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**Homework 1**

**Important dates:**

Submit report /presentation files before 6 November 2018

* Download data from <http://firebrowse.org/> and choose T1 PAAD (Pancreatic) and T2 LUAD (lung).
* Convert from MAF to PLINK using one of the following step
	+ I: Either you can convert maf to vcf and from vcf to PLINK .ped and. map file using tool available at https://github.com/mskcc/vcf2maf
	+ II : Copy the SNPs information from MAF and create .ped.
* PLINK to data analysis and perform the comparison
* GWAS comparision of T1 vs T2
* post-GWAS with DEPICT
* GenABEL / PLINK’ difference (software point of view)