

GENETICS, SOFTWARE

MY LINUX BOX

10.07.2007

When starting again analysis of pooled DNA samples – this time on the Affymetrix platform – I discovered not only the much improved “[Genotyping Console](#)” but also that my earlier [Genepool](#) binaries do not work anymore under [VMWARE/Ubuntu](#). I therefore tried [QEMU/-KNOPPIX](#) which is slow but runs reliable.

Unfortunately Genepool ends with a partitioning error; I have already asked the authors if the new 500K chip design may be responsible for that error (as it does not split Nsp and Sty chips anymore). To be continued ...

