iwidthvaluesBeforeLOwidth.min.bpfiltering
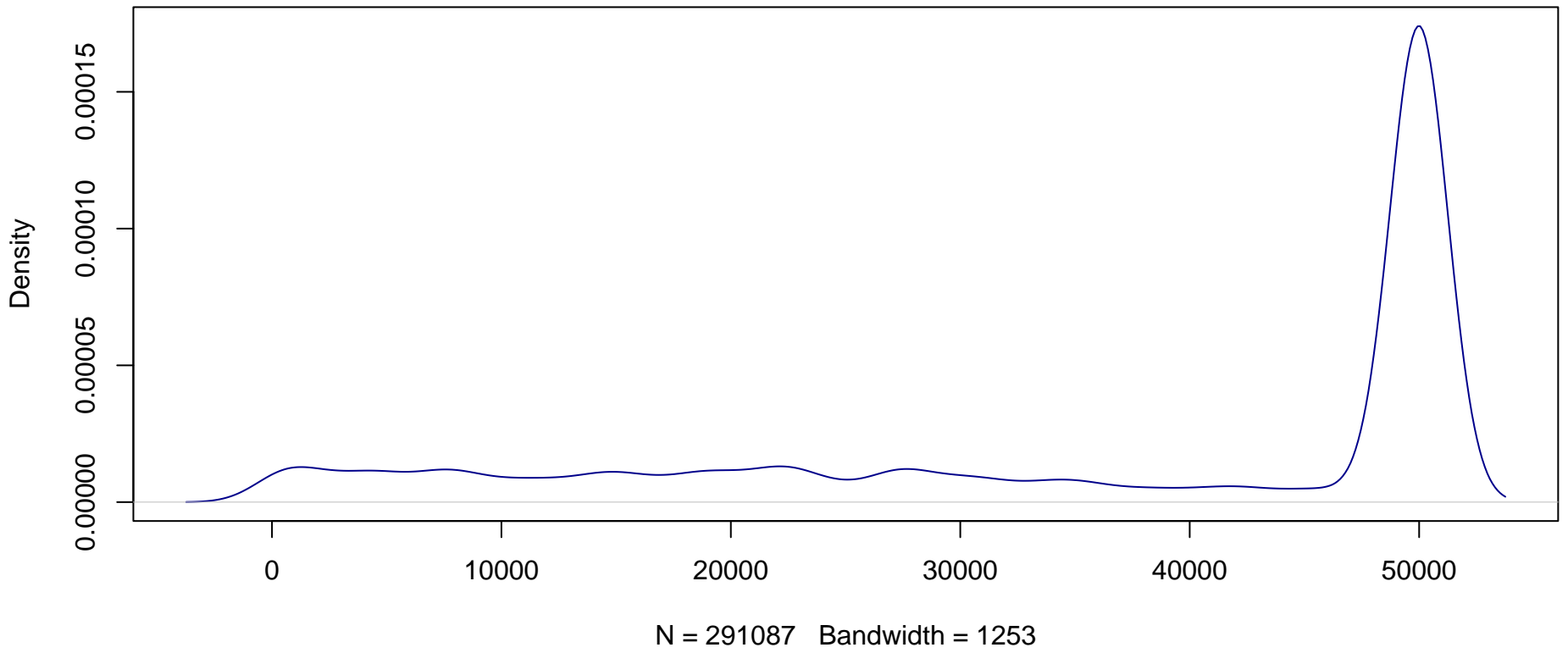**chr21_hg38ToHg19_LOwidth.min.bp30000_genome_state_Phylo−HMRF_mapping_contact50K_norm**

N = 291087   Bandwidth = 1253

jwidthvaluesBeforeLOwidth.min.bpfiltering
chr21_hg38ToHg19_LOwidth.min.bp30000_genome_state_Phylo−HMRF_mapping_contact50K_norm

N = 320774   Bandwidth = 1417