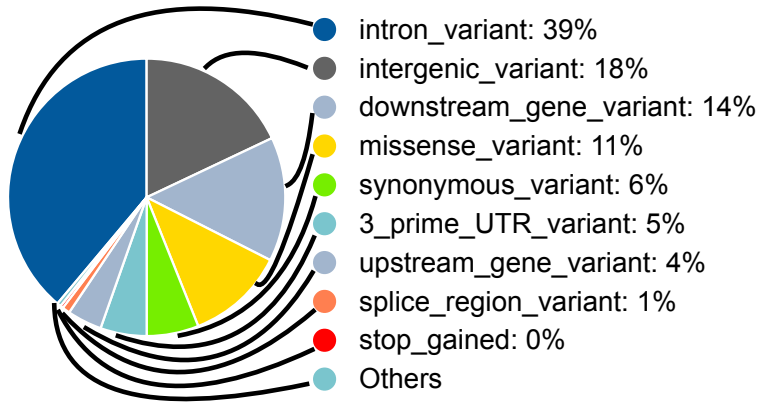
Category	Count
Variants processed	774
Variants filtered out	0
Novel / existing variants	773 (99.9) / 1 (0.1)
Overlapped genes	557
Overlapped transcripts	589
Overlapped regulatory features	-

Consequences (all)



Coding consequences