

## Reading and Writing Genomes.

George Church\*

Genetics Dept., Harvard Medical School, Boston, MA 02446 USA

Nucleic acid synthesis and sequencing technologies are intimately connected. In the past 4-5 years our team has helped develop 'second-generation' sequencing by a radical shift from electrophoresis to a system of multiplex fluidic cycles and imaging [1] bringing costs down by 10,000 fold to 25 Mbp/\$. We have also developed an analogous second generation approach to synthesis of genes from oligonucleotides on chips [2] such that DNA can be made for 30 Kbp/\$ (and much less if one counts combinatorial strategies). Finally we have integrated these with a third class of technology to program cells and viruses and greatly accelerate and monitor their lab evolution [3] -- Multiplexed Automated Genome Engineering (MAGE). We are constructing a (4.7 Mbp) bacterial genome with a new translational code that should result in resistance to multiple viruses (or all viruses). We have a related project to construct a mirror-image cell based on *E.coli* in vitro protein synthesis (a replicating 110 kbp genome [4]) resistant to most existing enzymes, drugs, predators. We have championed methods for safety and surveillance in synthetic biology.

**References** (see also <http://arep.med.harvard.edu/>)

1. Shendure, J, Porreca, GJ, Reppas, NB, Lin, X, McCutcheon, JP, Rosenbaum, AM, Wang, MD, Zhang, K, Mitra, RD, Church, GM (2005) **Accurate Multiplex Polony Sequencing of an Evolved Bacterial Genome**. *Science* 309(5741):1728-32.
2. Tian J, Gong H, Sheng N, Zhou X, Gulari E, Gao X, & Church GM (2004) **Accurate Multiplex Gene Synthesis from Programmable DNA Chips**. *Nature* 432: 1050-4.
3. Wang HH, Isaacs FJ, Forest CR, Sun ZZ, Xu G, Church GM (2009) **Multiplexed Genome Engineering and Directed Evolution**. *Nature* in press.
4. Forster, AC & Church, GM (2006) **Toward Synthesis of a Minimal Cell**. *Nature-EMBO-Molecular Systems Biology* 2 doi:10.1038/msb4100090