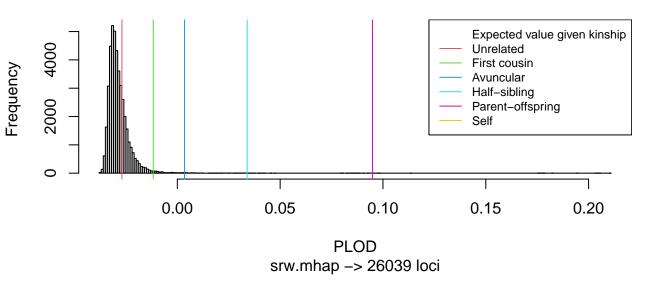
## **HSP vs UP PLODs for all samples**



## Uncommon values suggest likely close-kin

