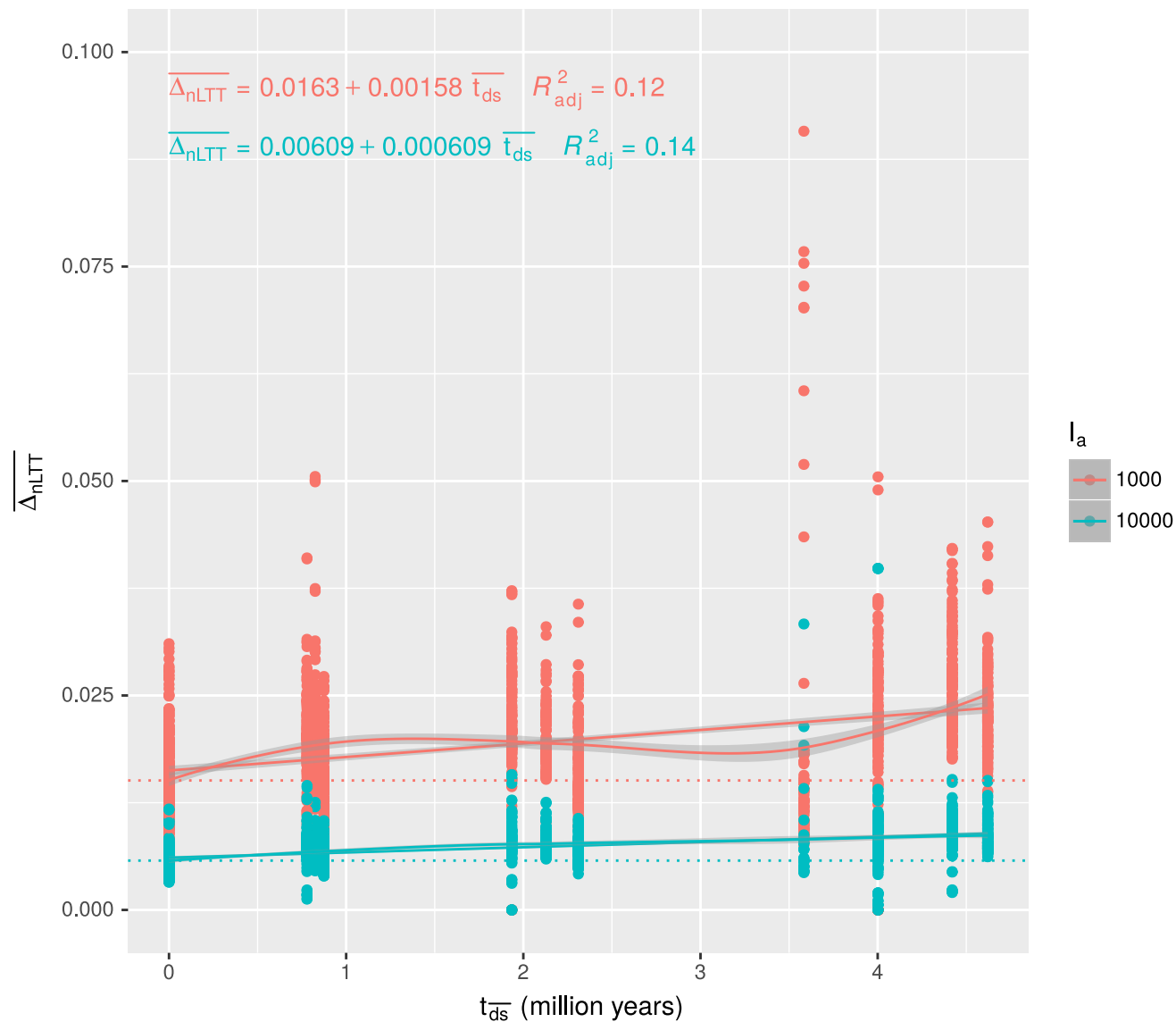


# Mean nLTT statistic for different expected mean duration of speciation, for different DNA alignment lengths



figure\_error\_expected\_mean\_dur\_spec\_mean\_alignment\_length