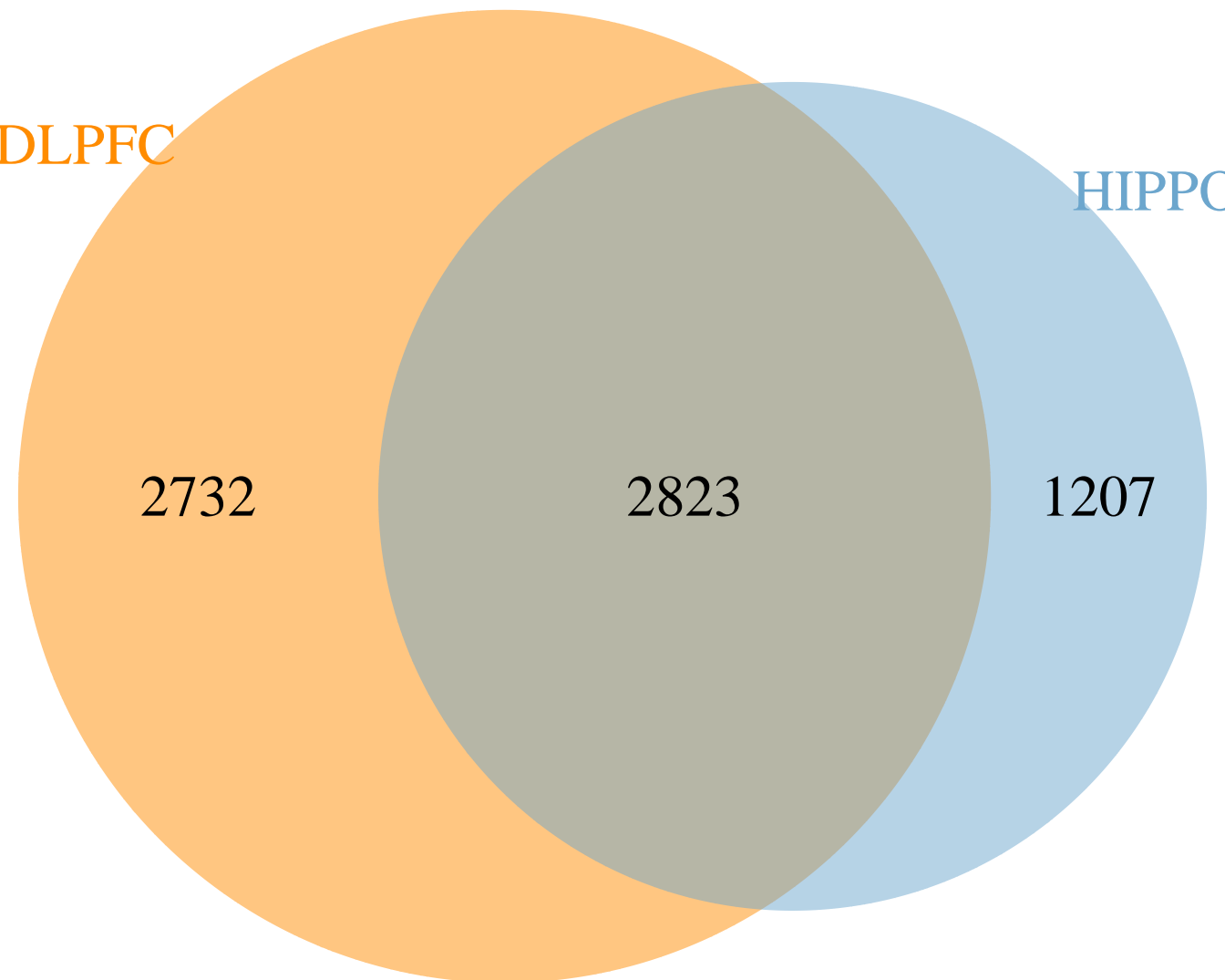
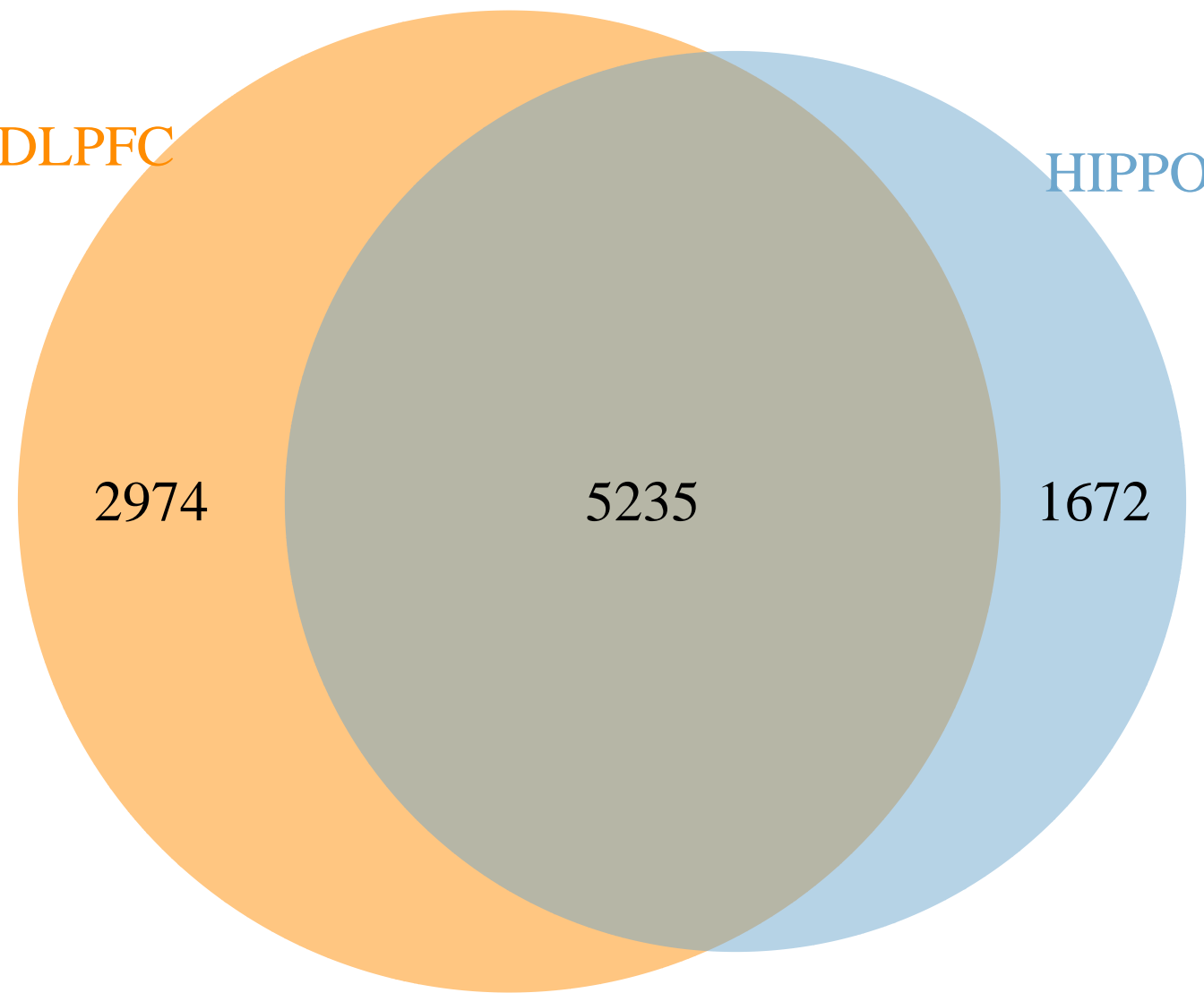


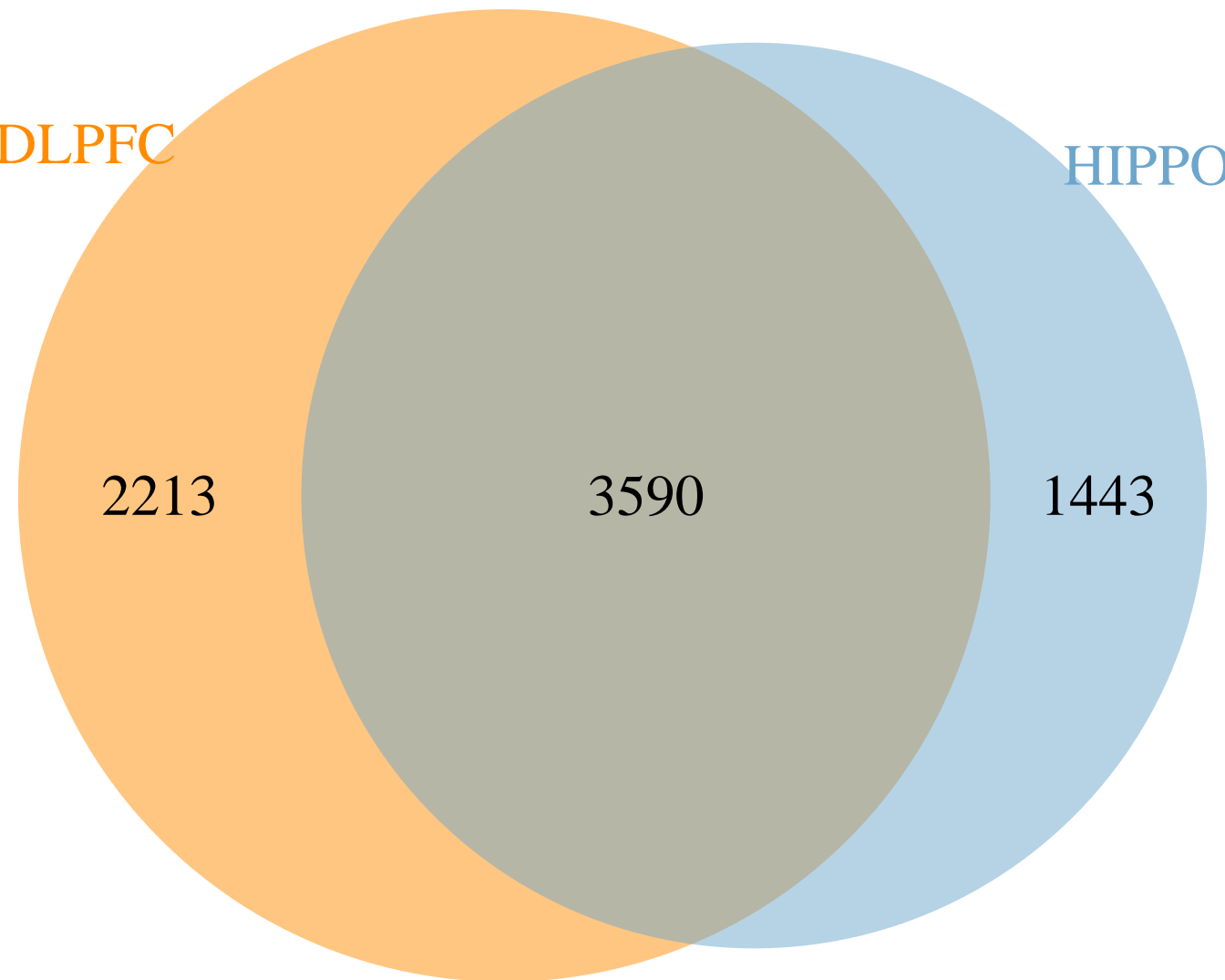
# Features with TWAS weights (by gene ID) – gene



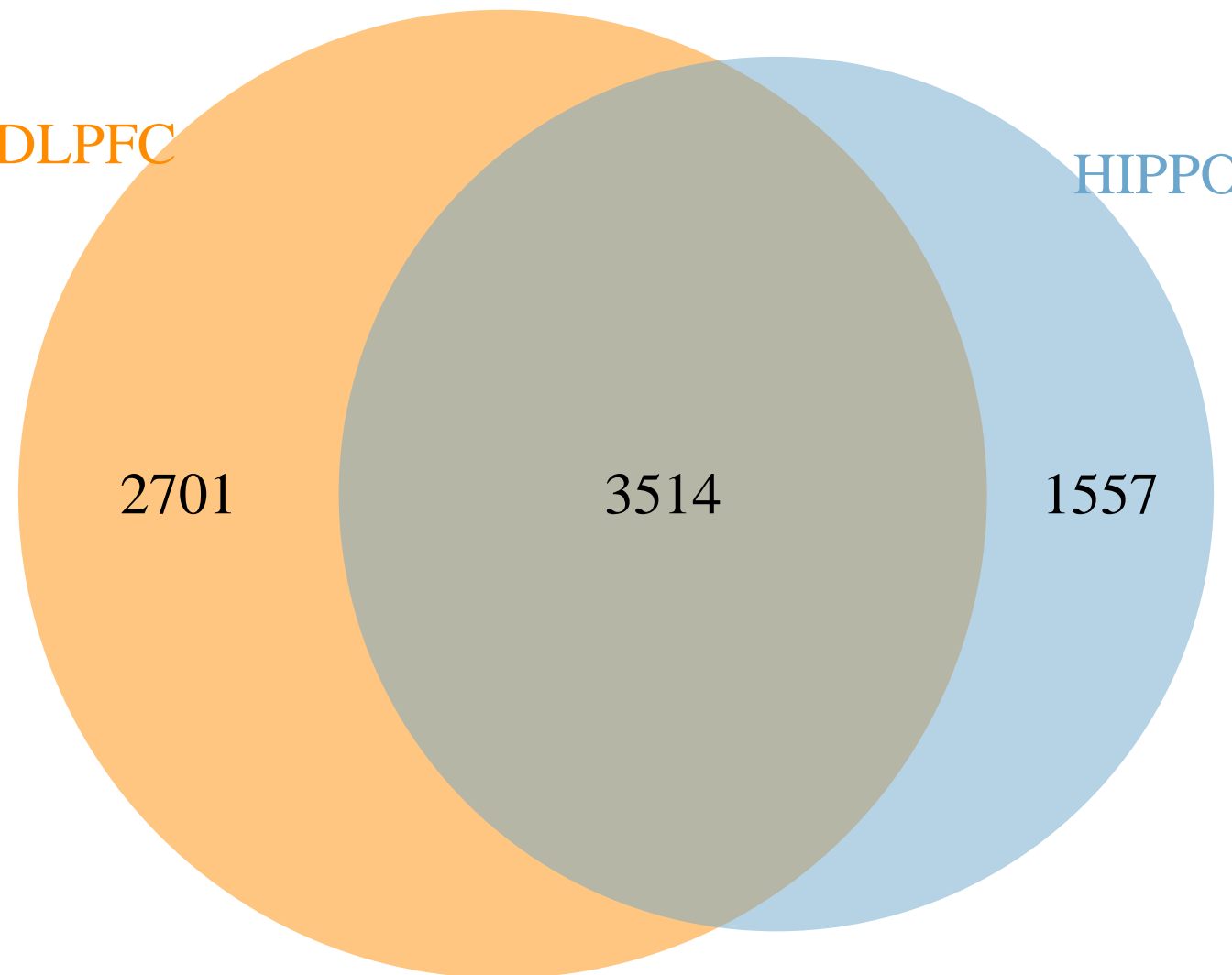
# Features with TWAS weights (by gene ID) – exon



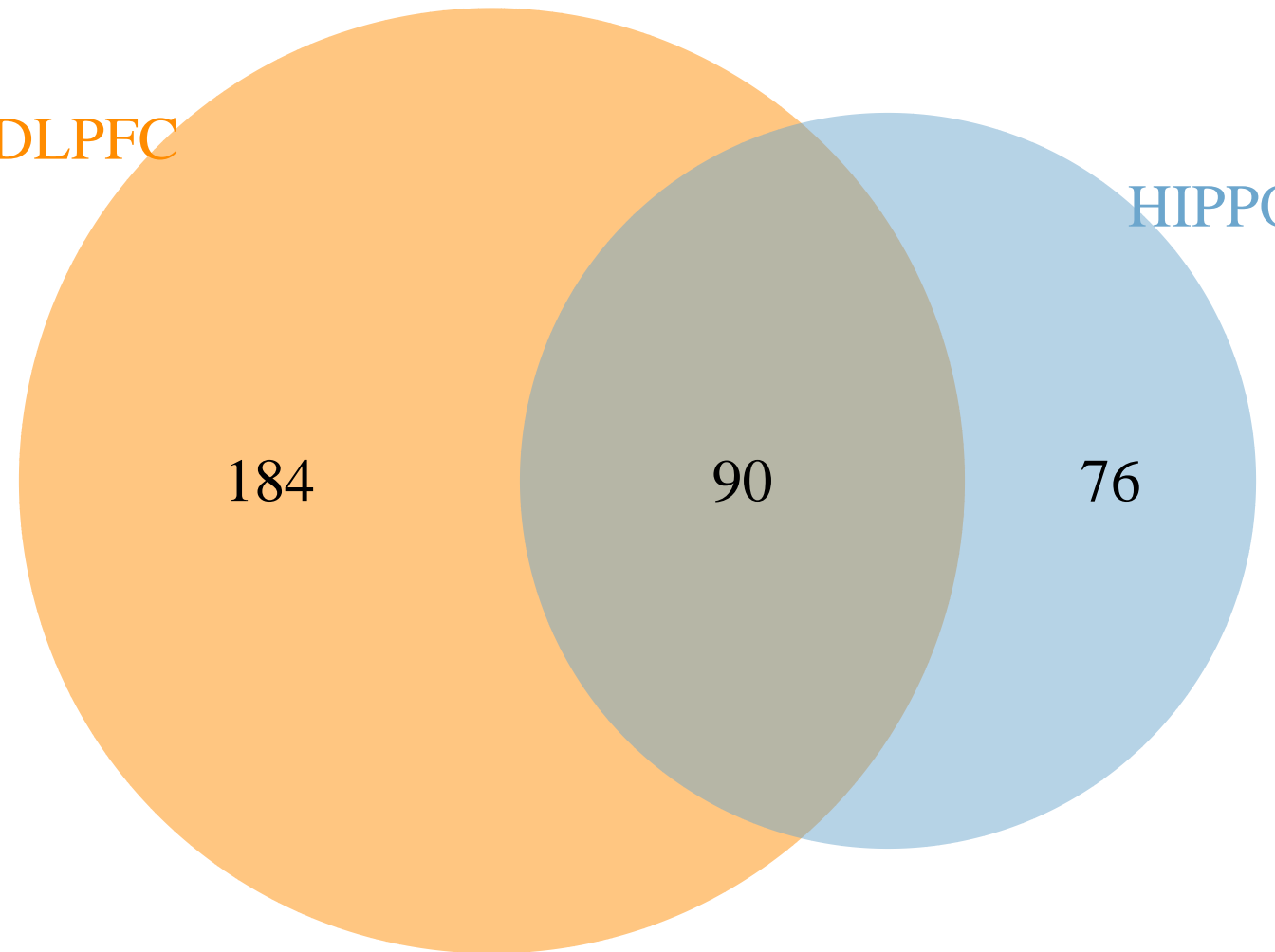
# Features with TWAS weights (by gene ID) – jxn



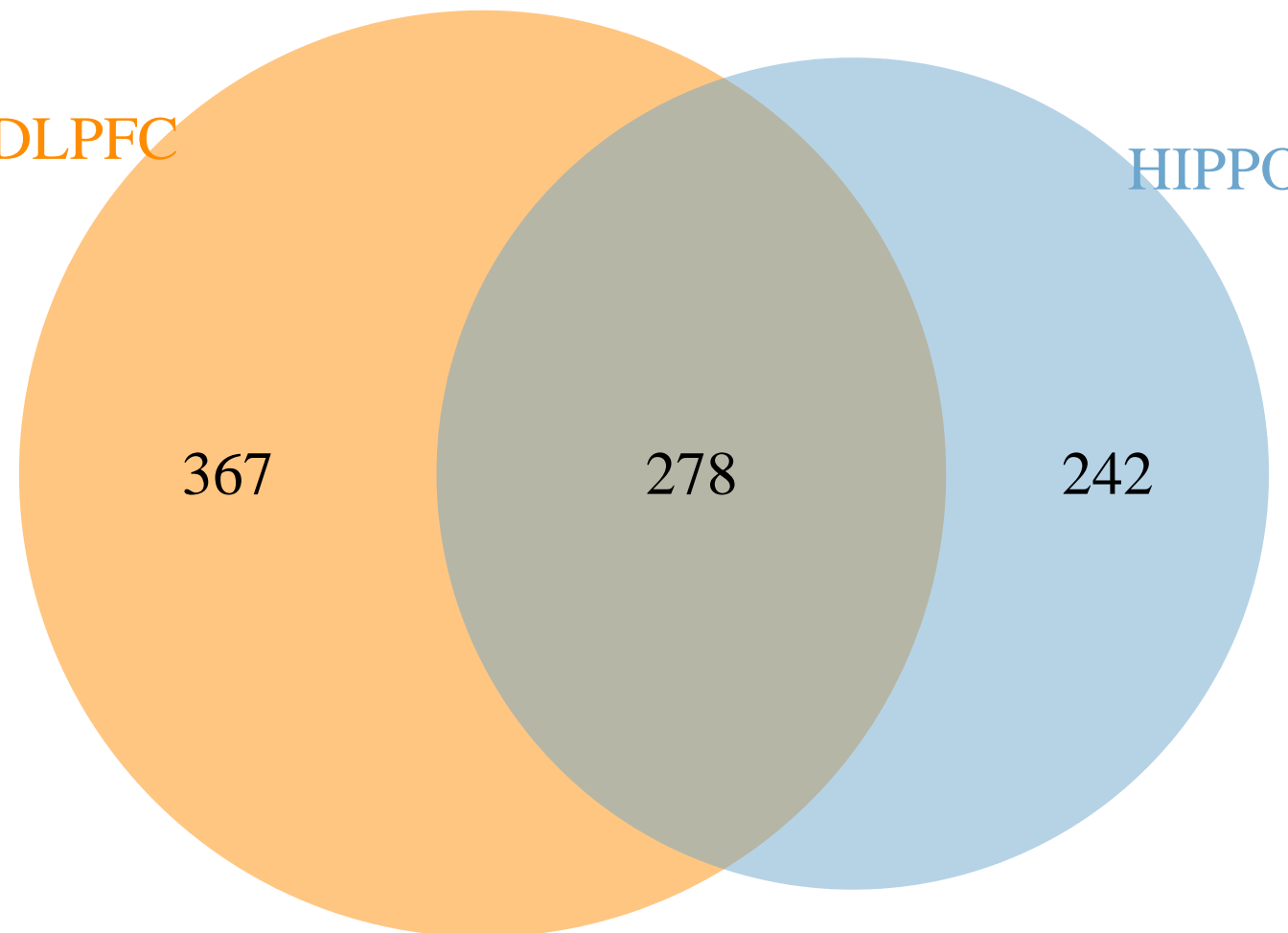
# Features with TWAS weights (by gene ID) – tx



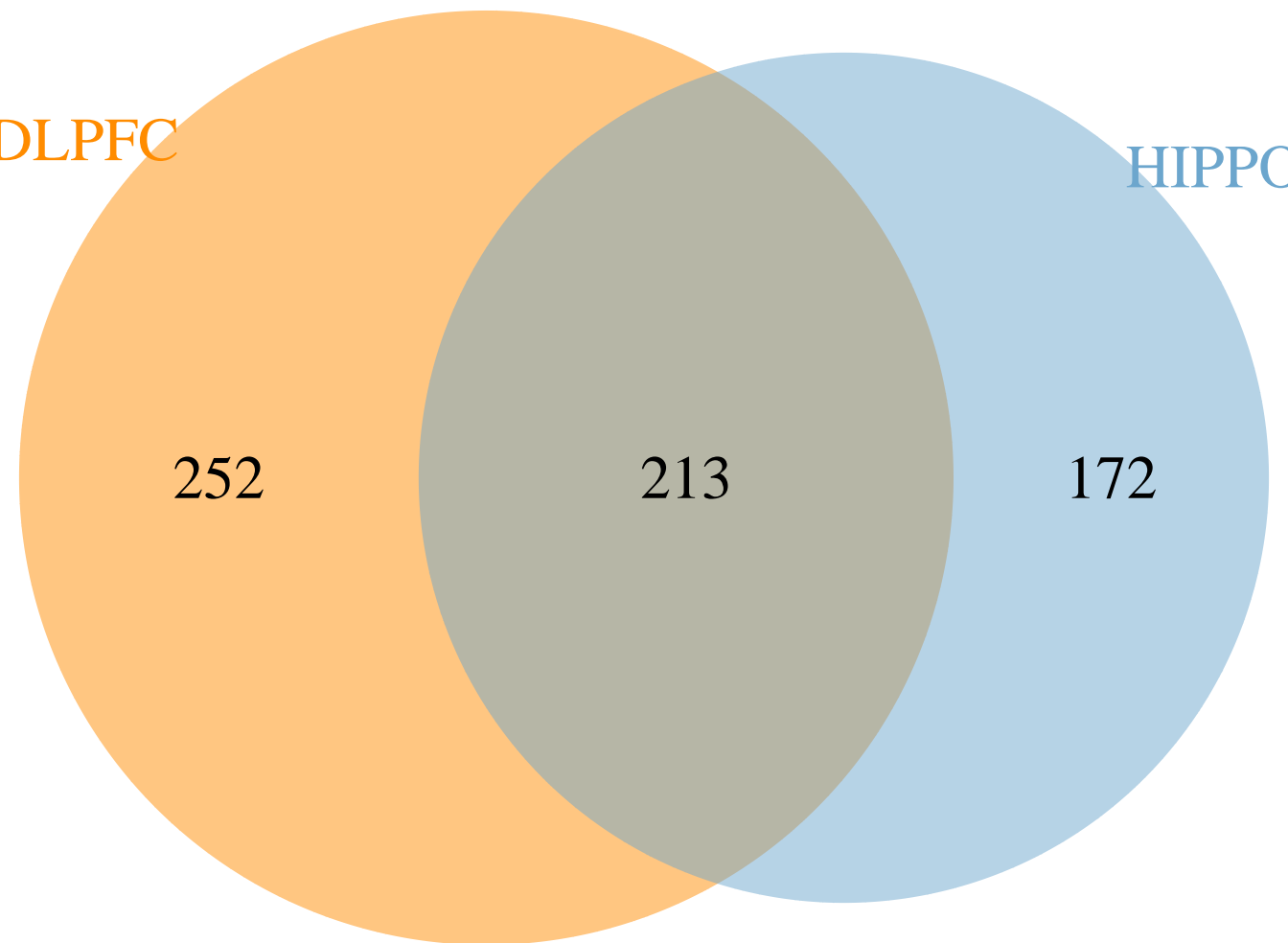
Features with TWAS FDR<5% (by gene ID) – gene



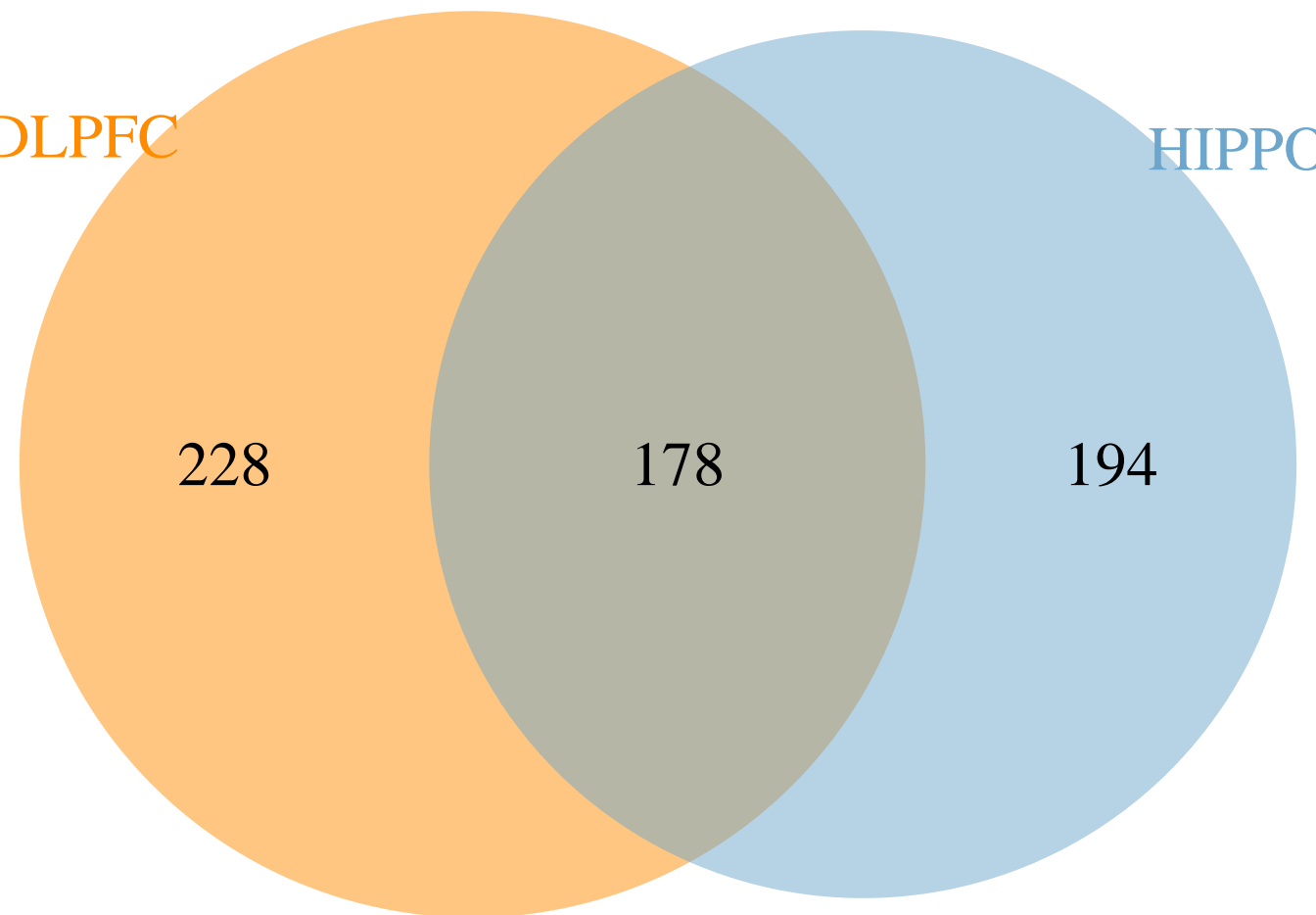
Features with TWAS FDR<5% (by gene ID) – exon



Features with TWAS FDR<5% (by gene ID) – jxn

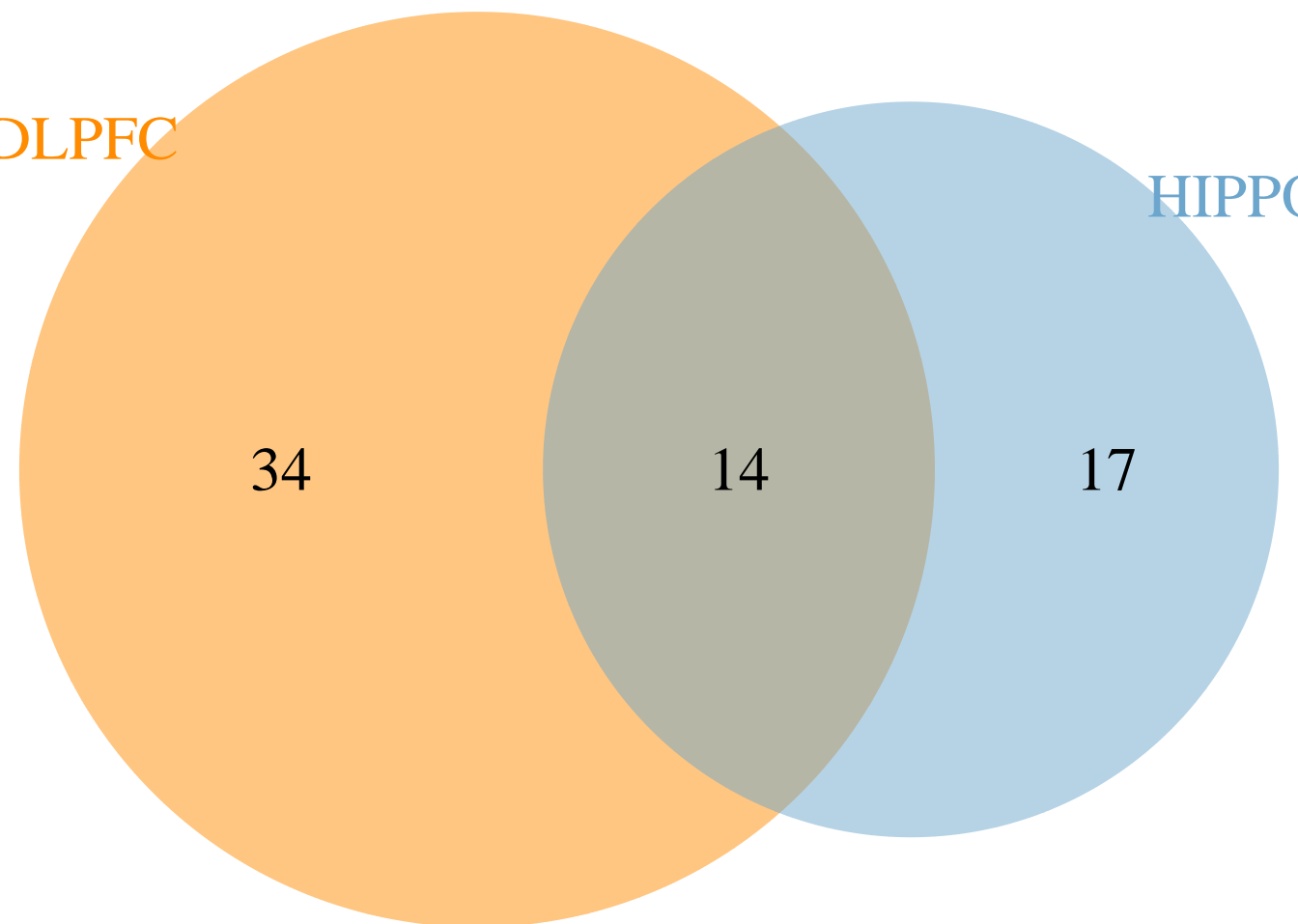


Features with TWAS FDR<5% (by gene ID) – tx

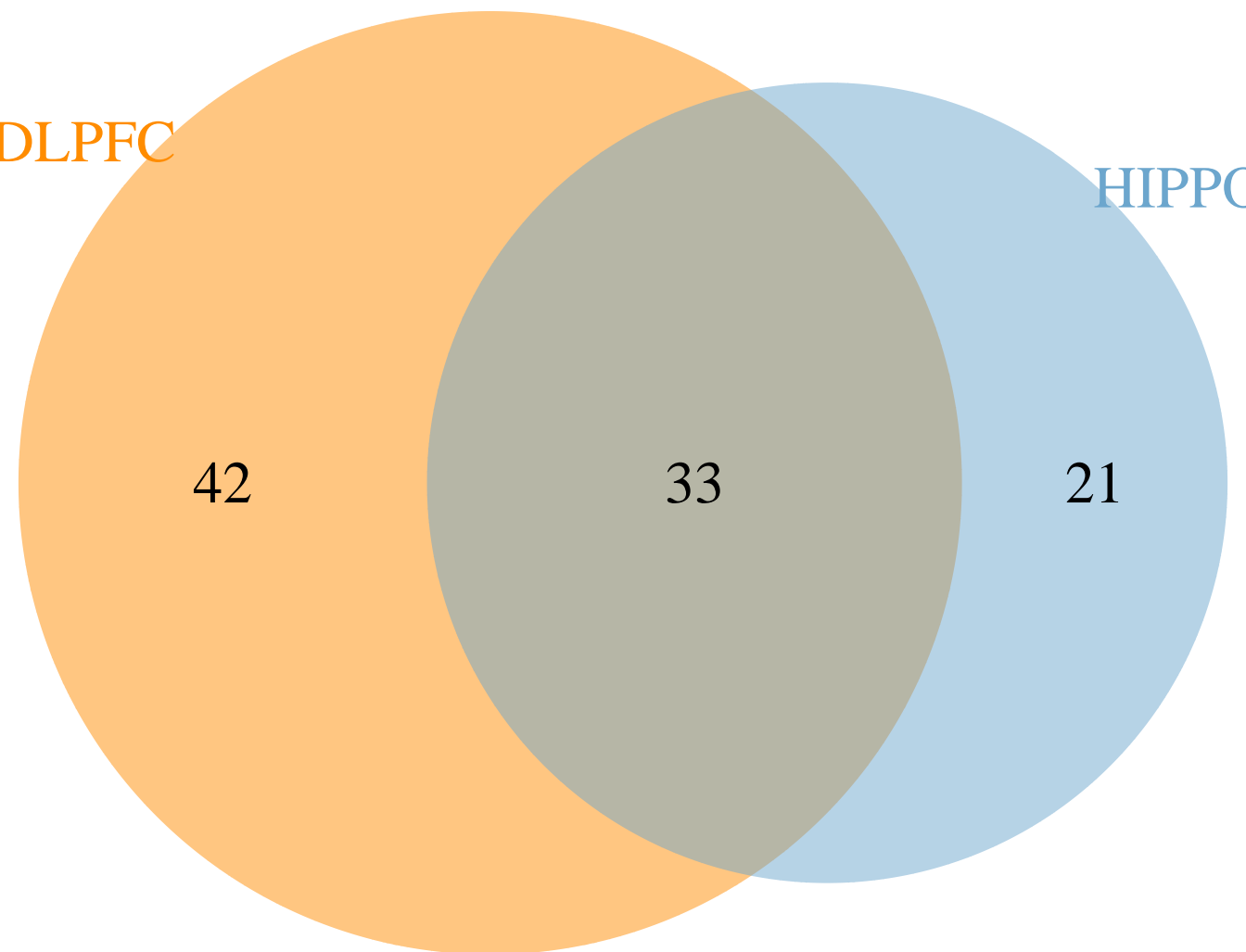




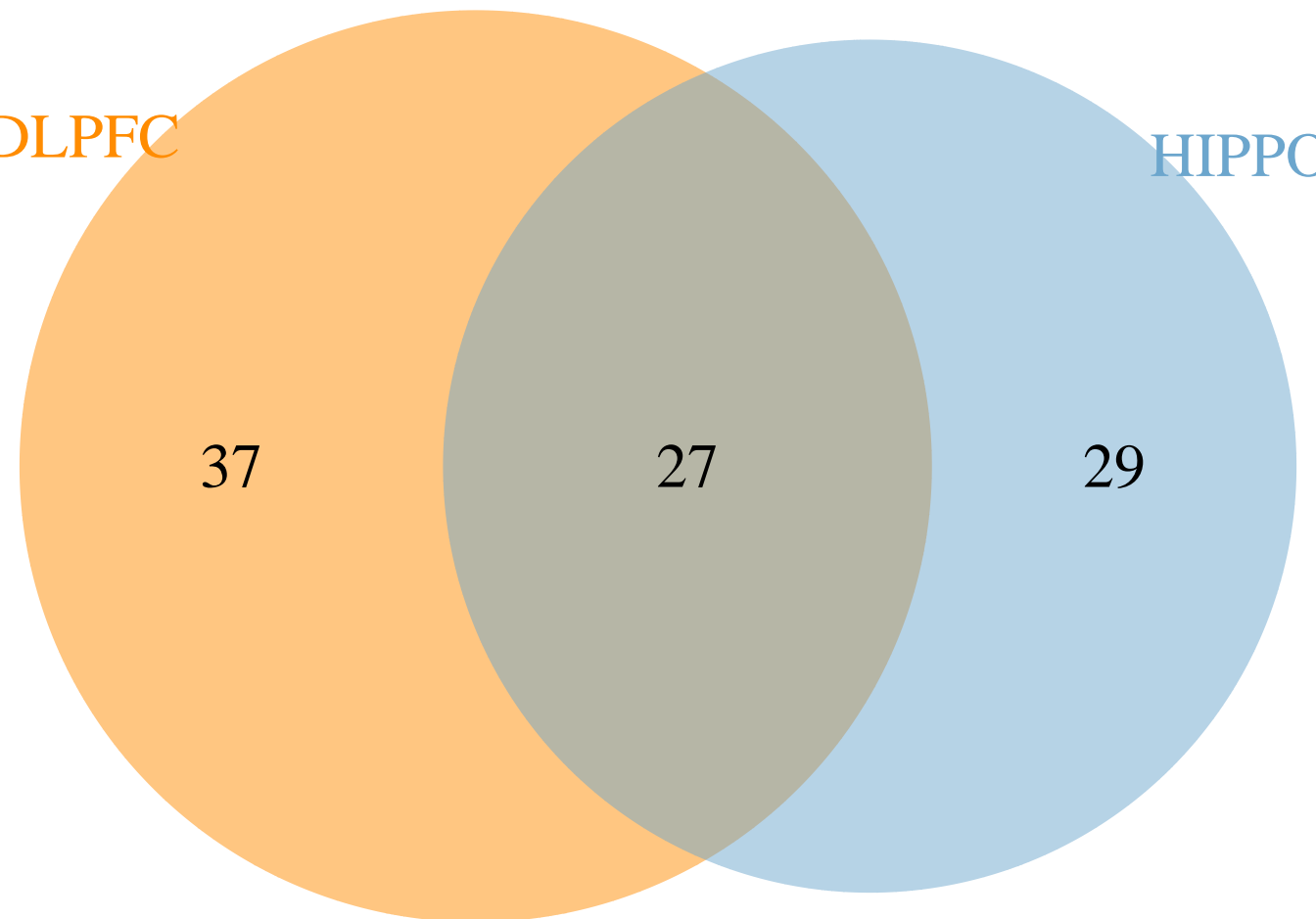
Features with TWAS Bonf<5% (by gene ID) – gene



Features with TWAS Bonf<5% (by gene ID) – exon



Features with TWAS Bonf<5% (by gene ID) – jxn



Features with TWAS Bonf<5% (by gene ID) – tx

