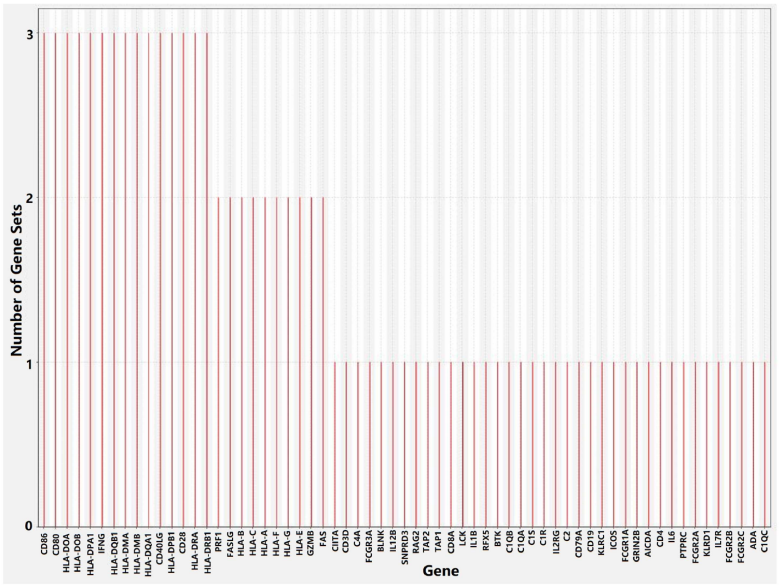
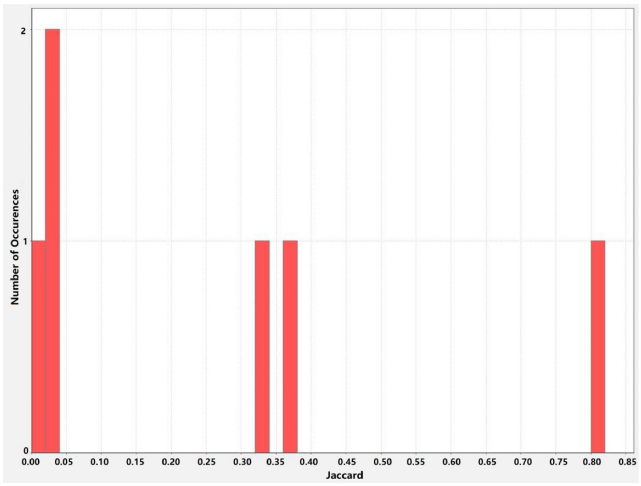


A) Overlapping genes in Gal-9-related pathways.



B) The leading edge analysis.