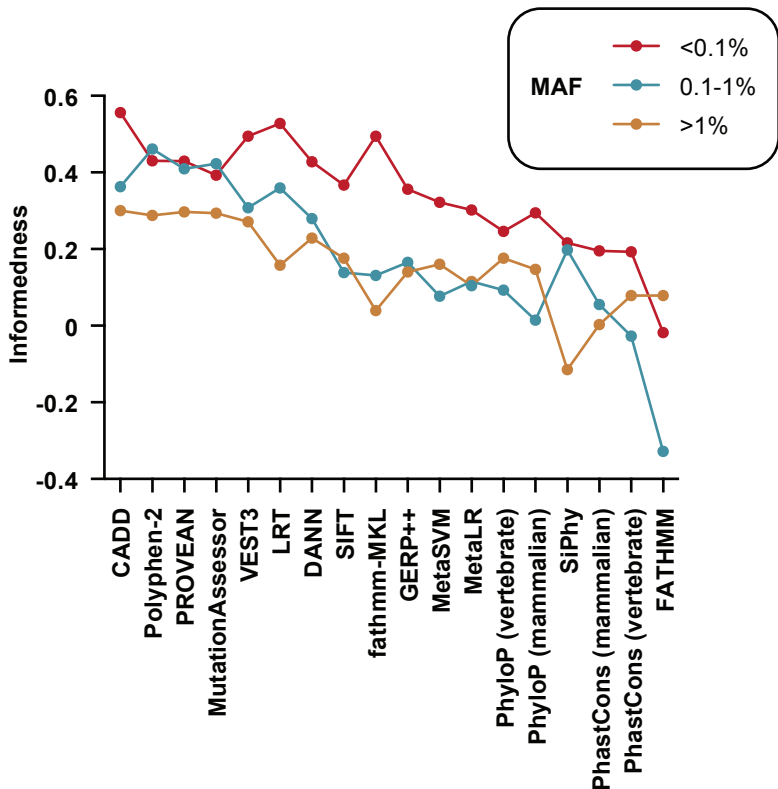# Supplementary Figure 2

Supplementary Figure 2: Functionality prediction algorithms and scores perform overall better on rare compared to common pharmacogenetic variants. Informedness values are shown for variants binned by their minor allele frequency (MAF):<0.1% (n=232, red), 0.1-1% (n=53, blue) and >1% (n=52, yellow)