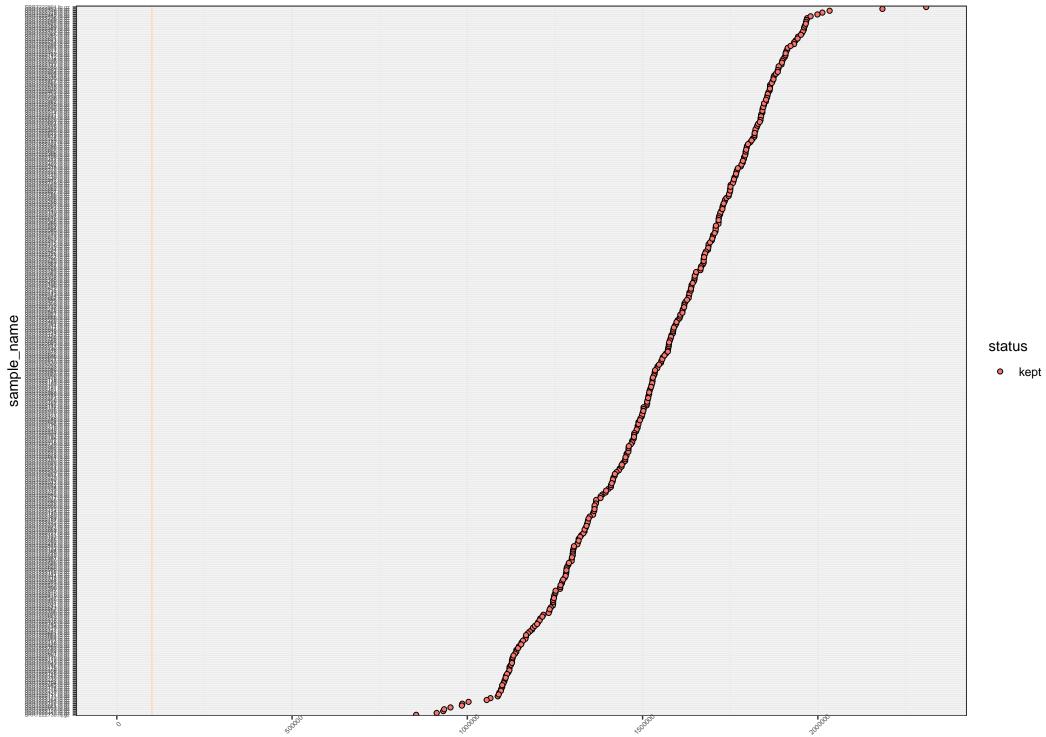
Dot plot of pre-assembly read counts run_name = bioprj_PRJNA473221_Bombus-bifarius



Number of reads per indiv. sample (at end of cleaning/filtering pipeline)