GENETICS

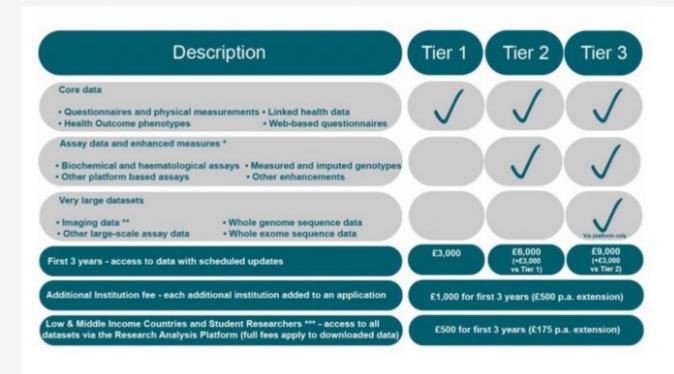
HAPMAP

21.03.2022

I find it increasingly difficult to get the haplotype structure of a given gene. While the are <u>patches to the human genome</u> at UCSC, this isn't the same like good old hapmap.org by the <u>HapMap consortium that has been retired</u> back in 2016.

A recent computer security audit has revealed security flaws in the legacy HapMap site that require NCBI to take it down immediately. We regret the inconvenience, but we are required to do this. That said, NCBI was planning to decommission this site in the near future anyway (although not quite so suddenly), as the 1,000 genomes (1KG) project has established itself as a research standard for population genetics and genomics. NCBI has observed a decline in usage of the HapMap dataset and website with its available resources over the past five years and it has come to the end of its useful life.

It seems that also its successor, the 1000 genomes, will be retired during the next month. Unfortunately, the phasing server in Michigan does not allow any logon for whatever reason, UK biobank is too expensive so haplotypes basically vanished in the nirvana.



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