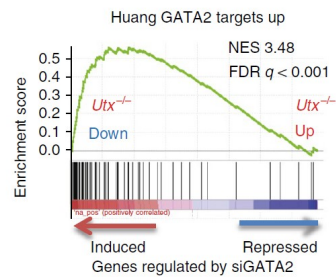
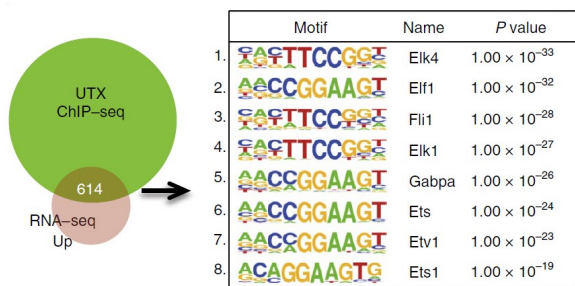
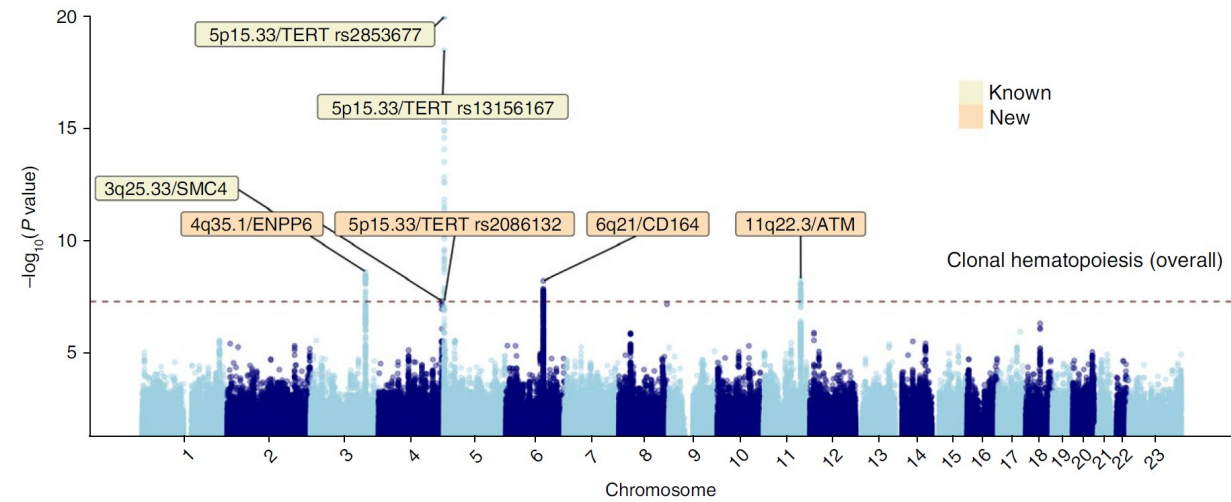
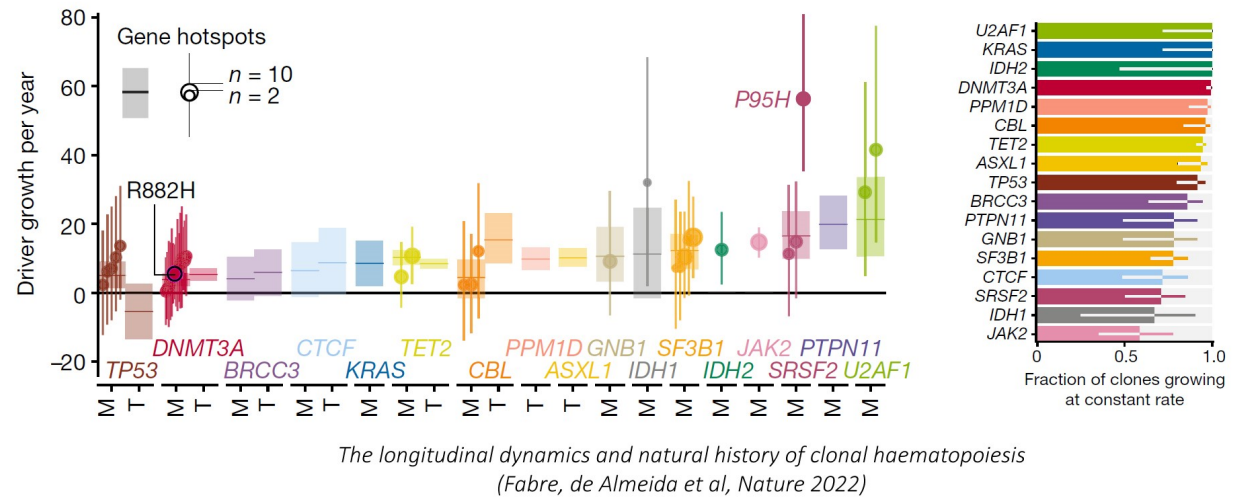


Multiparameter prediction of myeloid neoplasia risk
(Gu et al, Nature Genetics 2023)



(UTX-mediated enhancer and chromatin remodeling suppresses myeloid leukemogenesis through noncatalytic inverse regulation of ETS and GATA programs (Gozdecka et al, Nature Genetics 2018)



Genome-wide analyses of 200,453 individuals yield new insights into the causes and consequences of clonal hematopoiesis
(Kar, Quiros et al, Nature Genetics 2022)