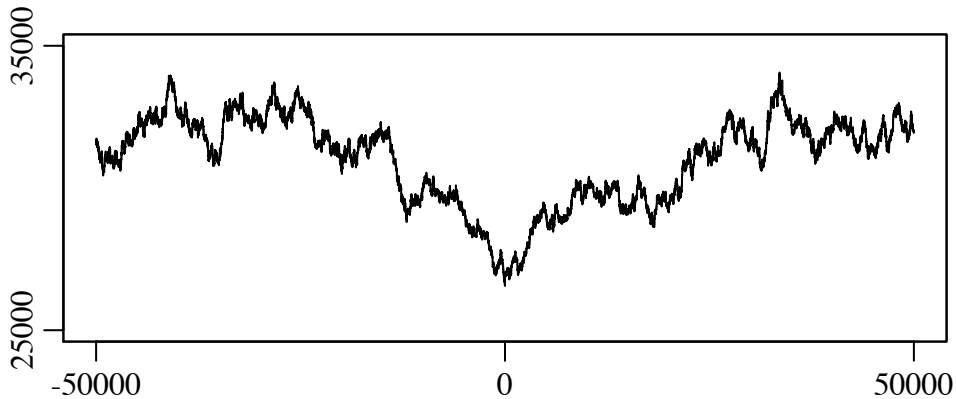


mean tree height



distance from gene