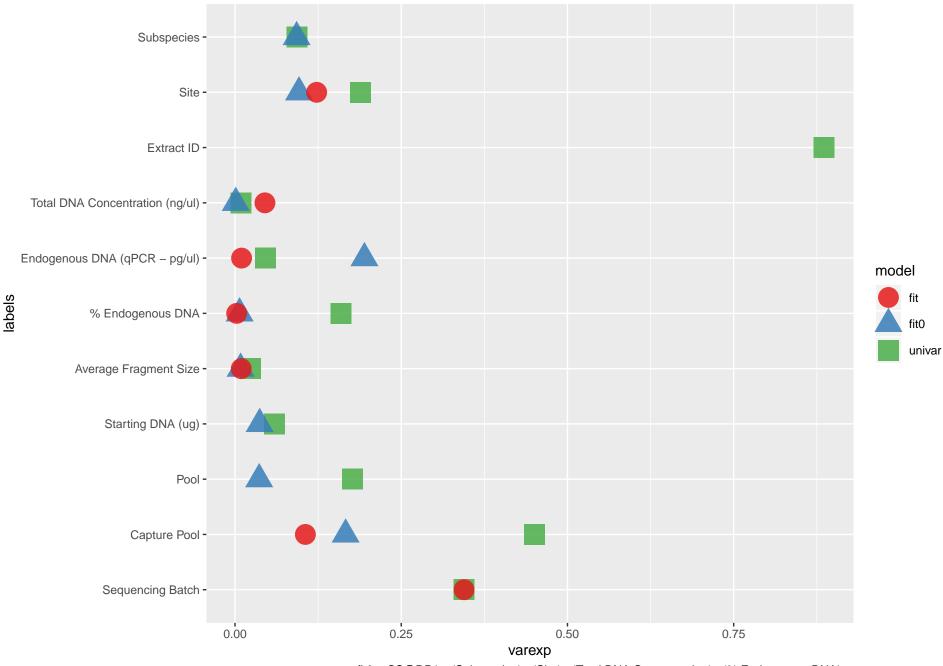
Variance explained in capture sensitivity derived from univariate and type I multivariate ANOVAs



fit0 = CS@DP1 ~ 'Subspecies' + 'Site' + 'Total DNA Concentration' + '% Endogenous DNA' + 'Average Fragment Size' + 'Starting DNA' + 'Pool' + 'Capture Pool' + 'Sequencing Batch'