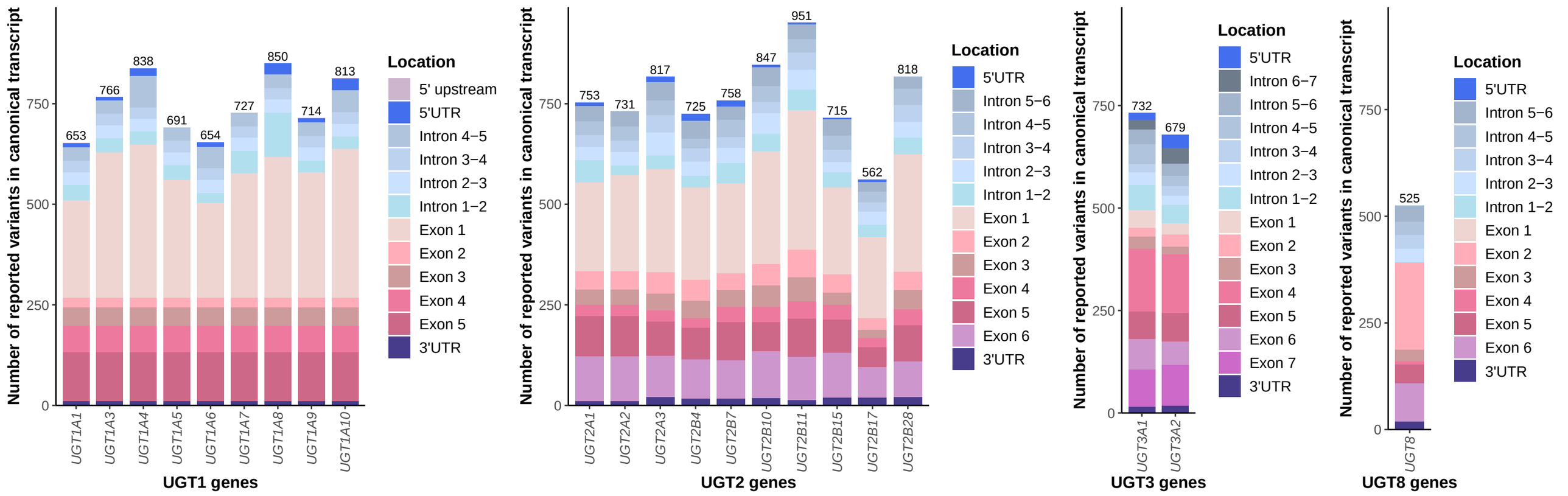
Number of reported variants in canonical transcript
0
250
500
750
653
766
838
691
654
727
850
714
813
UGT1A1
UGT1A3
UGT1A4
UGT1A5
UGT1A6
UGT1A7
UGT1A8
UGT1A9
UGT1A10
UGT1 genes
Location
5' upstream
5'UTR
Intron 4−5
Intron 3−4
Intron 2−3
Intron 1−2
Exon 1
Exon 2
Exon 3
Exon 4
Exon 5
3'UTR
753
731
817
725
758
847
951
715
562
818
UGT2A1
UGT2A2
UGT2A3
UGT2B4
UGT2B7
UGT2B10
UGT2B11
UGT2B15
UGT2B17
UGT2B28
UGT2 genes
Location
5'UTR
Intron 5−6
Intron 4−5
Intron 3−4
Intron 2−3
Intron 1−2
Exon 1
Exon 2
Exon 3
Exon 4
Exon 5
Exon 6
3'UTR
732
679
UGT3A1
UGT3A2
UGT3 genes
Location
5'UTR
Intron 6−7
Intron 5−6
Intron 4−5
Intron 3−4
Intron 2−3
Intron 1−2
Exon 1
Exon 2
Exon 3
Exon 4
Exon 5
Exon 6
Exon 7
3'UTR
525
UGT8
UGT8 genes
Location
5'UTR
Intron 5−6
Intron 4−5
Intron 3−4
Intron 2−3
Intron 1−2
Exon 1
Exon 2
Exon 3
Exon 4
Exon 5
Exon 6
3'UTR