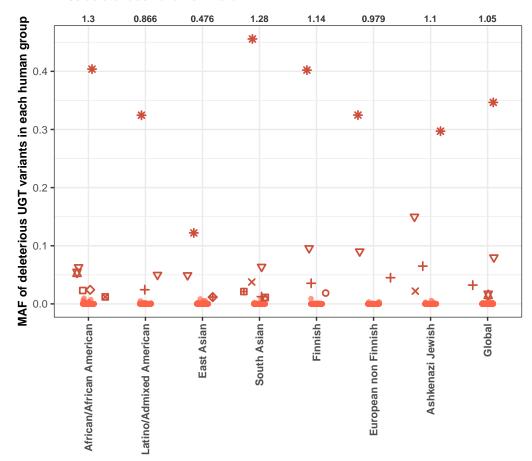
Deleterious variants in all UGT genes per population

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 (UGT2A[1-2])
- × 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)