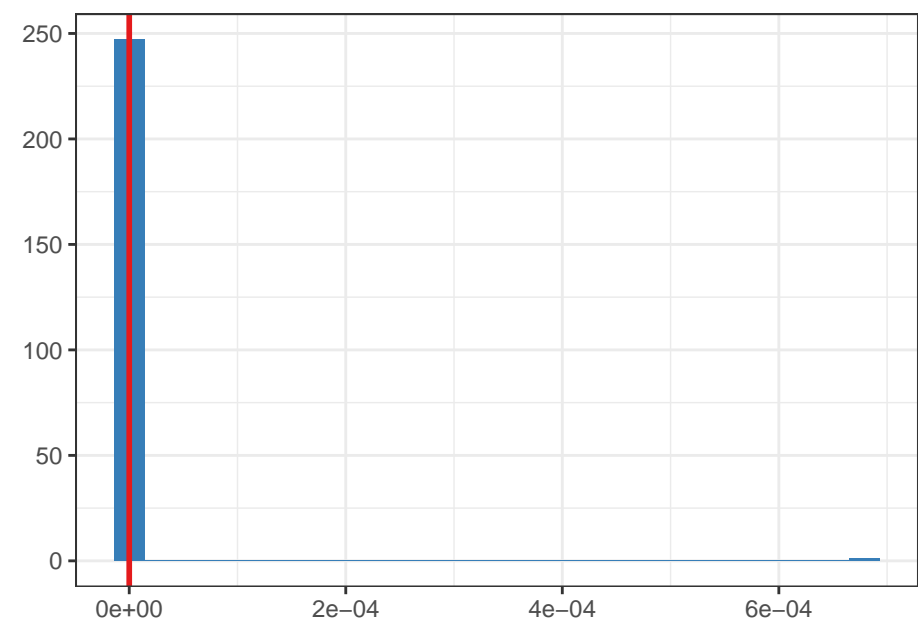
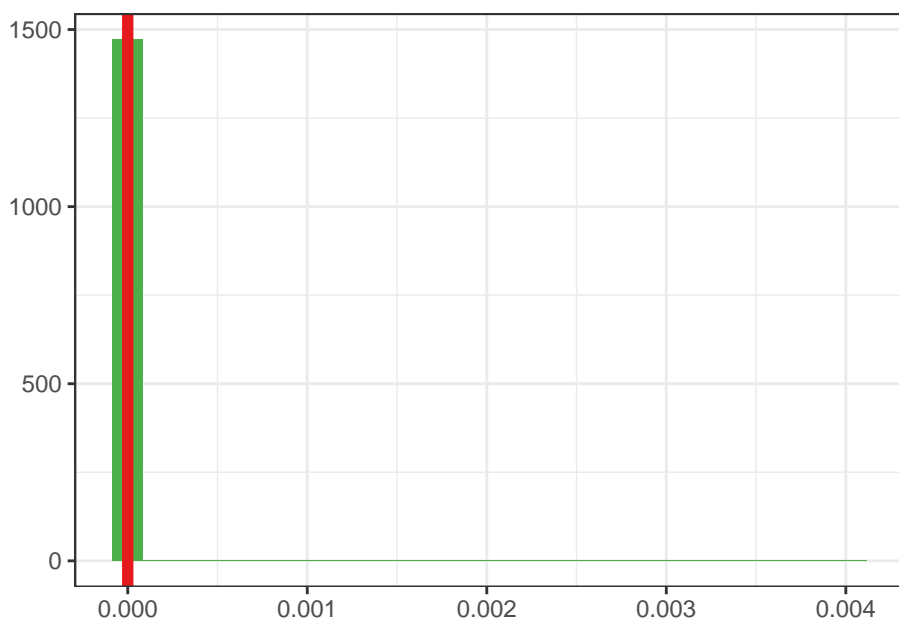


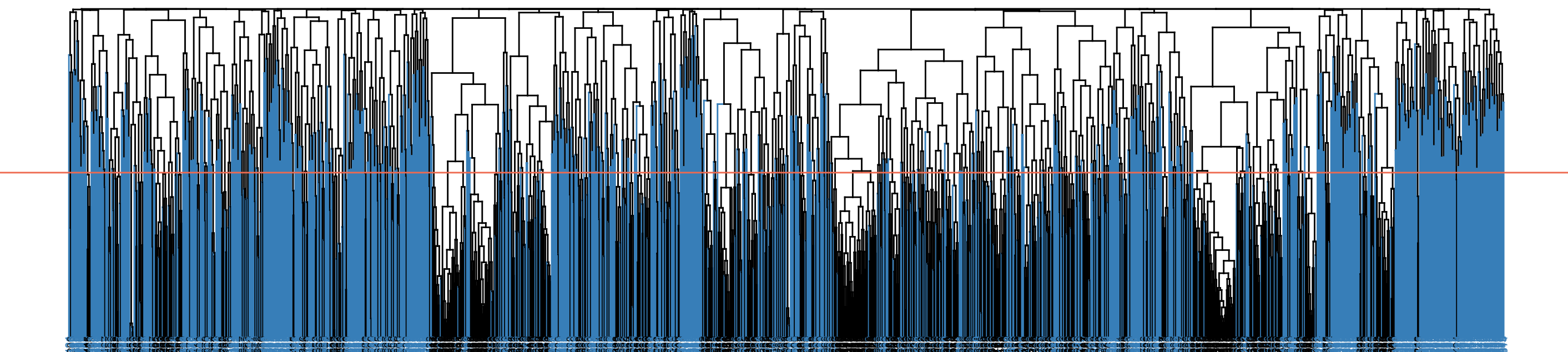
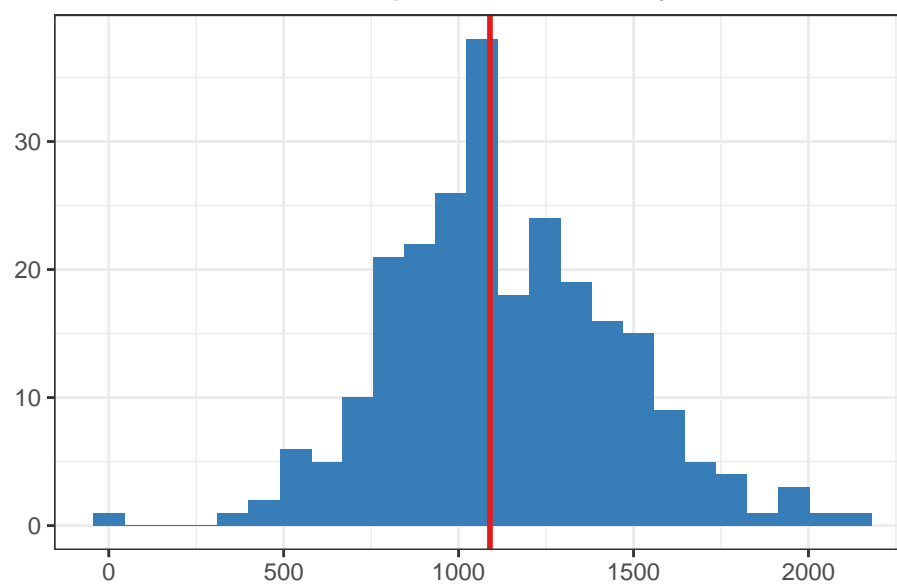
sample missingness



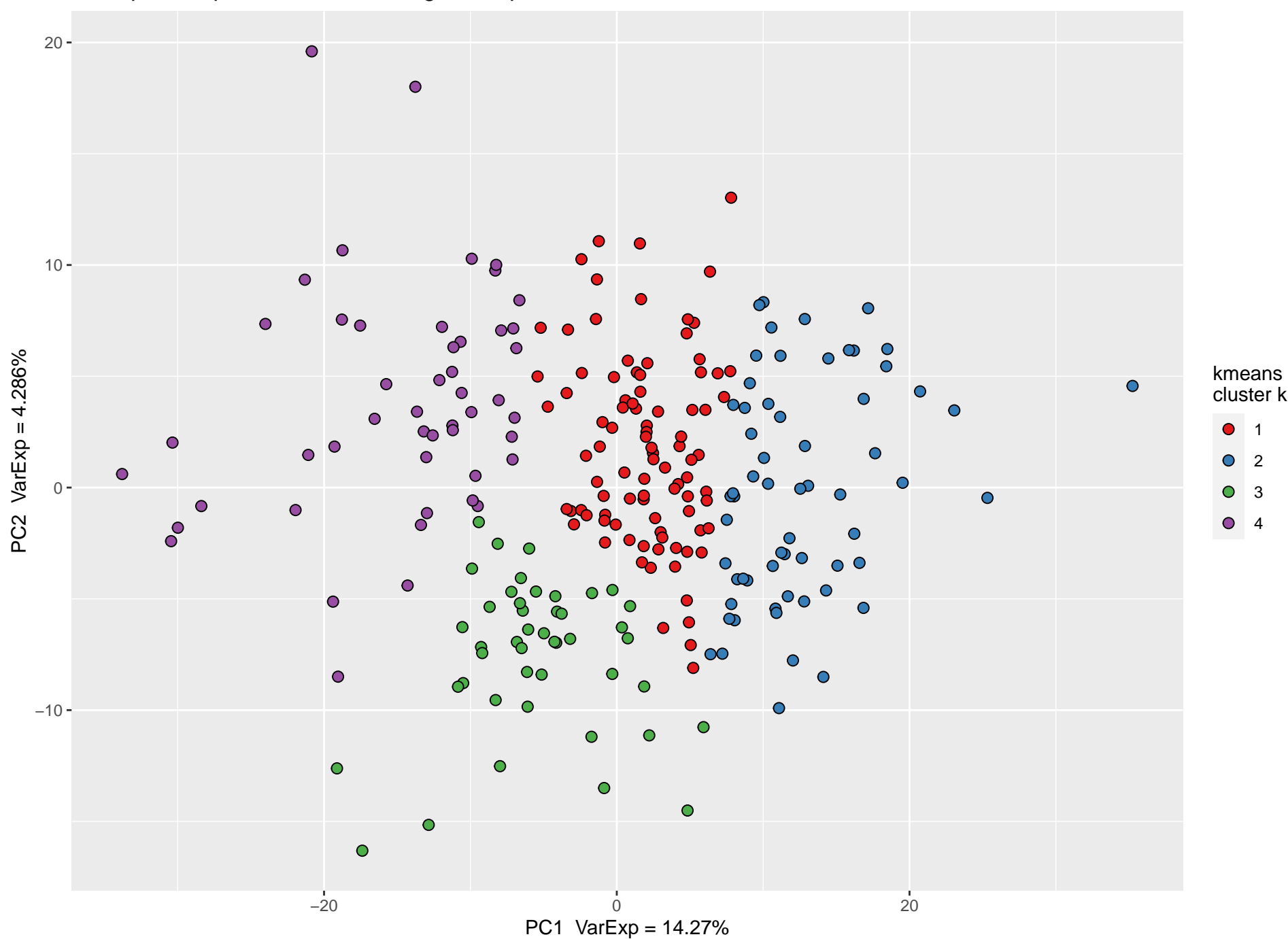
feature missingness



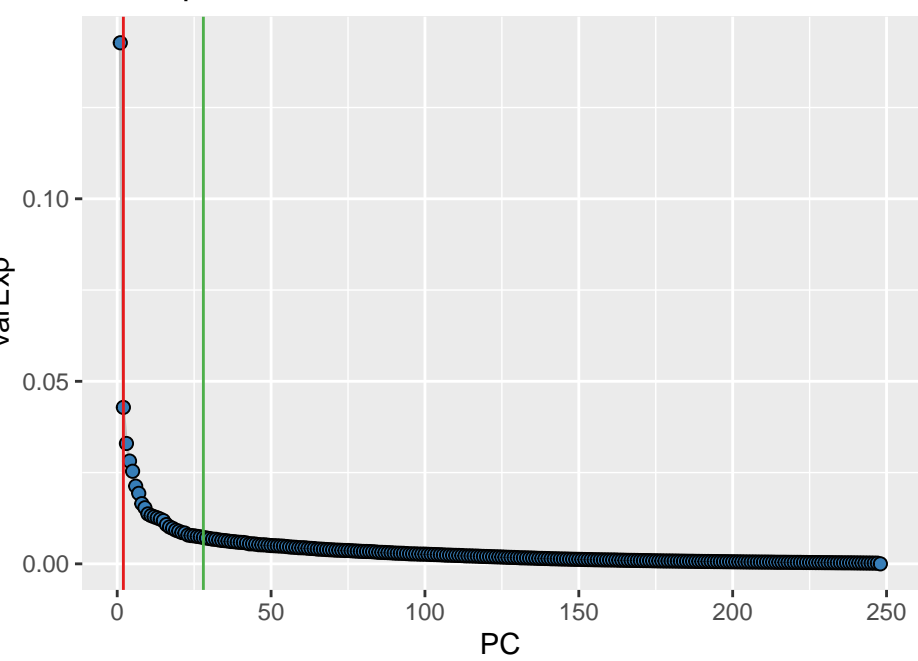
total abundance of samples
at complete features only



Principal components 1-&-2 using 808 representative metabolites



Scree plot



data.reduction	count
total metabolite count	1472
metabolites included in data reduction	1472
number of metabolite clusters	808
number of representative metabolites	808