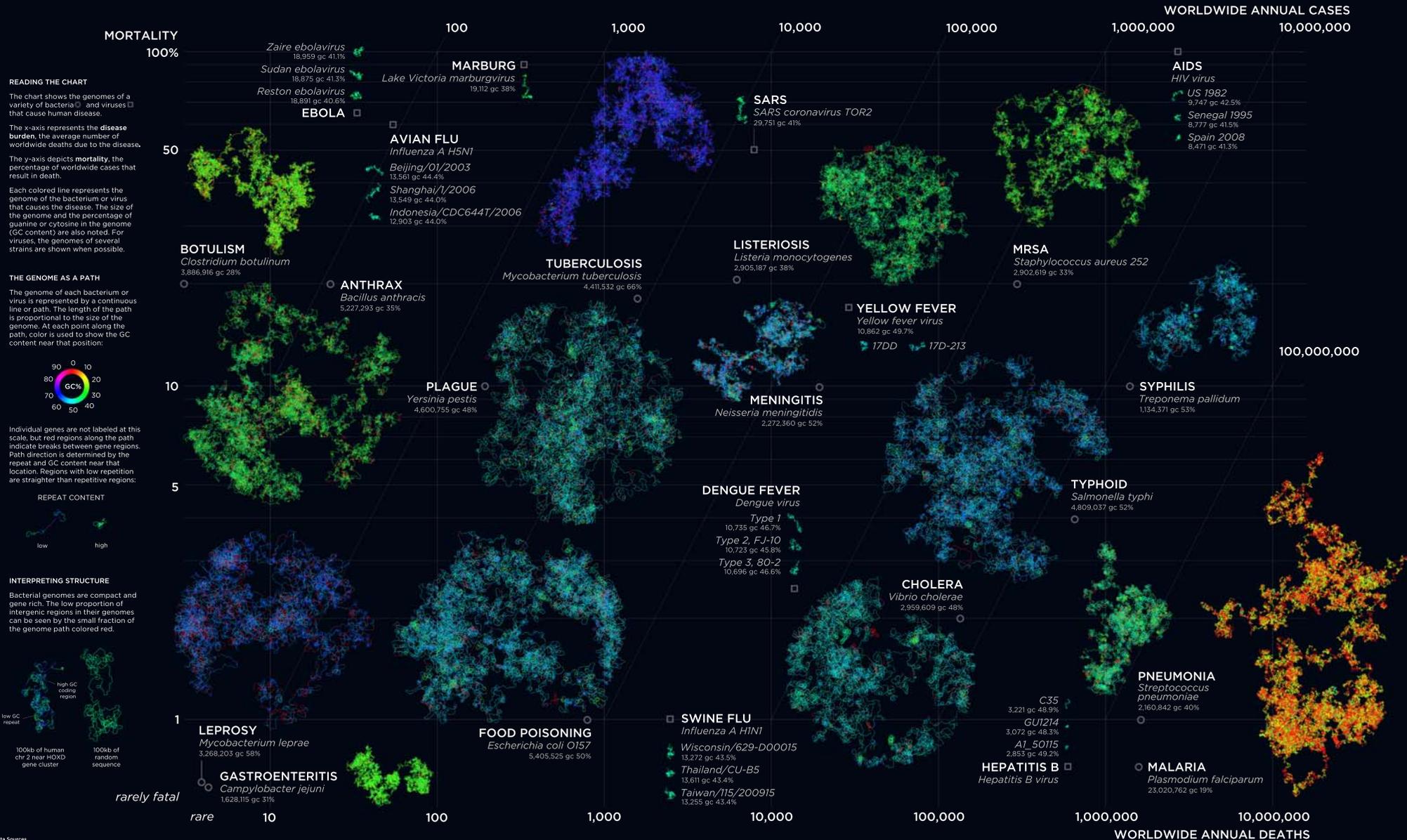


THE DEADLY GENOMES

GENOME STRUCTURE AND SIZE OF HARMFUL BACTERIA AND VIRUSES



Data Sources
Incidence and mortality data were obtained from the World Health Organization (<http://www.who.int>). In cases where mortality is variable (e.g. mortality of SARS ranges from 1-50%, depending on age), the highest value is reported. For some outbreak diseases, such as Ebola, statistics were averaged over several years. The distribution of cases and mortality across world regions for most diseases shown here is not uniform. For example, haemorrhagic fevers like Marburg, Dengue and Ebola are largely limited to Africa. For more information about this poster, the data and to download genome paths for each disease, visit <http://mikeweb.bcgsc.ca/deadlygenomes>.

Credits
Martin Krzywinski, Sydney Nielsen, Ian Bosdet (Canada's Michael Smith Genome Sciences Center), Jonathan Corum (136t)