Summary

- "Sanger" sequencing widely used up through 2006 or 2007, including for the human genome project.

- Won Sanger his second Nobel prize.

- Lander-Waterman statistics estimate the number of islands you will get for a given coverage.

  - Used as a way to guess how much sequencing you need to do for a given technology and genome size.

  - Often hard in practice to guess the genome size g before you've sequenced it.