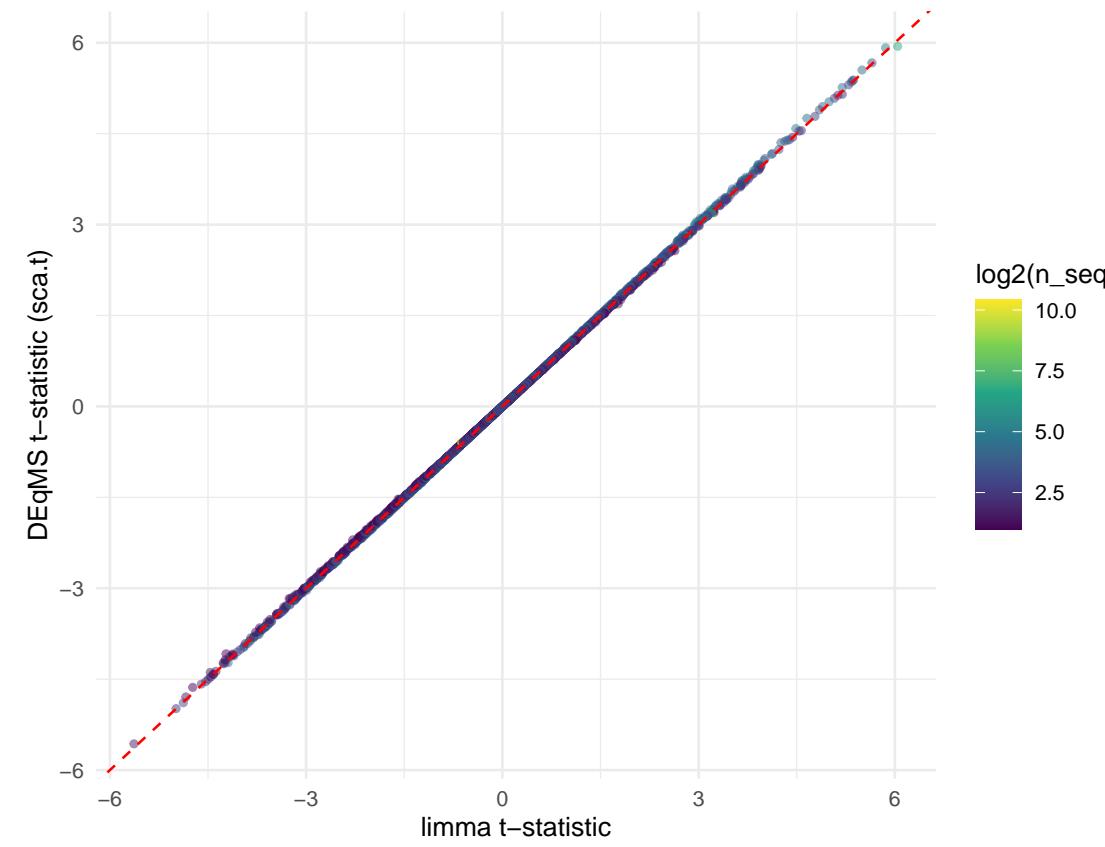


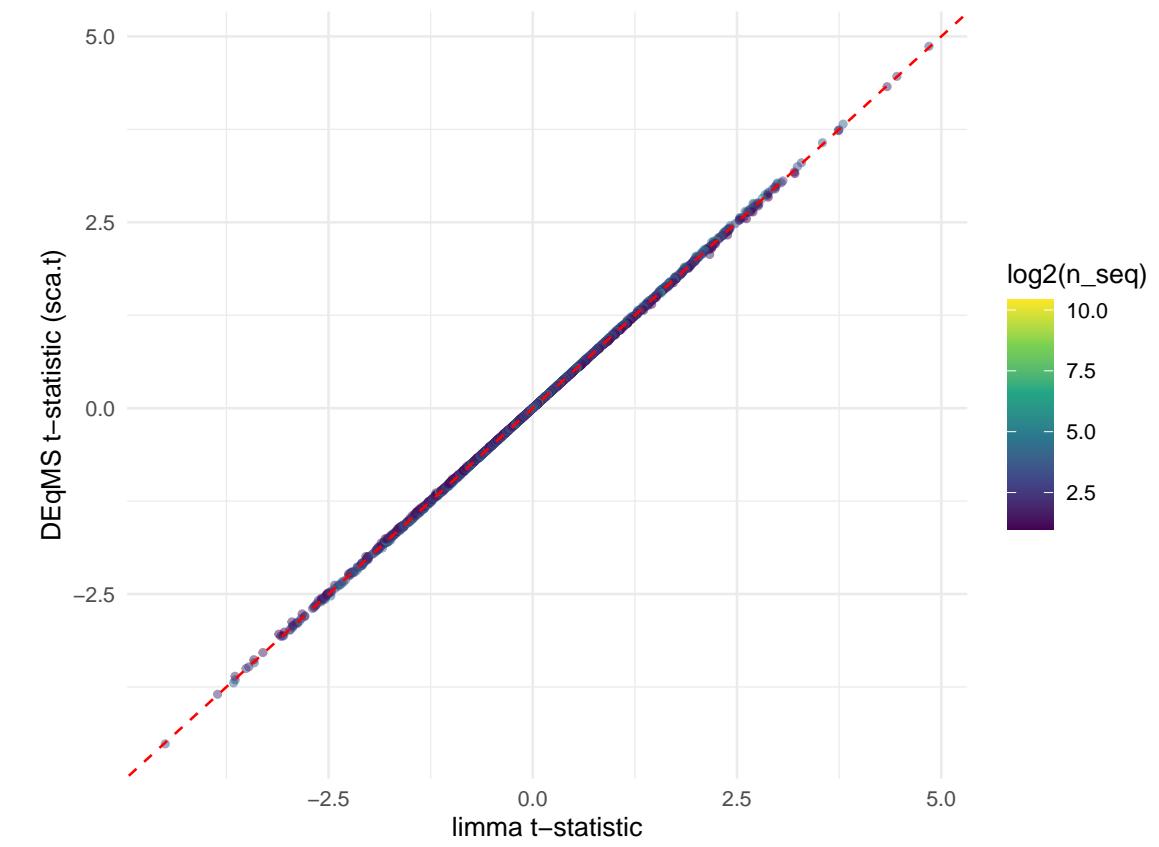
DEqMS vs limma t-statistics

Points colored by log2(peptide count); dashed = identity line

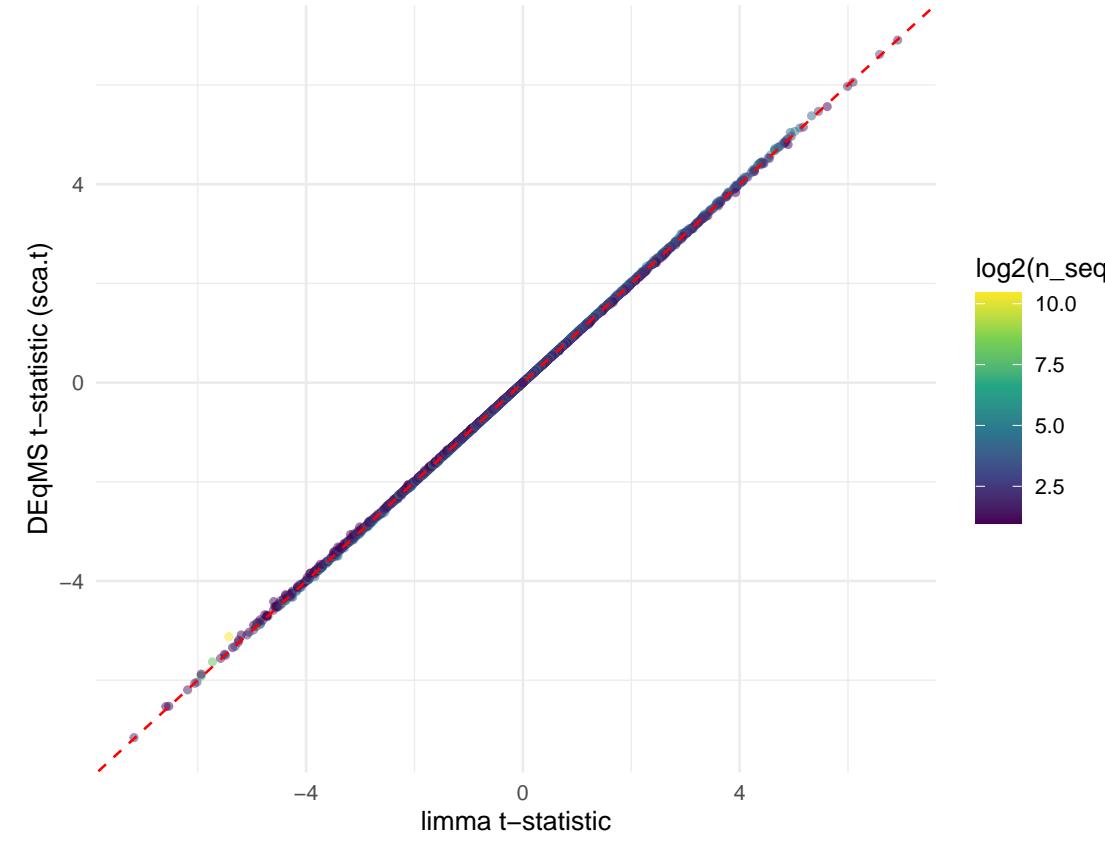
Training_Young



Training_Old



Aging



Interaction

