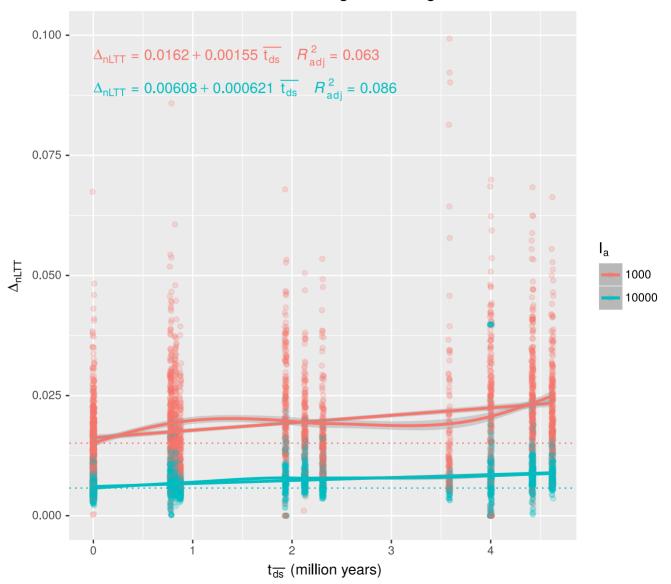
nLTT statistic for different expected mean duration of speciation, for different DNA alignment lengths



 $n = 5000 \ / \ 2742400, figure\_error\_expected\_mean\_dur\_spec\_alignment\_length$