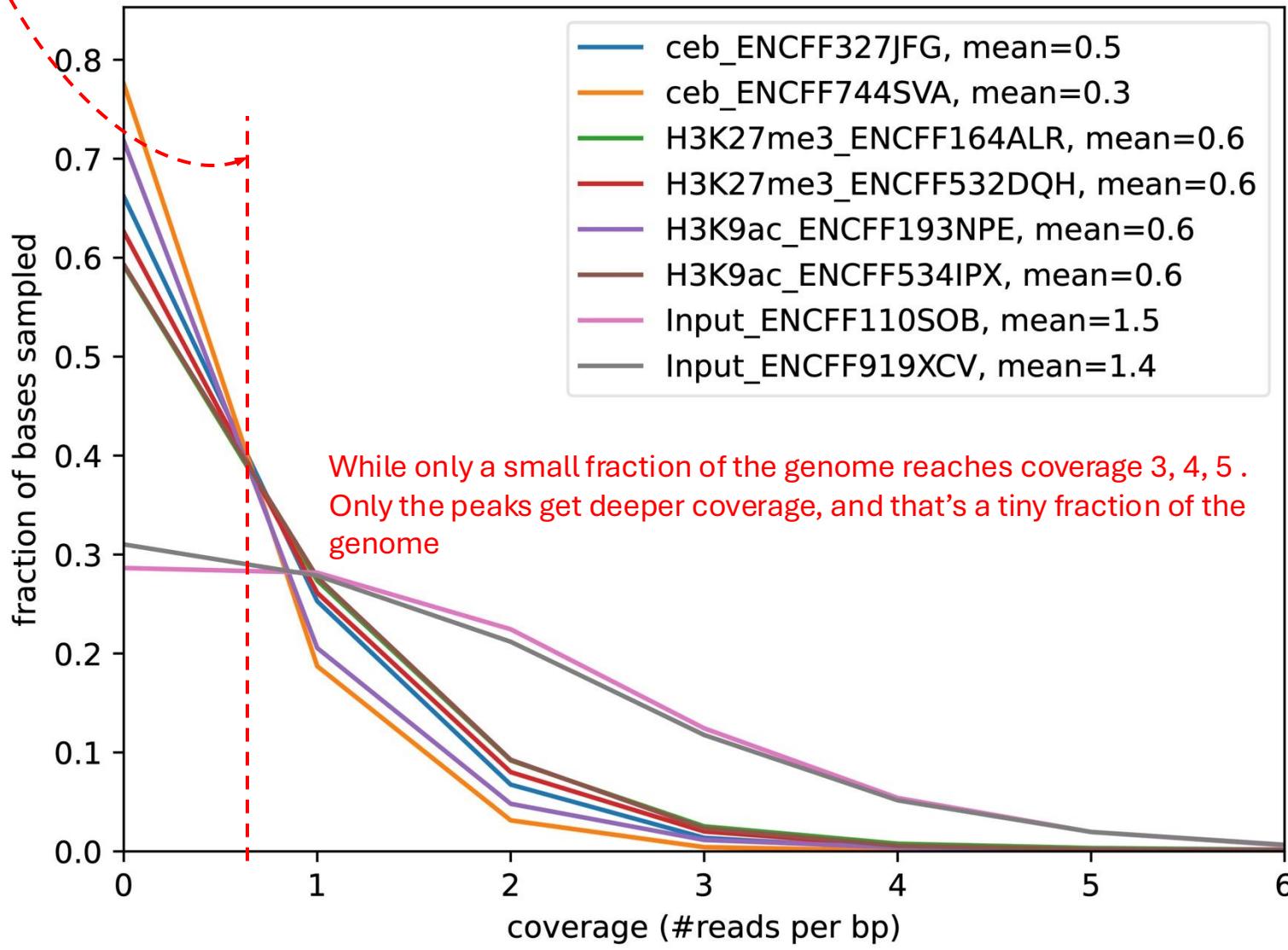


In IP most bases are only lightly covered between 0 and 1



Higher coverage means more reads piling up at that genomic position.