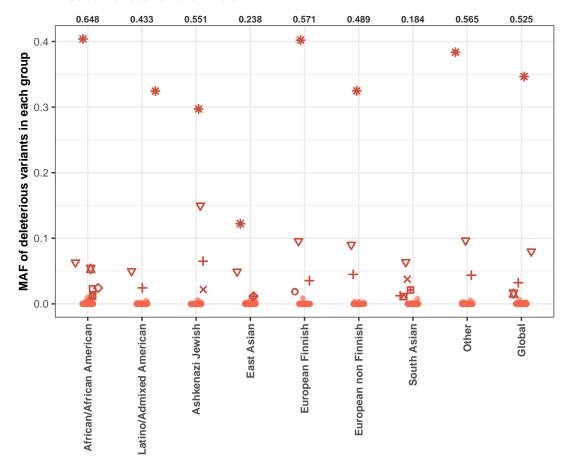
## **Deleterious variants in all UGT genes**

1206 deleterious variants in total



## Variant ID (MAF>0.01) & Gene(s)

- \* 2-234668879-C-CAT (UGT1A1)
- 2-234668879-C-CATAT (UGT1A1)
- 2-234638282-G-GT (UGT1A3)
- 2-234622331-GC-G (UGT1A5)
- 2-234676872-C-T (UGT1A[1-10])
- + 4-70512787-A-T (UGT2A1)
- ▼ 4-70462042-C-T (UGT2A1, UGT2A2)
- × 4–69811110–A–C (UGT2A3)
- ♦ 4–69693141–GT–G (UGT2B10)
- □ 4-70078393-C-T (UGT2B11)
- 4-69536234-G-T (UGT2B15)
- 4-69512937-T-A (UGT2B15)