HSP vs UP PLODs within SG_ATL

Frequency (y-axis) vs **PLOD** (x-axis)

Expected value given kinship
Unrelated
First cousin
Avuncular
Half−sibling
Parent−offspring
Self

srwqc −> 3879 loci
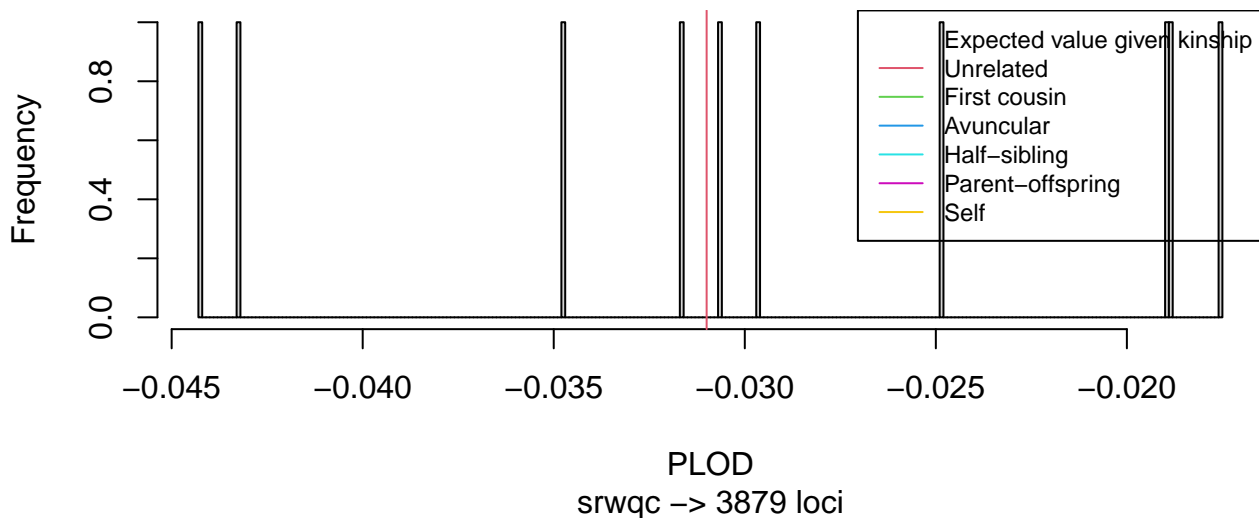
Uncommon values suggest likely close−kin

Frequency (y-axis) vs **PLOD** (x-axis)