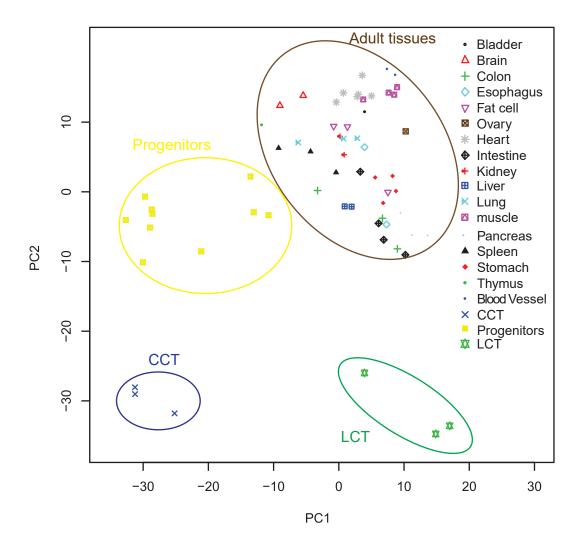
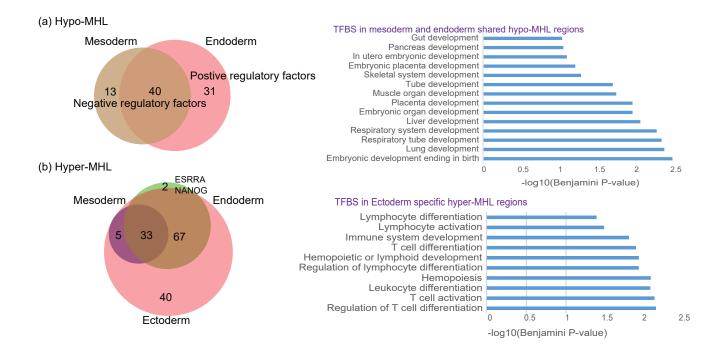
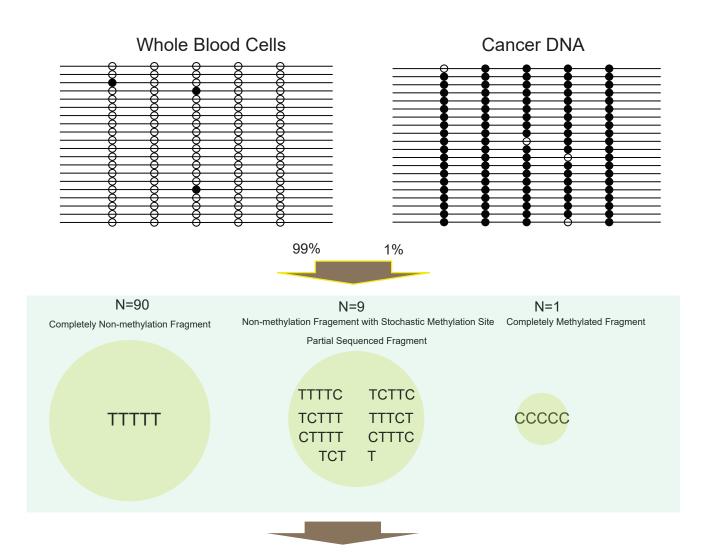


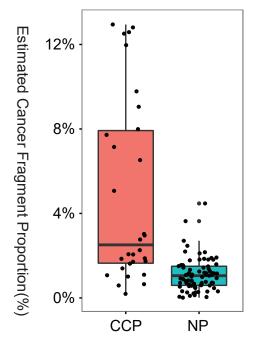
Distance from center of methylation haplotype blocks in base pairs

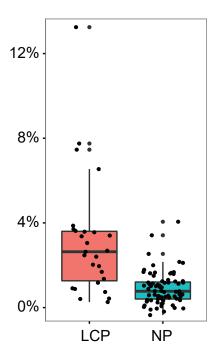


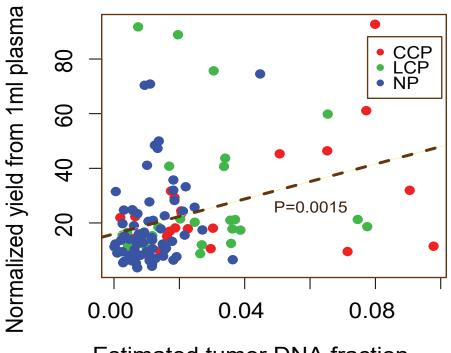




Average 5mC level = 0.03 Methylation haplotype Load(MHL) = 0.13







Estimated tumor DNA fraction

