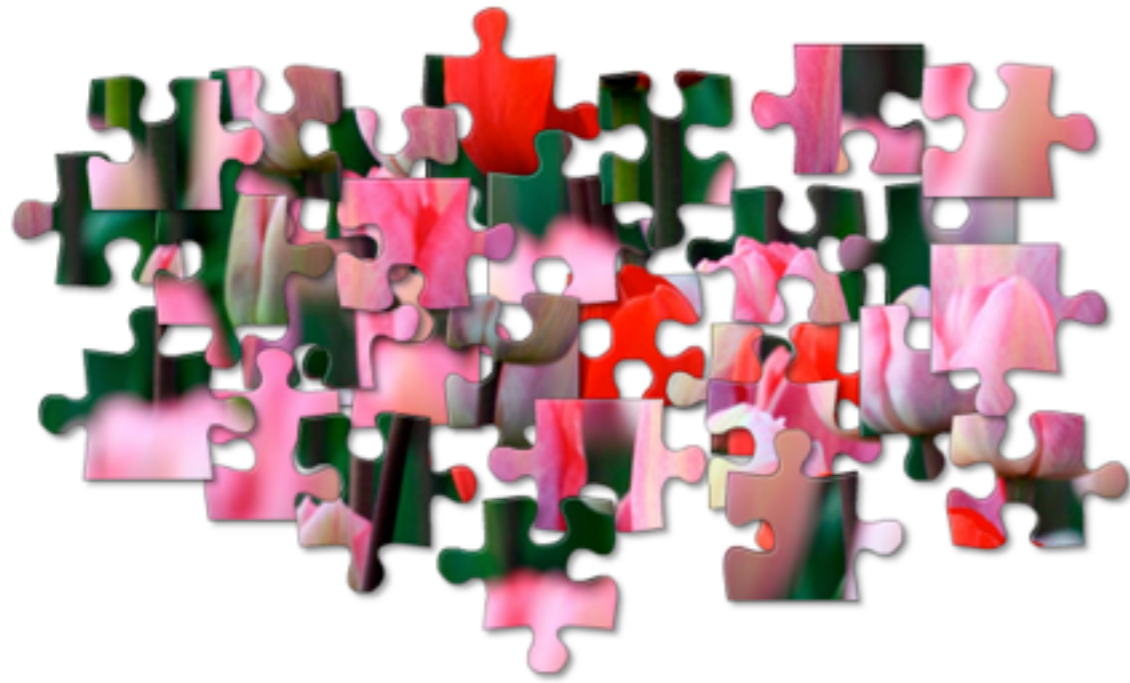
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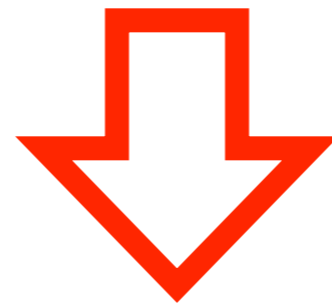
Your genome



100,000,000 nt



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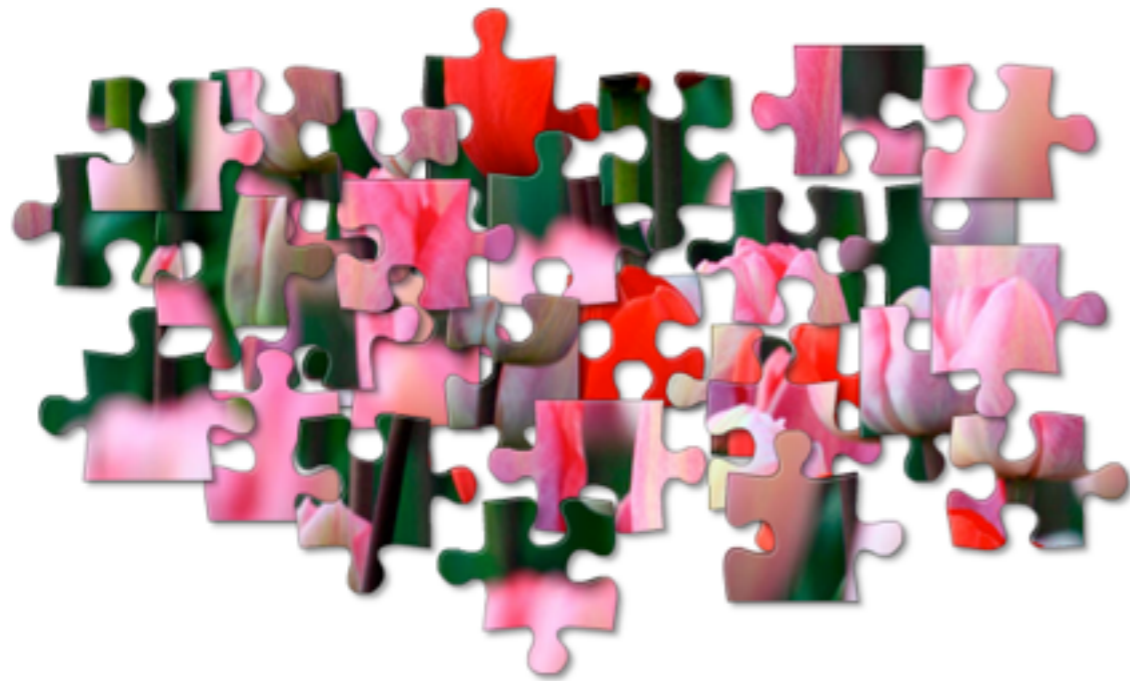
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Unrelated humans have genomes that are 99.8-99.9% similar

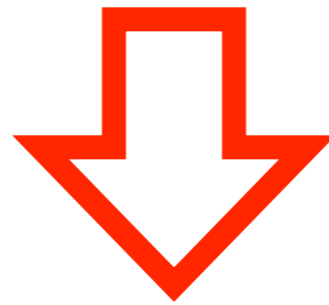
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Alignment



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Assembly

