Altered in 952 (74.32%) of 1281 samples. 19 TMB 184 No. of samples PER3 14% PER2 14% GPR50 13% CYP2C19 13% MTNR1B 9% CLOCK 8% CYP1A2 8% **RORA** 7% **ARNTL** 7% MTNR1A 7% **ASMT** 6% Cancer Cancer Missense_Mutation Splice_Site Nonsense Mutation Nonstop_Mutation ACC LUSC PCPG • SKCM COAD HNSC LAML THYM Frame_Shift_Ins Translation_Start_Site BLCA DLBC KICH LGG MESO PRAD STAD UCEC In_Frame_Del Multi_Hit BRCA **ESCA** KIRC LIHC OV ■ READ ■ TGCT UCS Frame_Shift_Del CESC **GBM** KIRP LUAD PAAD SARC THCA NA