1206 deleterious variants across all UGT genes 1e-01 log10(GMAF) of deleterious variants 1e-02 · Variant ID (GMAF>0.01) 2-234668879-C-CAT 2-234668879-C-CATAT 1e-03 -4-70462042-C-T 4-70512787-A-T 1e-04 · 1e-05 34 51 29 127 134 128 UGT1A3 UGT1A6 UGT1A7 UGT1A8 UGT1A9 UGT2A2 UGT2A3 UGT2B4 UGT2B7 UGT3A1 UGT3A2 UGT2A1 UGT1A4 UGT1A5 UGT1A10 UGT2B10 UGT2B15 UGT2B28 UGT1A1 UGT2B11 UGT2B17