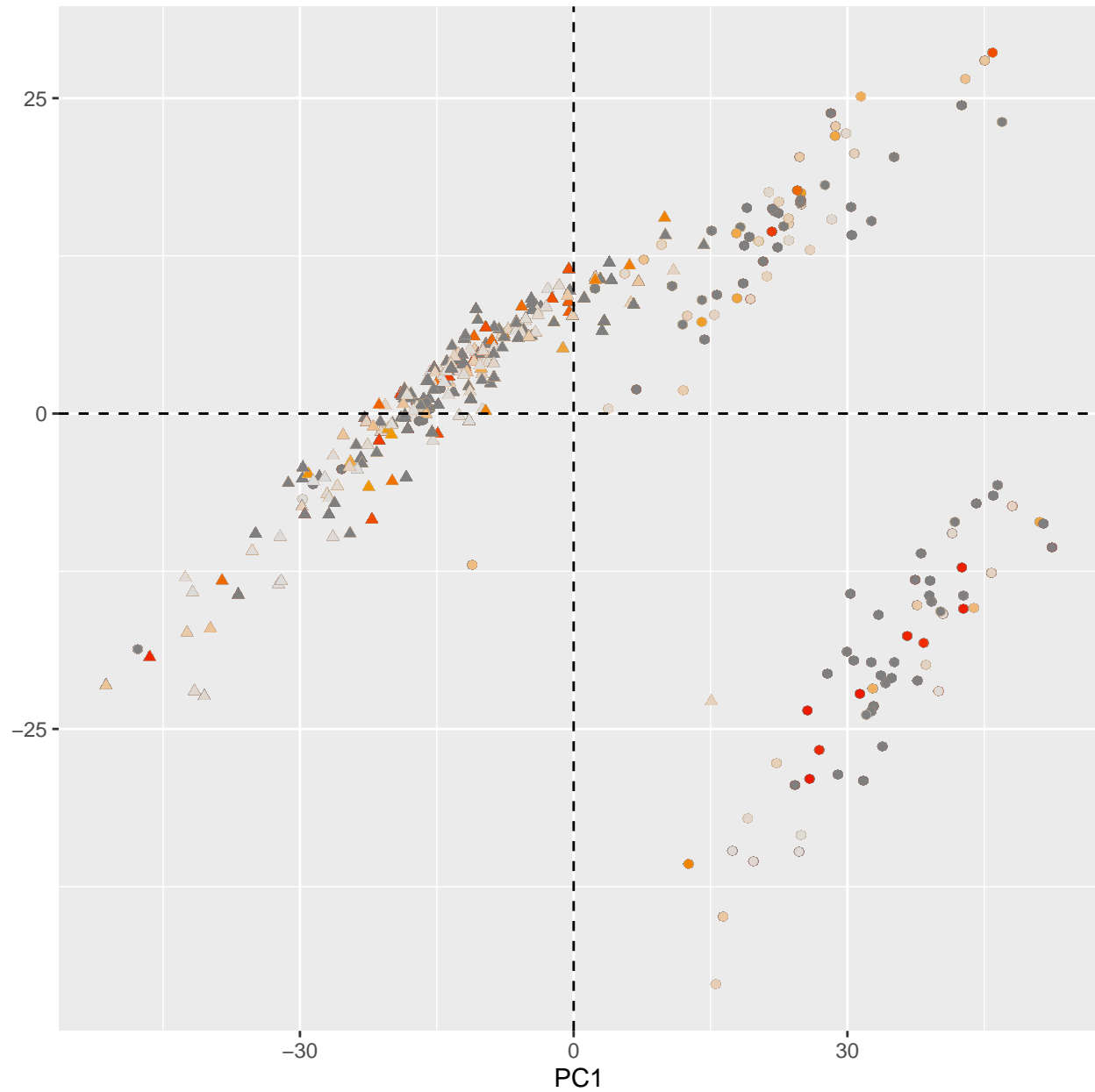


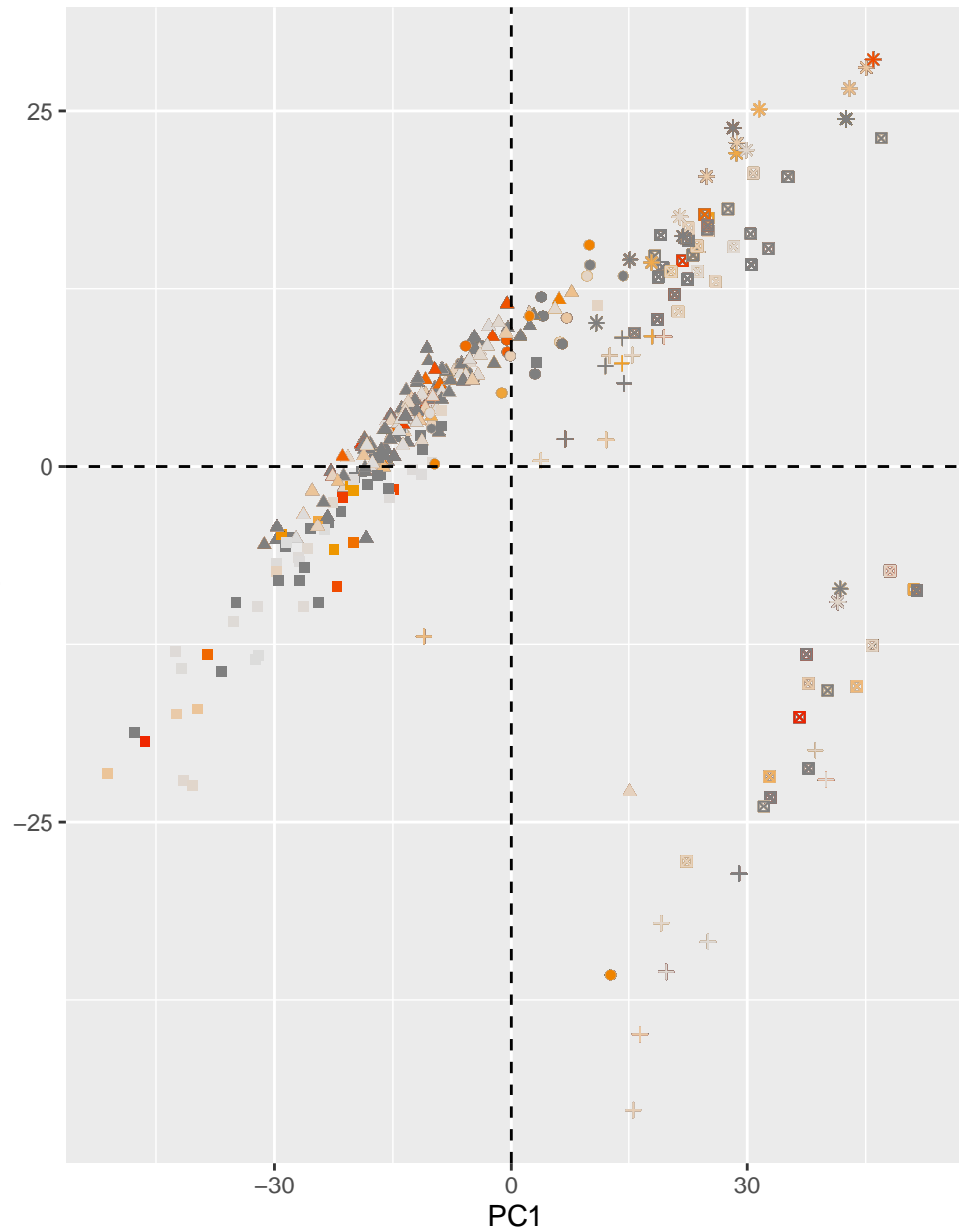
n.samples: 345



n.samples: 345



n.samples: 345



prob

0.75

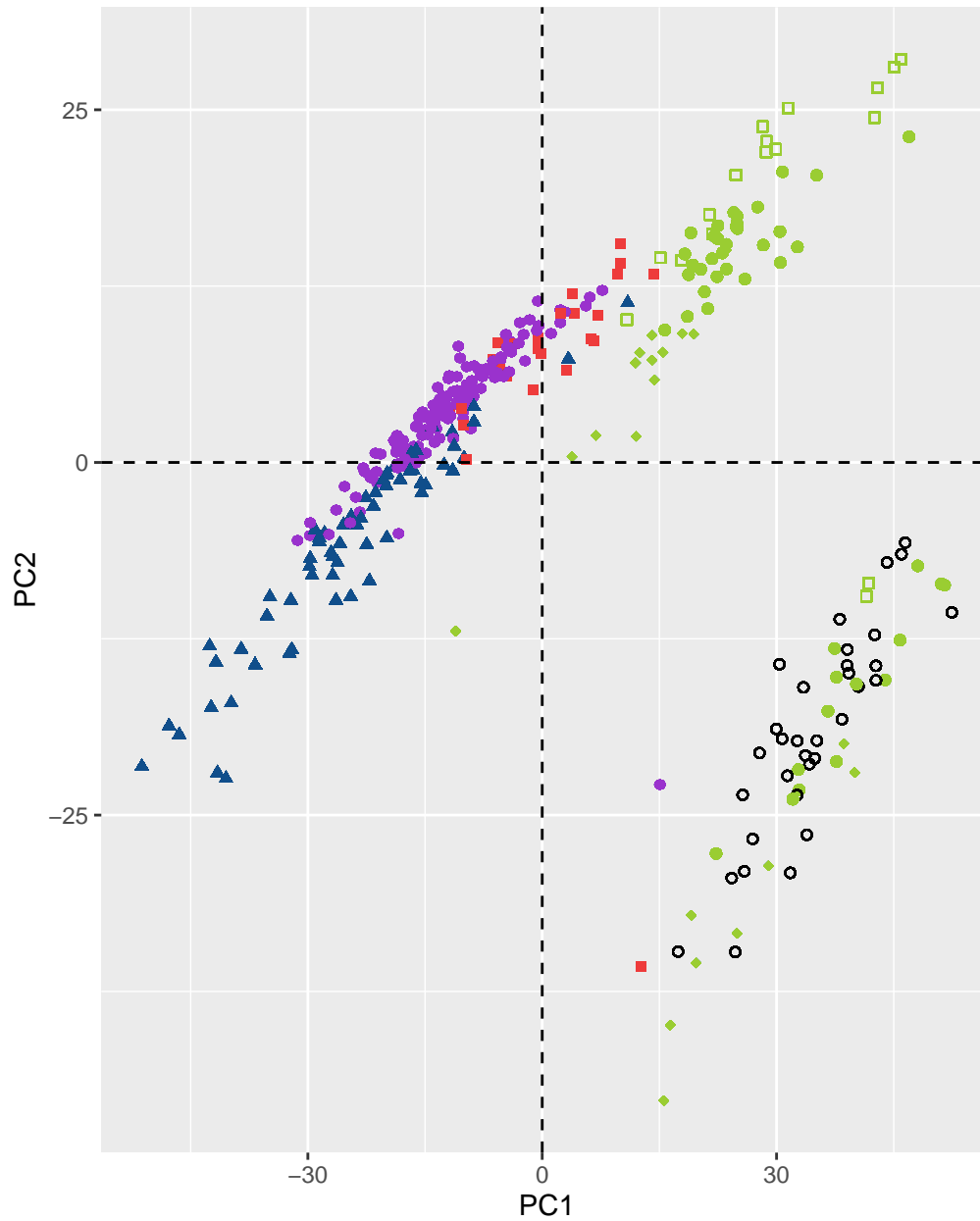
0.50

0.25

factor(DNA_methylation_clusters)

- LGM1
- ▲ LGM2
- LGM3
- + LGM4
- ⊠ LGM5
- * LGM6
- LGr_Unkown

n.samples: 345



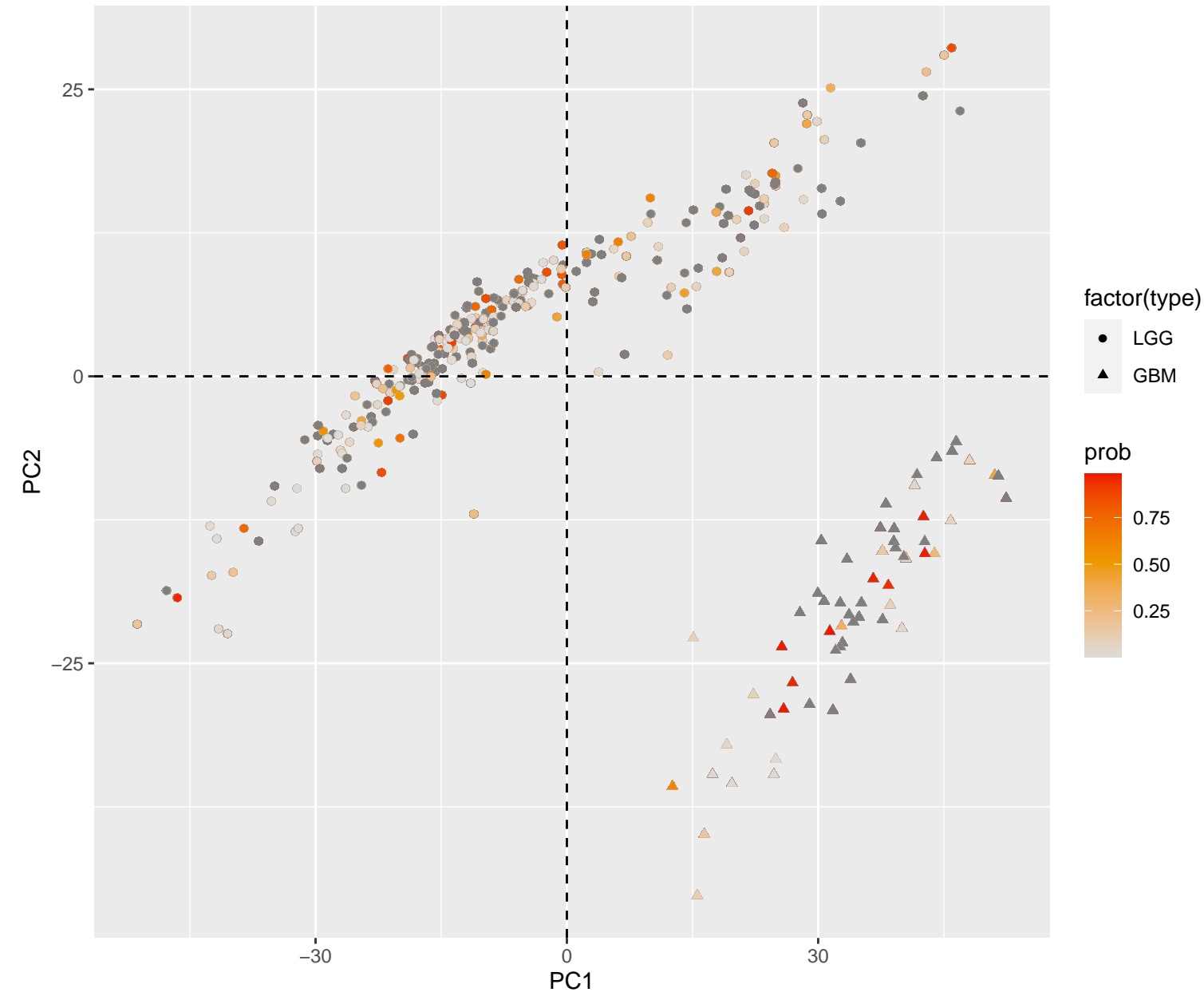
factor(DNA_methylation_clusters)

- LGM1
- LGM2
- ▲ LGM3
- ◆ LGM4
- LGM5
- LGM6
- LGr_Unkown

DNA_methylation_clusters

- LGM1
- LGM2
- LGM3
- LGM4
- LGM5
- LGM6
- LGr_Unkown

n.samples: 345



n.samples: 345

