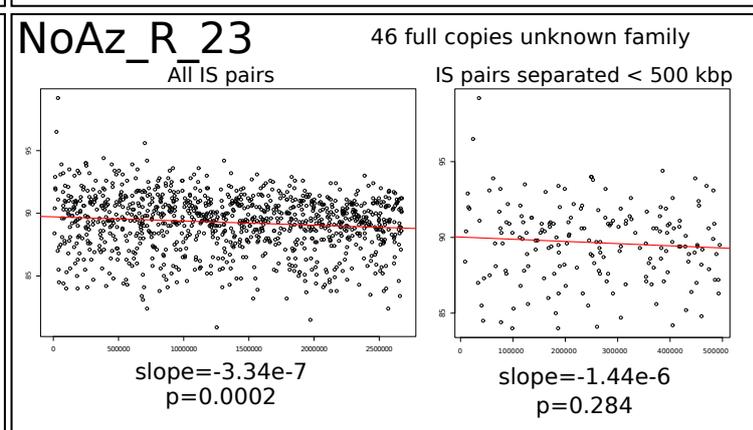
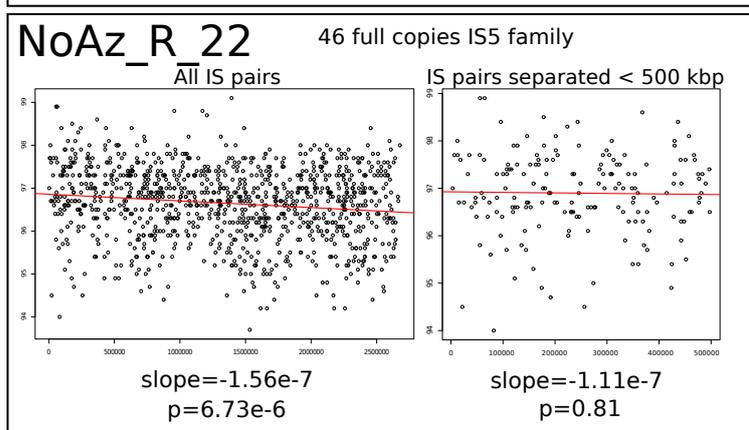
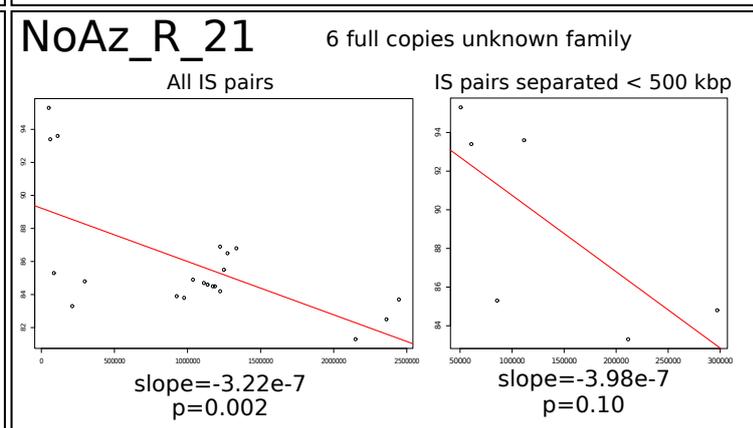
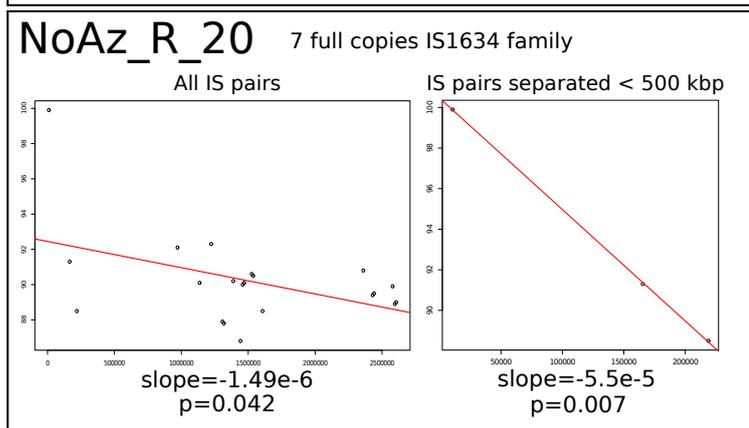
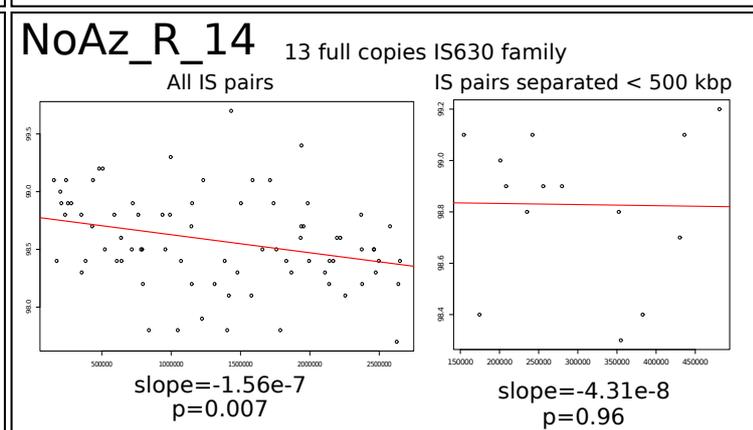
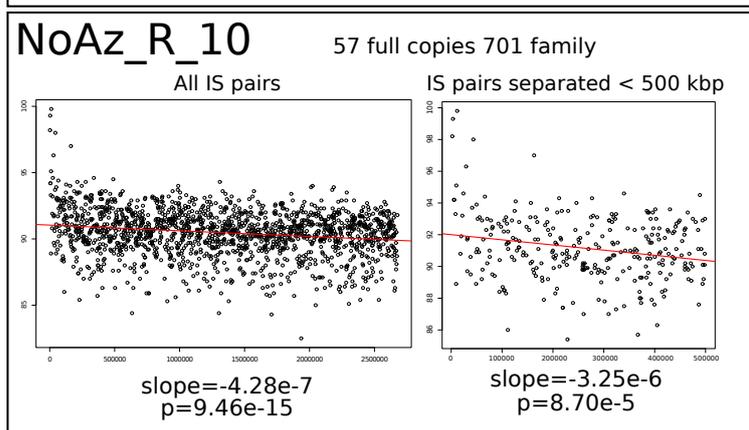
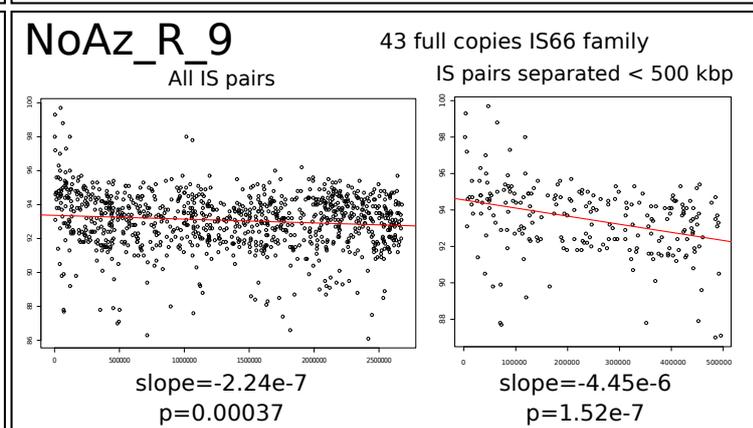
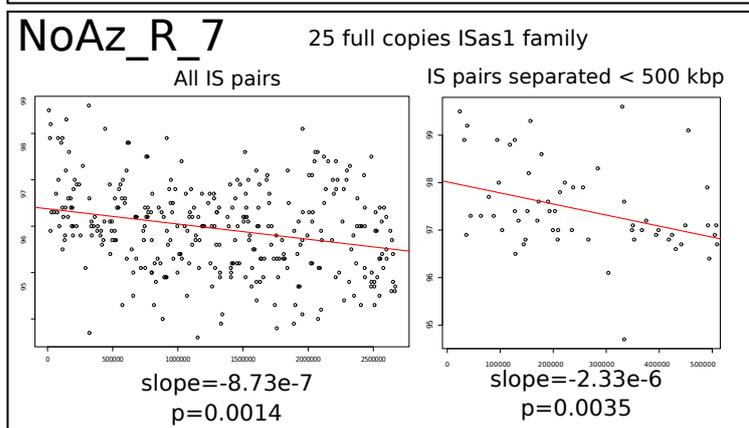
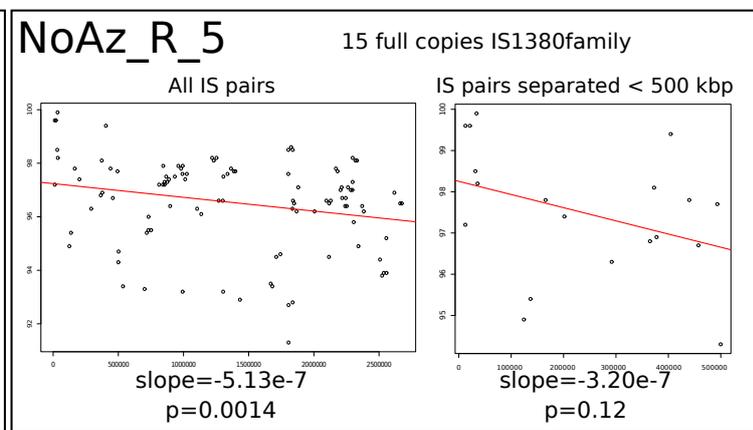
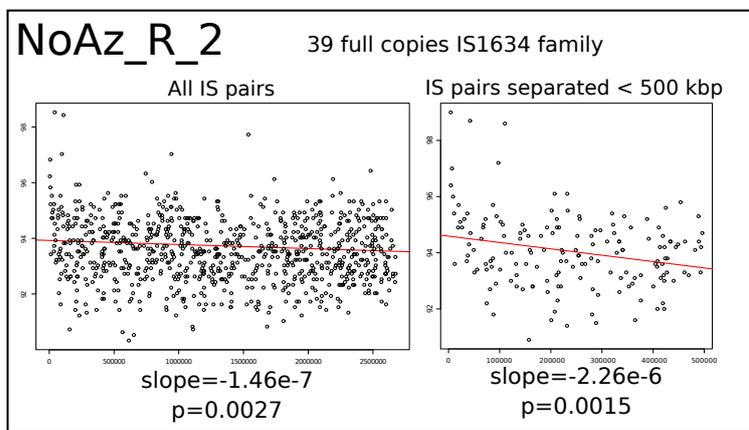


# Repeat alignment identities versus distance on chromosome for some repeats in NoAZ

alignment, % identical positions



Repeat distance on chromosome (nts)