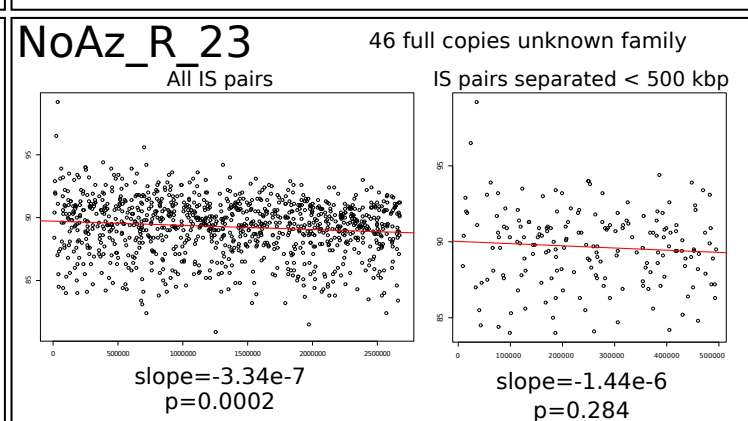
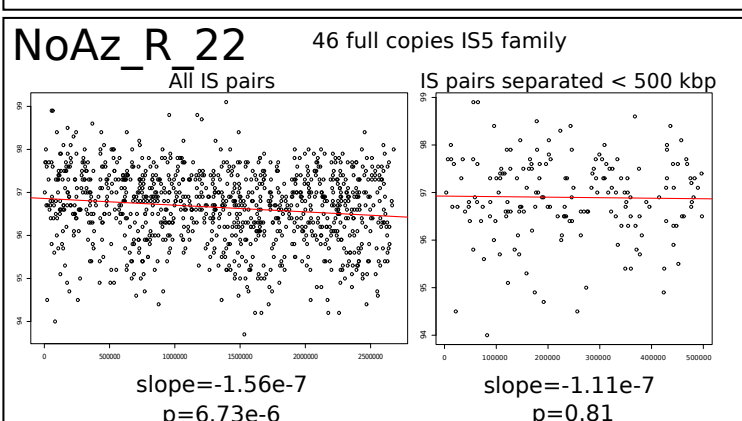
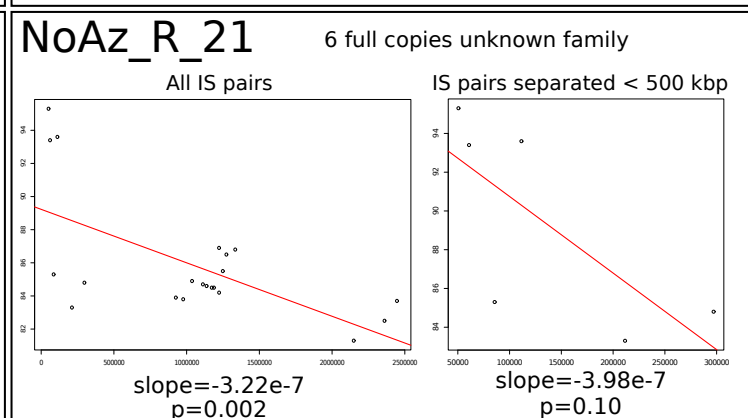
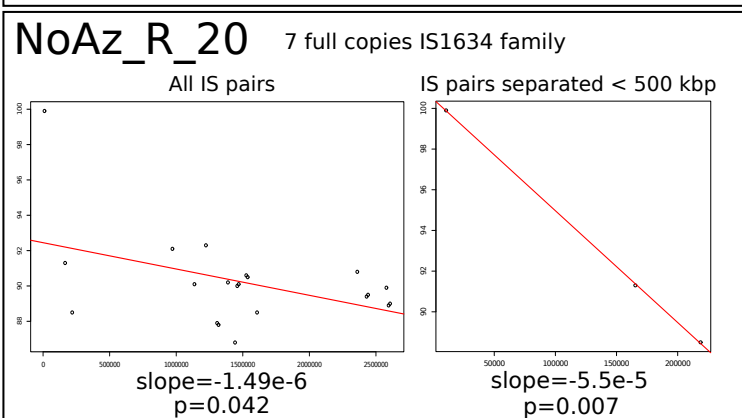
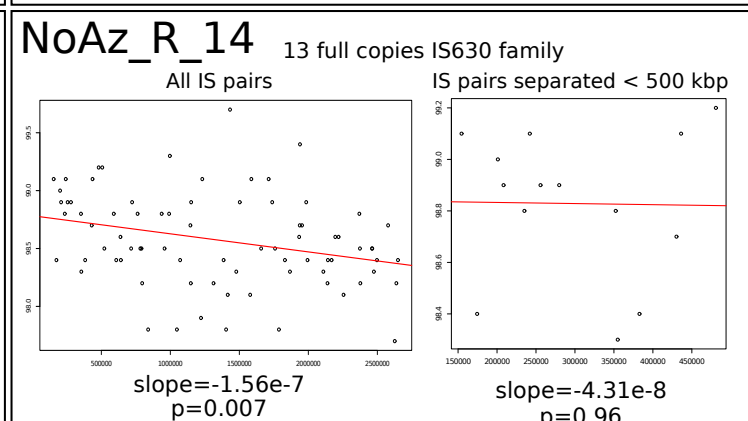
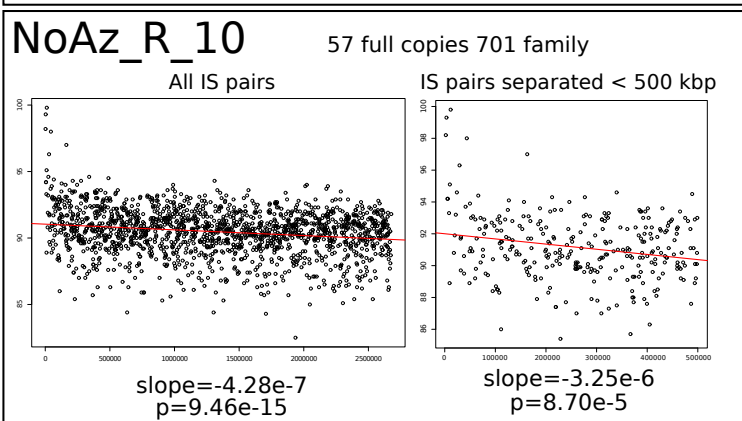
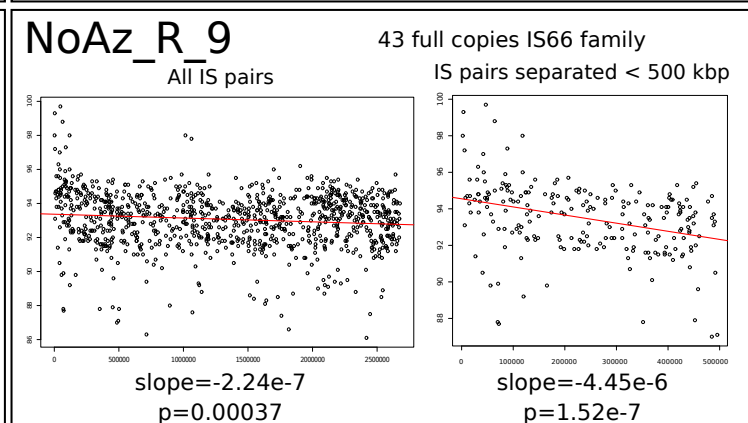
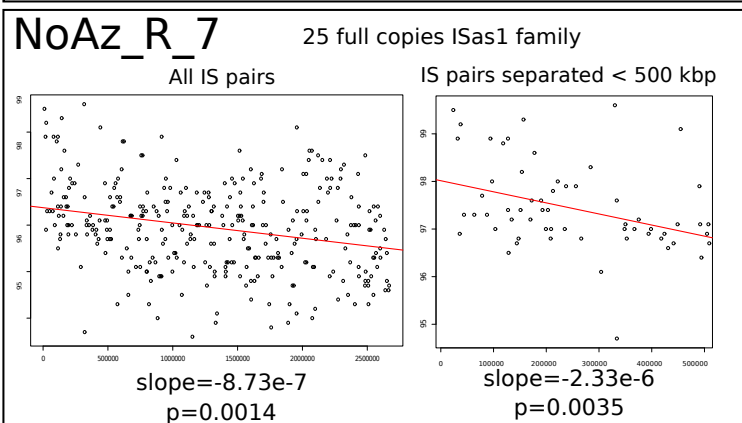
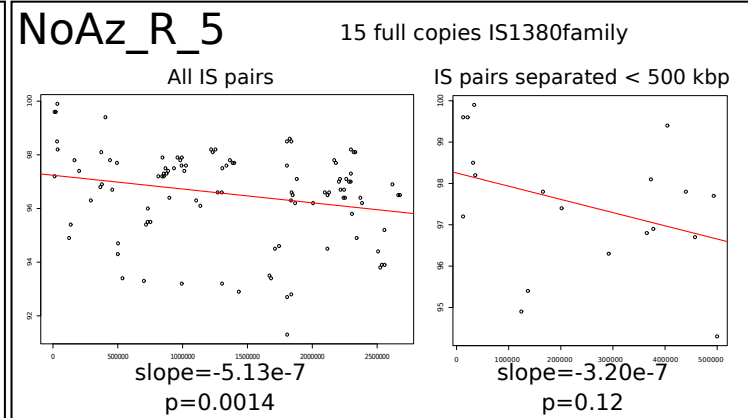
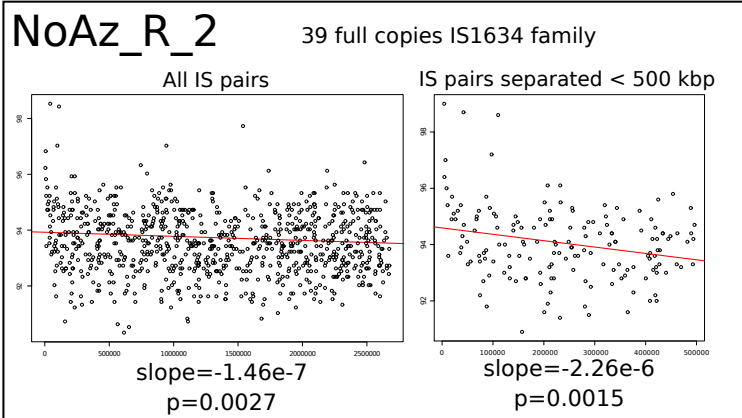


Repeat alignment identities versus distance on chromosome for some repeats in NoAZ



Repeat distance on chromosome (nts)

alignment, % identical positions