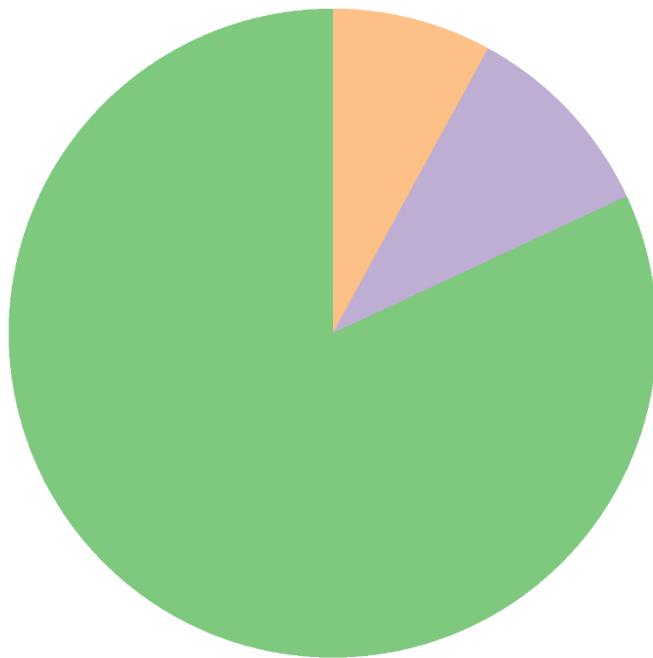
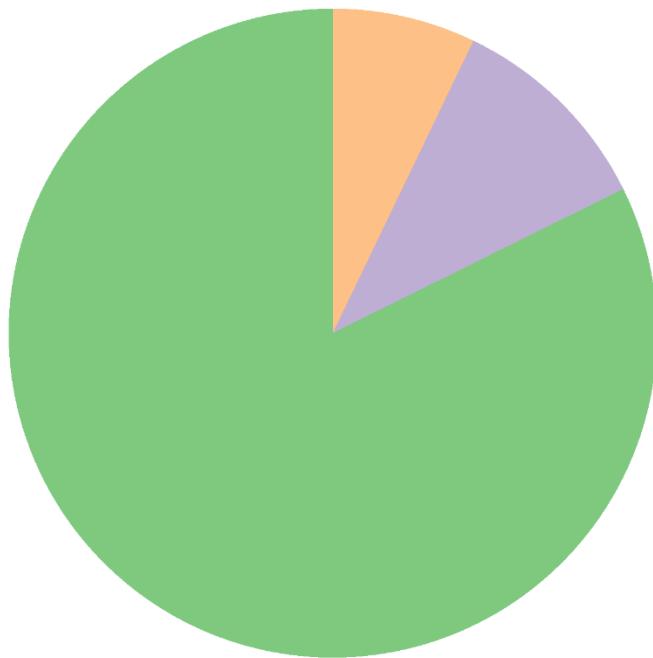


Percent of genome regions (M1)



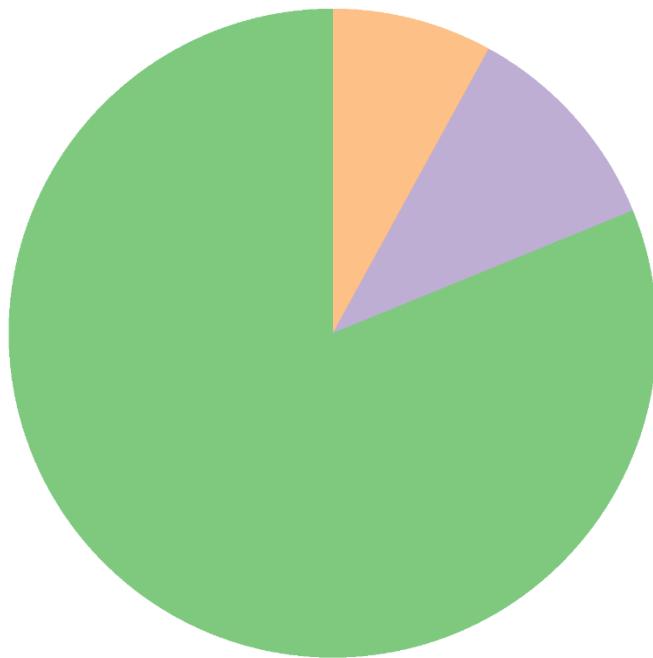
exon	(4939096430, 81.94%)
intron	(478984523, 7.95%)
intergenic	(609917039, 10.12%)

Percent of genome regions (M2)



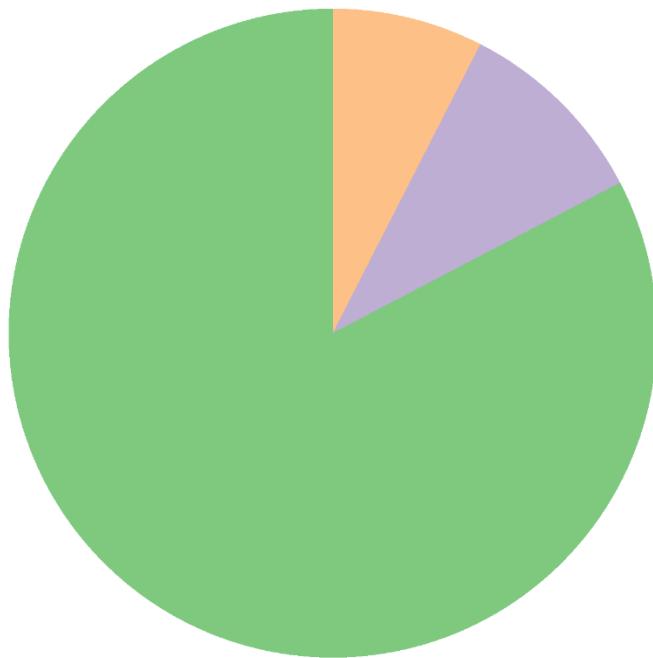
exon	(4595927734, 82.33%)
intron	(398816678, 7.14%)
intergenic	(587834840, 10.53%)

Percent of genome regions (M3)



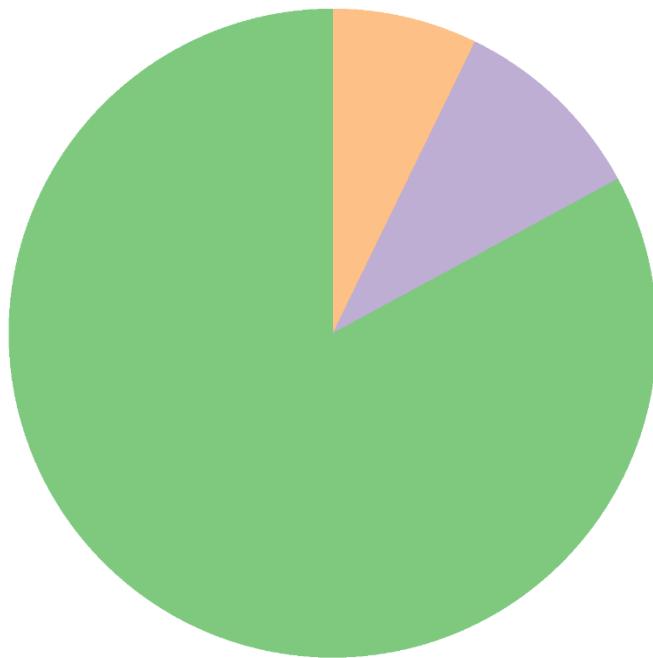
exon	(5094775364, 81.17%)
intron	(501171659, 7.98%)
intergenic	(680547853, 10.84%)

Percent of genome regions (N1)



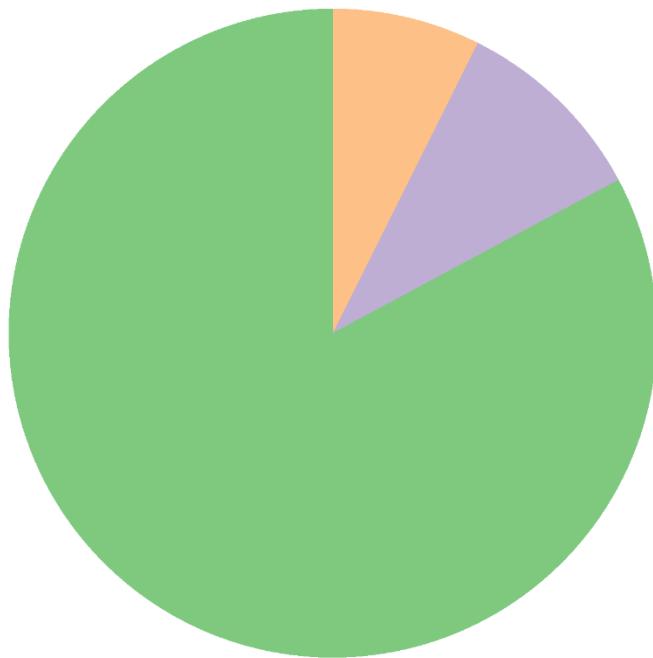
[Green square]	exon (5027731318, 82.67%)
[Orange square]	intron (457433097, 7.52%)
[Purple square]	intergenic (596388960, 9.81%)

Percent of genome regions (N2)



[Green square]	exon (4879161053, 82.90%)
[Orange square]	intron (425038501, 7.22%)
[Purple square]	intergenic (581653553, 9.88%)

Percent of genome regions (N3)



[Green square]	exon (5337477366, 82.83%)
[Orange square]	intron (475541333, 7.38%)
[Purple square]	intergenic (630581129, 9.79%)