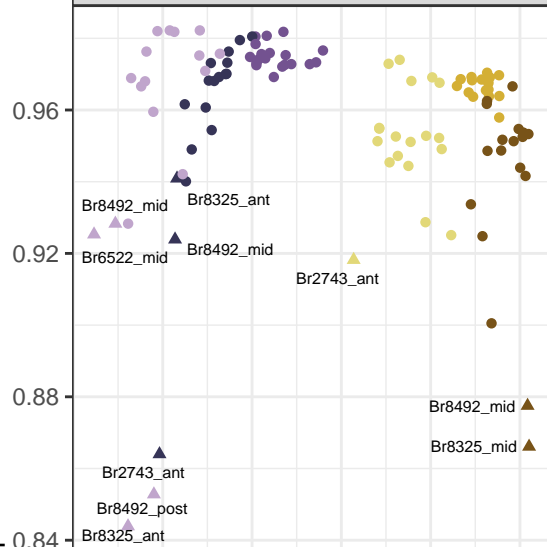


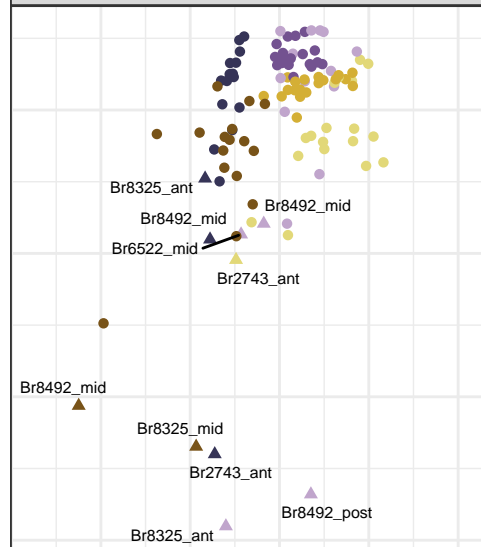
concordMapRate

concordMapRate

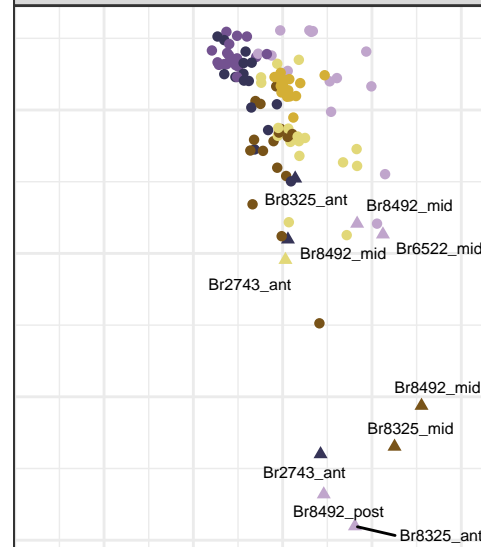
PC1: 59.1% Var Expl



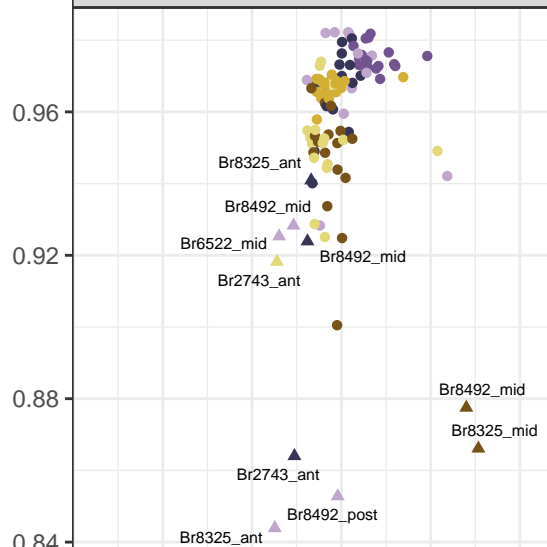
PC2: 9.17% Var Expl



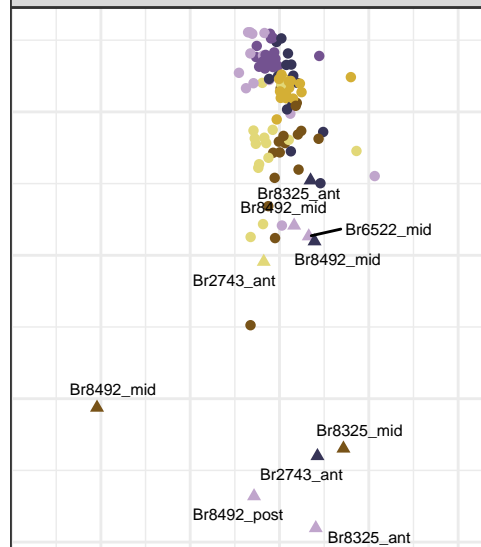
PC3: 5.94% Var Expl



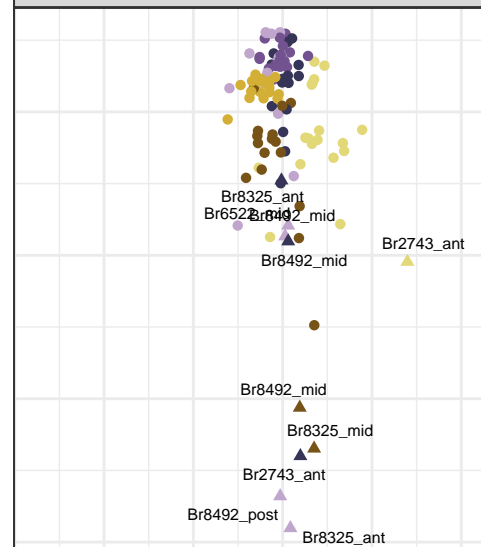
PC4: 3.75% Var Expl



PC5: 2.72% Var Expl



PC6: 2.27% Var Expl



library_combo

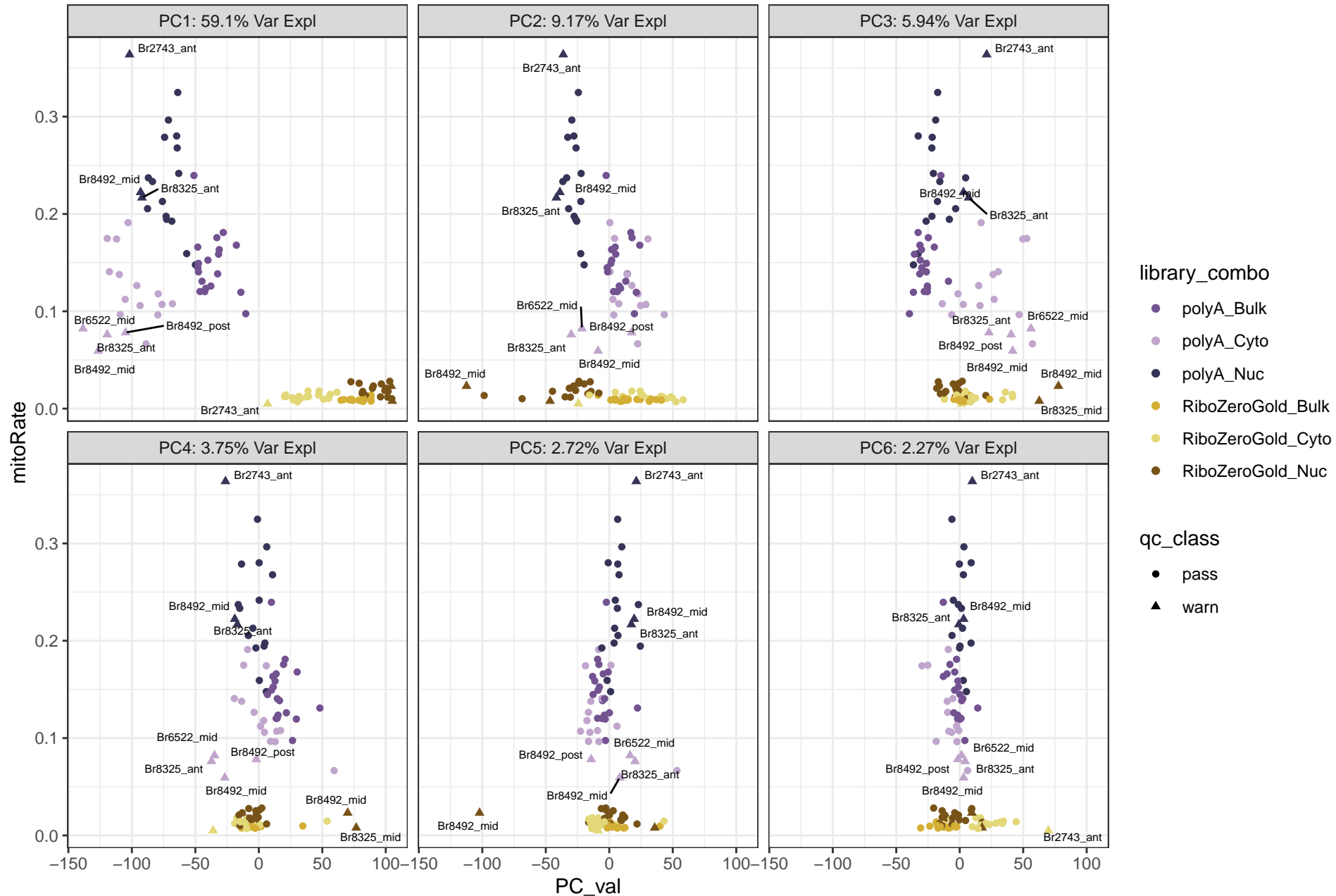
- polyA_Bulk
- polyA_Cyto
- polyA_Nuc
- RiboZeroGold_Bulk
- RiboZeroGold_Cyto
- RiboZeroGold_Nuc

qc_class

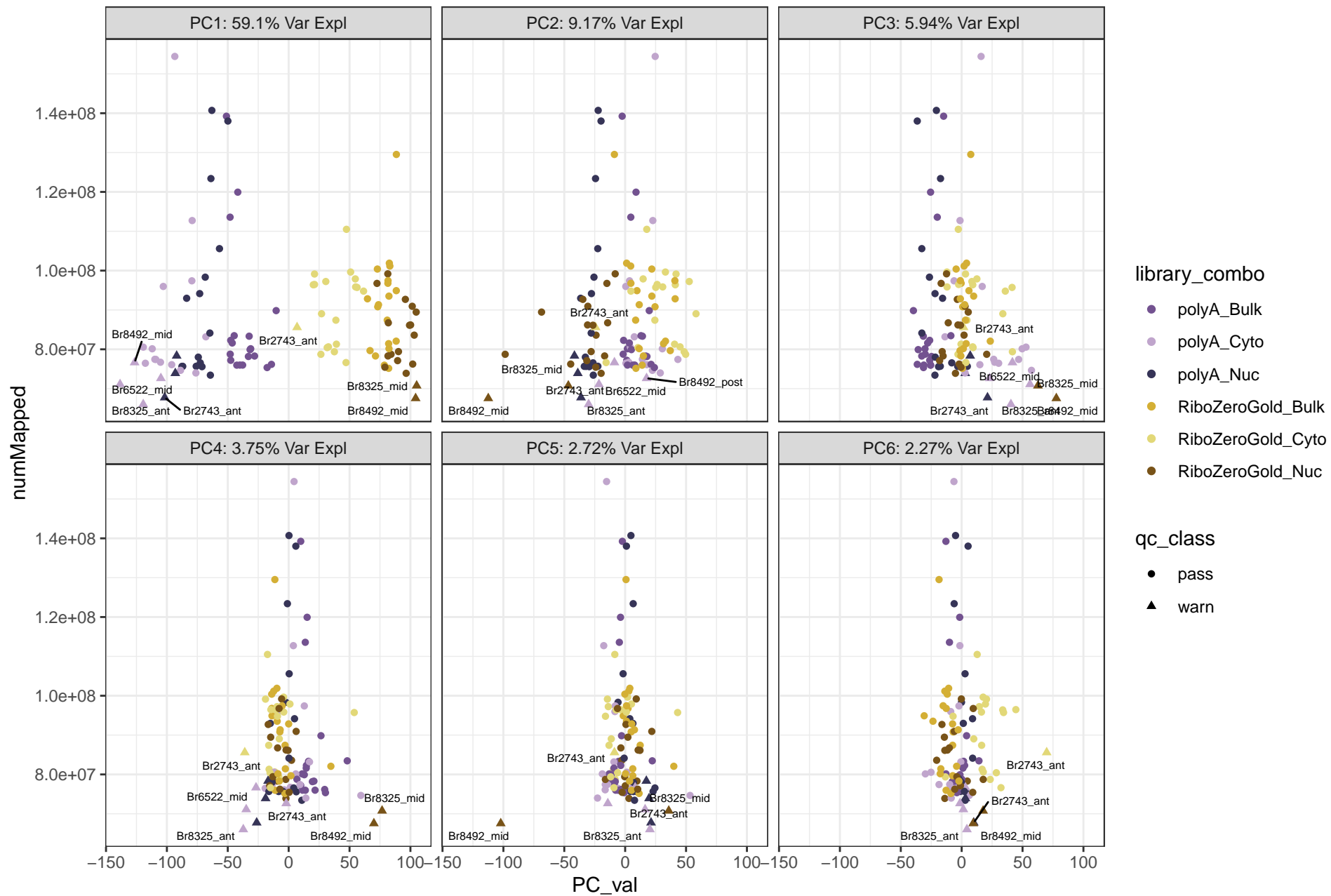
- pass
- warn

PC_val

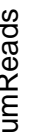
mitoRate



numMapped



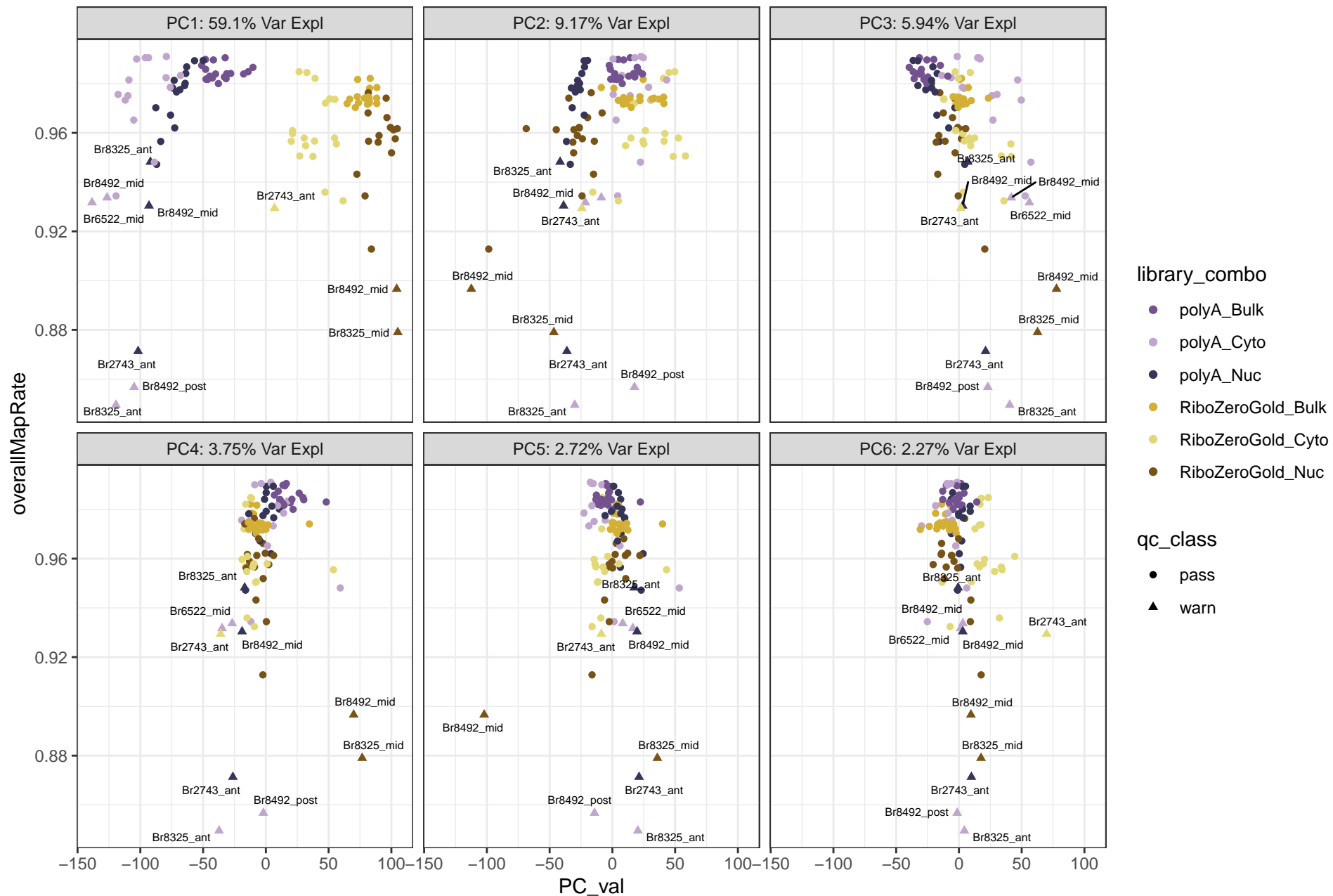
numReads



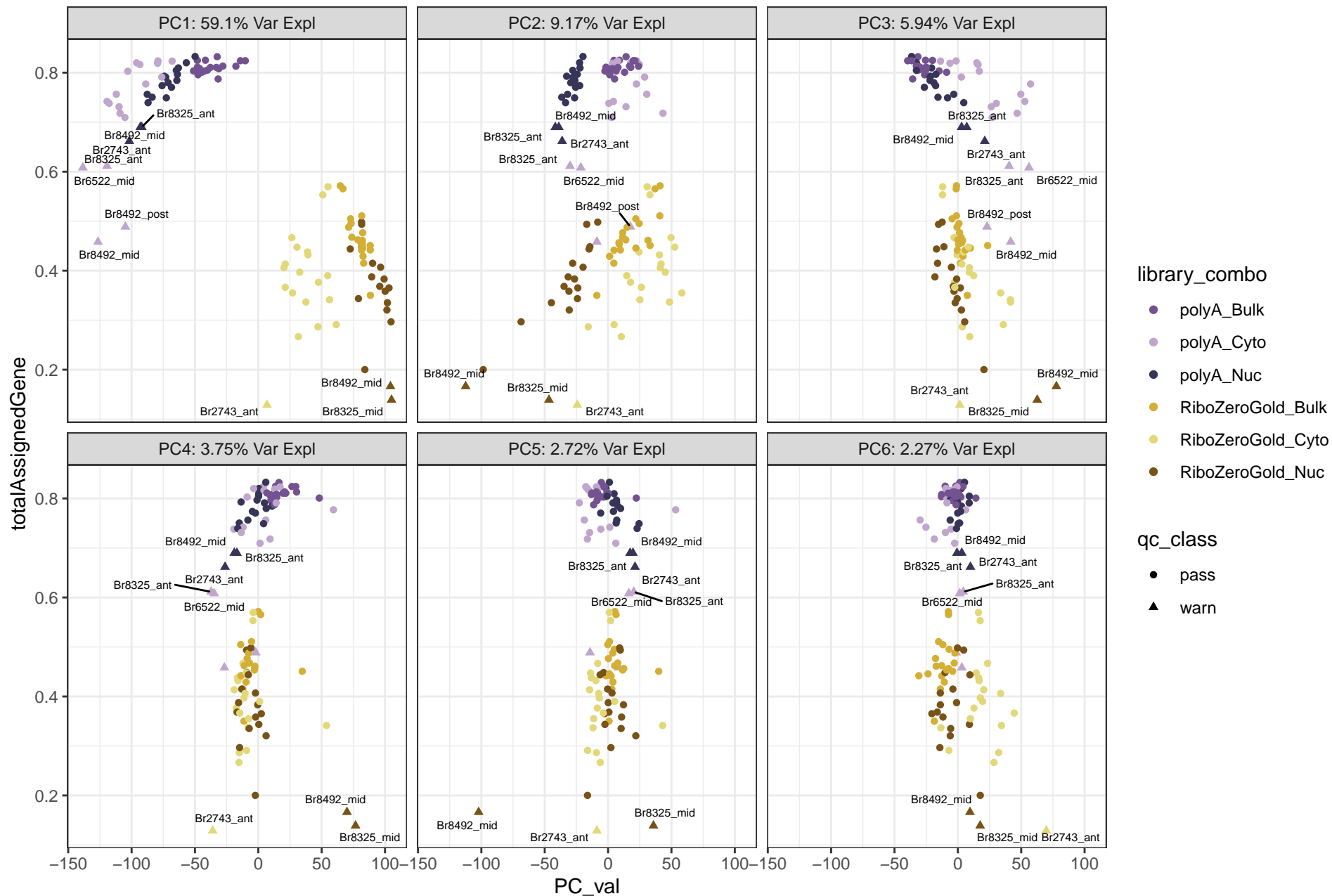
- polyA_Bulk
- polyA_Cyto
- polyA_Nuc
- RiboZeroGold_Bulk
- RiboZeroGold_Cyto
- RiboZeroGold_Nuc

- pass
- ▲ warn

overallMapRate



totalAssignedGene



totalMapped

