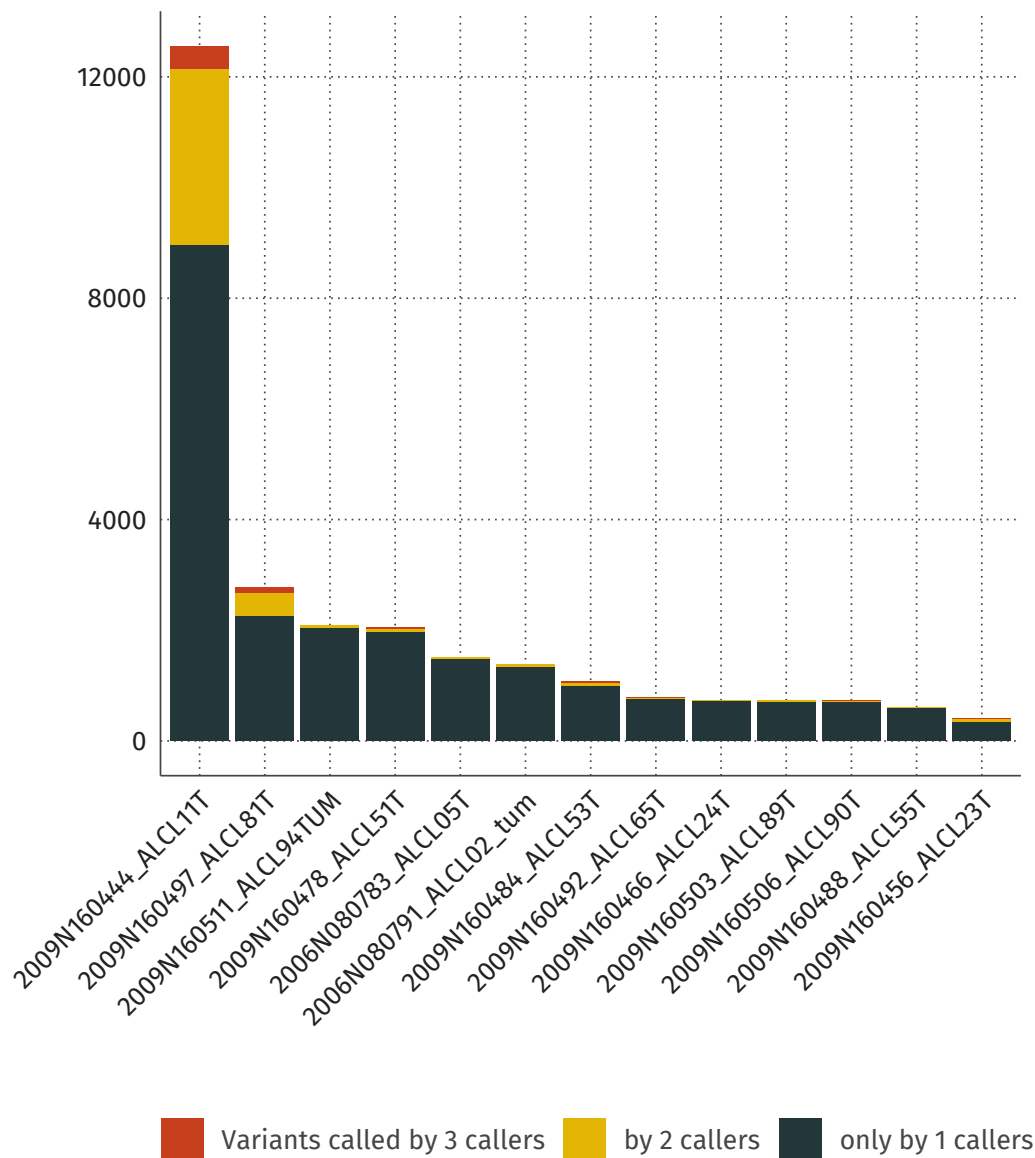
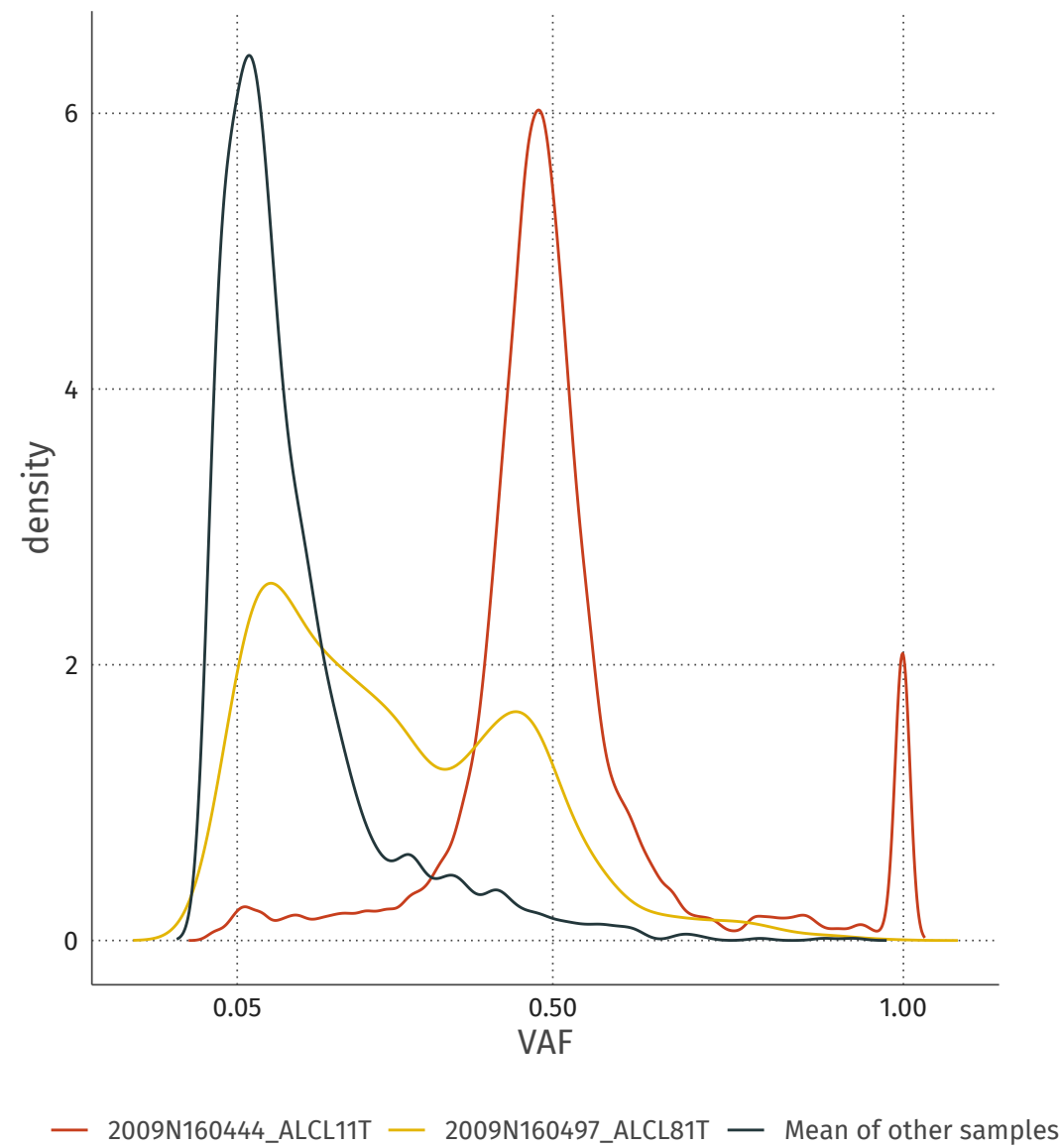


Number of uniques mutations



VAF of samples 11 and 81 compared to others



Incidence moyenne dans la population normale des variants detectés (gnomAD)

