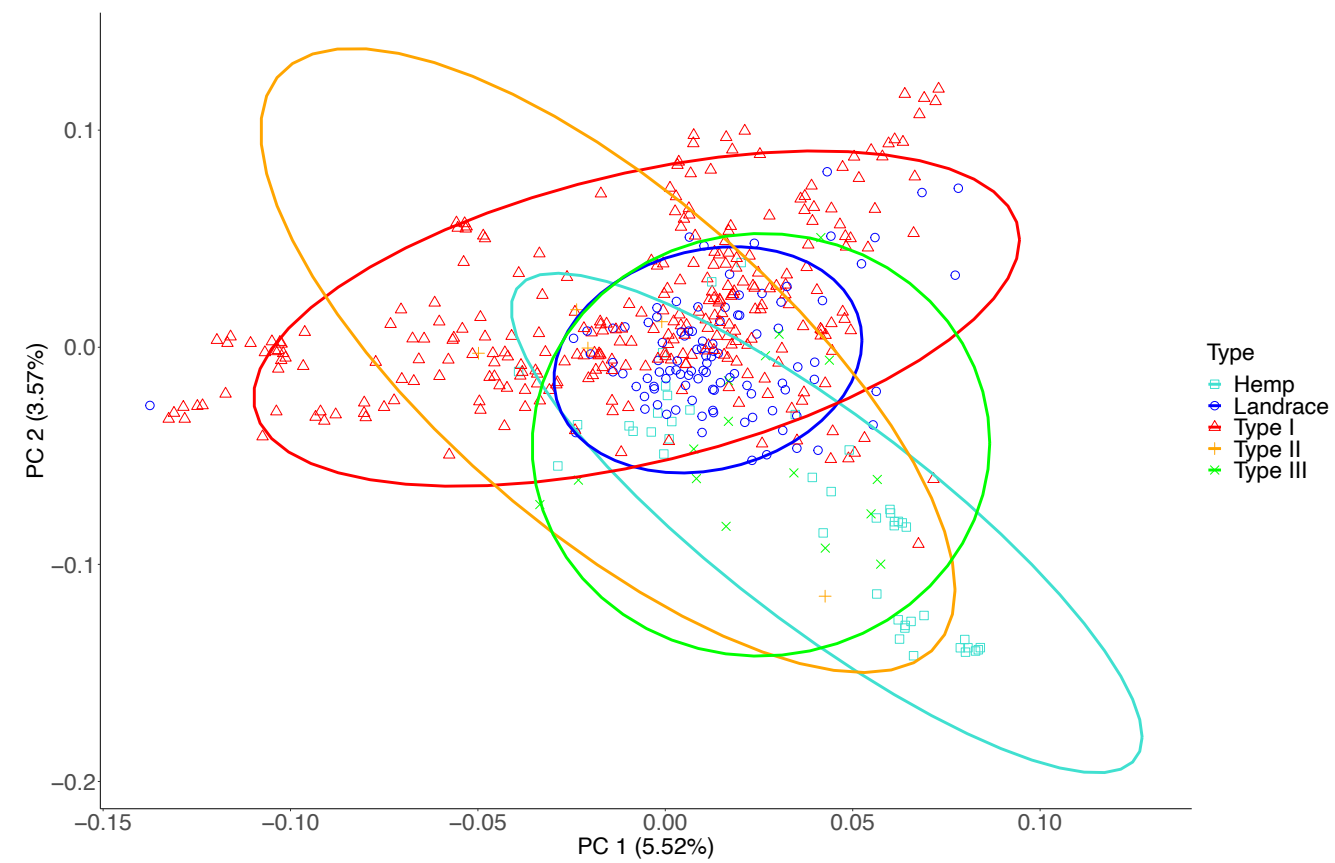
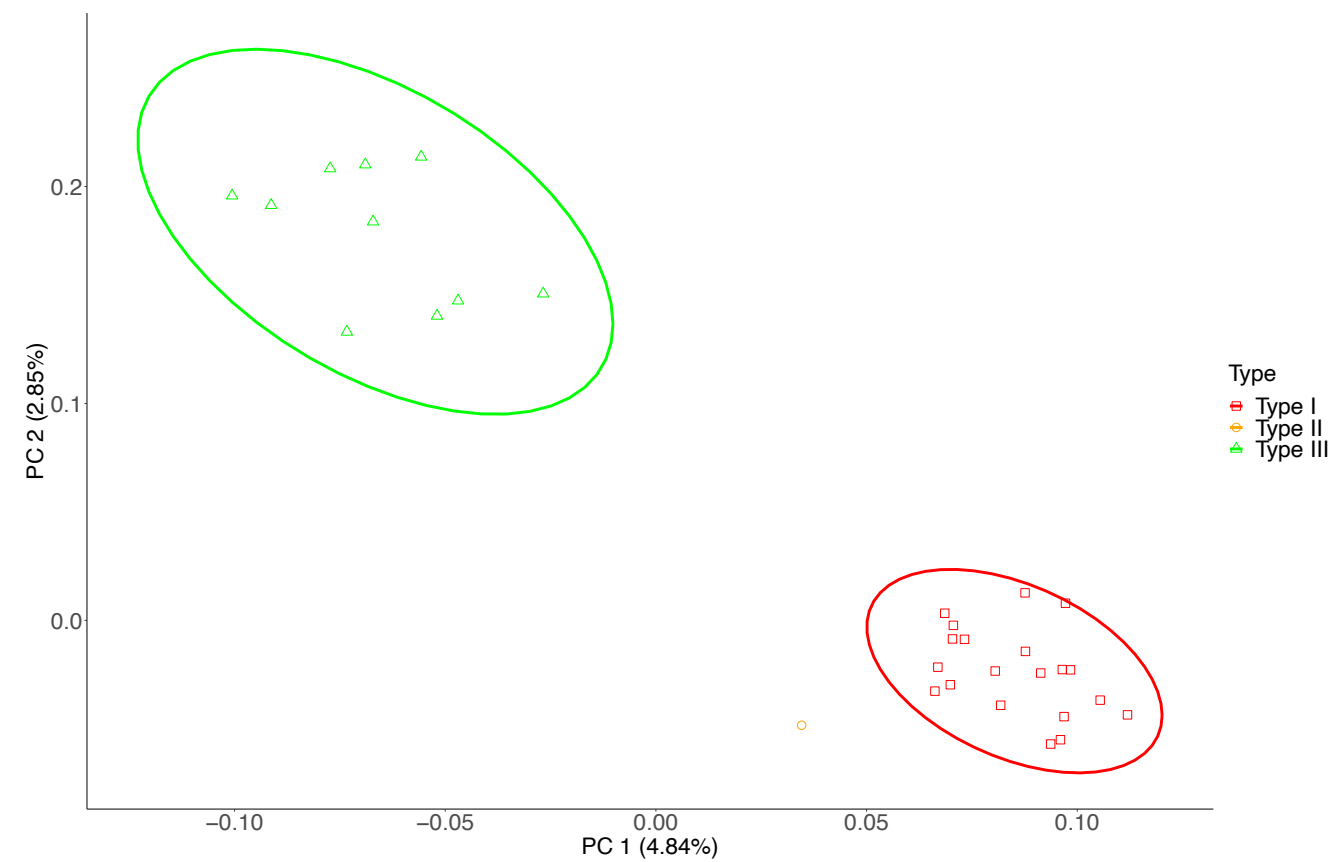


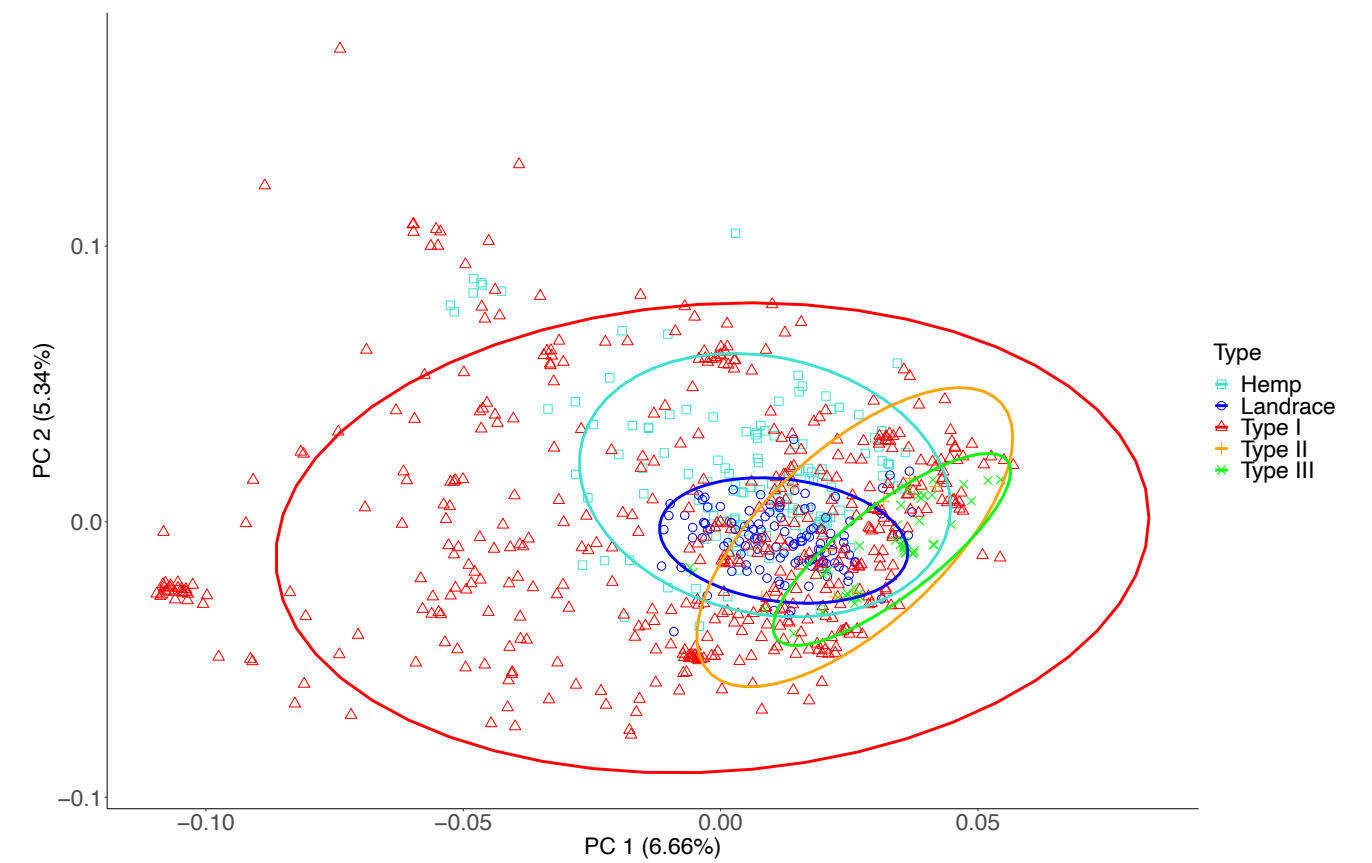
A



B



C



D

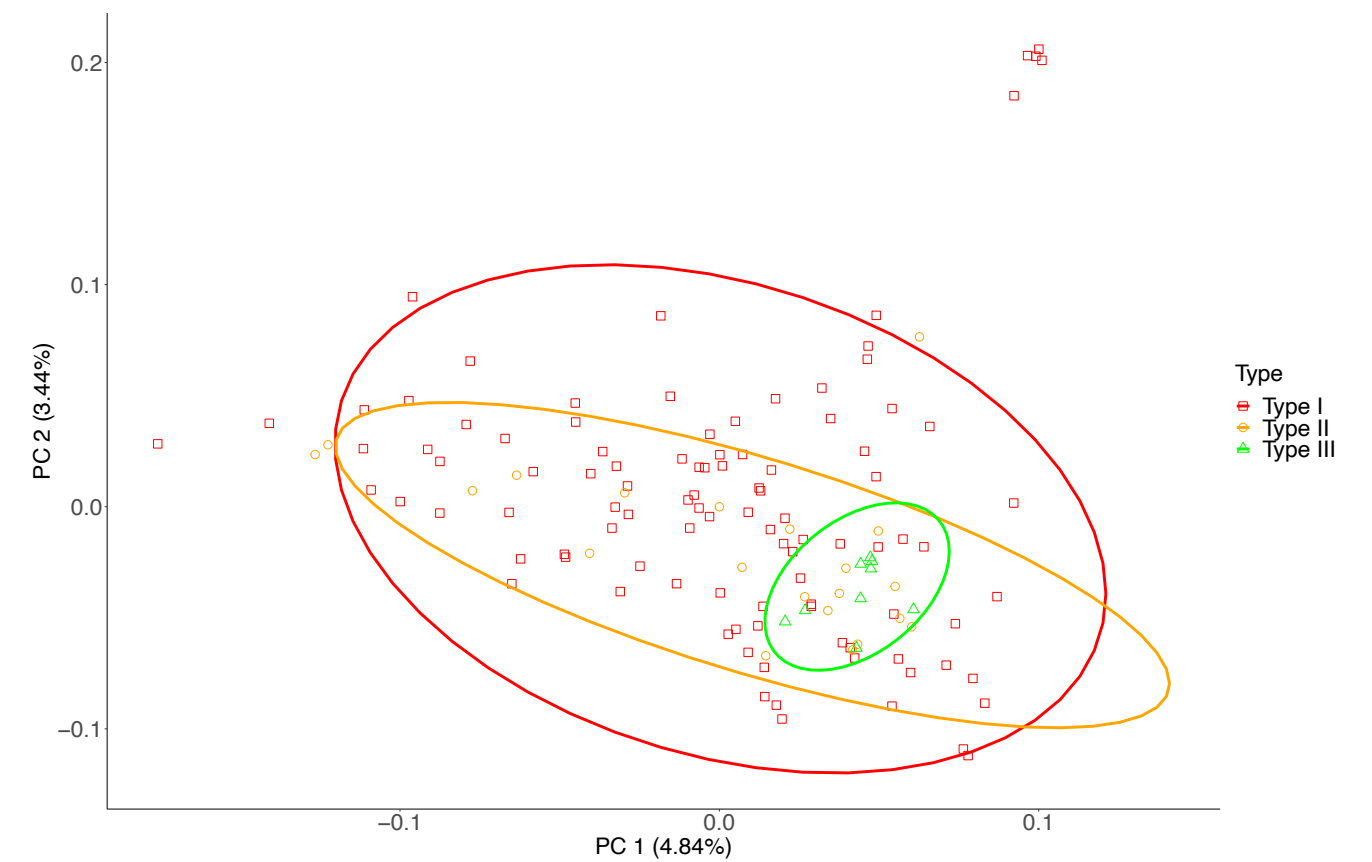


Figure 3 Examination of use-type association across datasets **(A)** Principal component analysis (PCA) from 520 nuclear SNPs for the LeafWorks Inc. dataset **(B)** PCA from 213 SNPs Phylos Biosciences(n=845) dataset **(C)** PCA from 6,865 nuclear SNPs for the Soorni et al., 2017 dataset where cannabinoid content could be determined due to recent publication for 31/94 samples. **(D)** PCA from 5,045 nuclear SNPs for the Medicinal Genomics StrainSEEK V1 dataset.