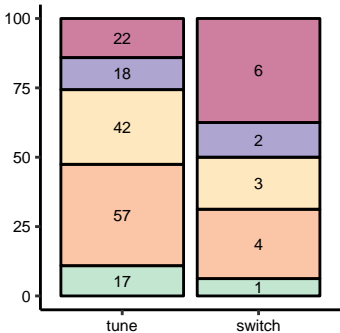


% of Variants



accessible



coding



intergenic



intronic