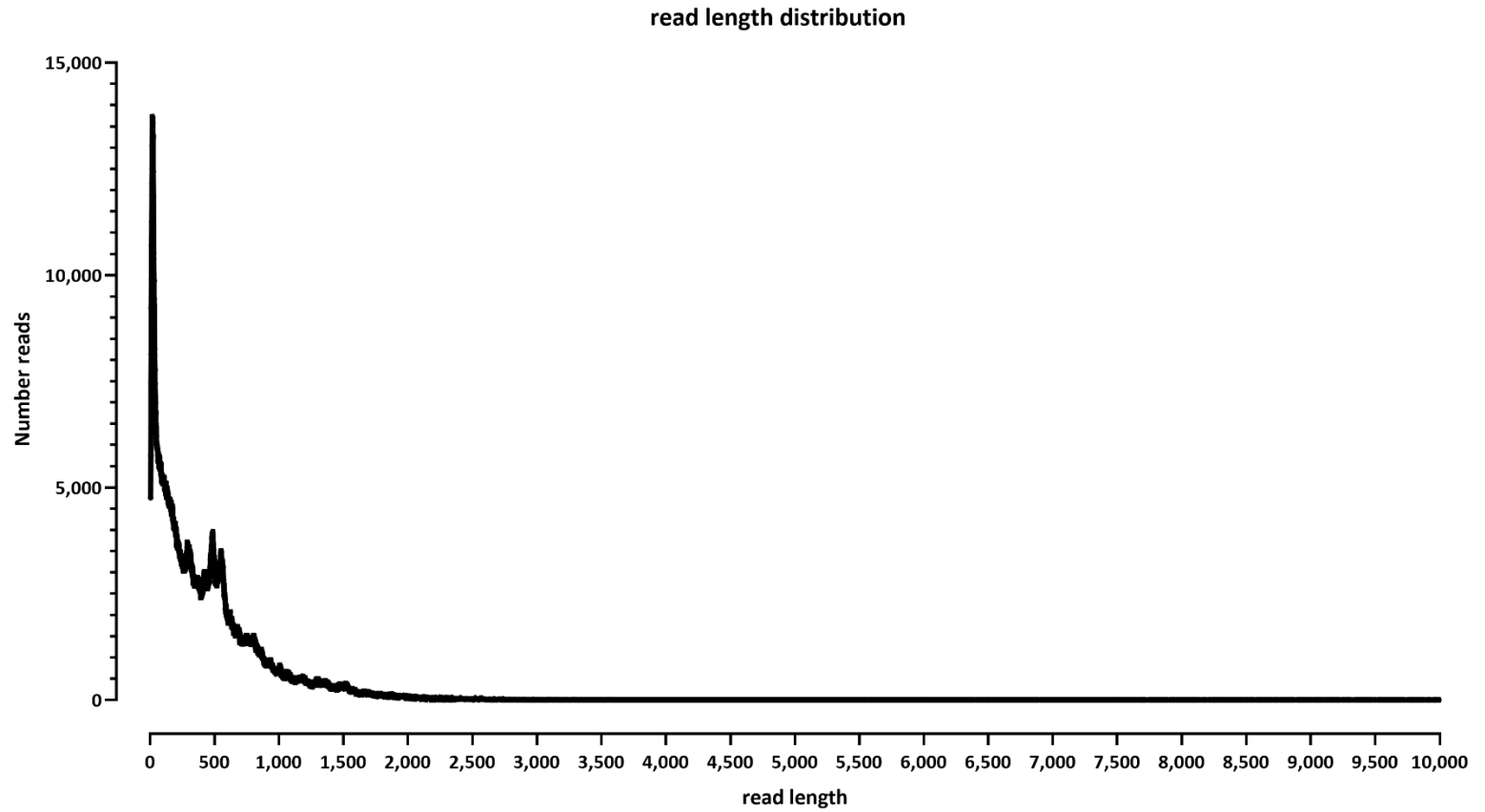
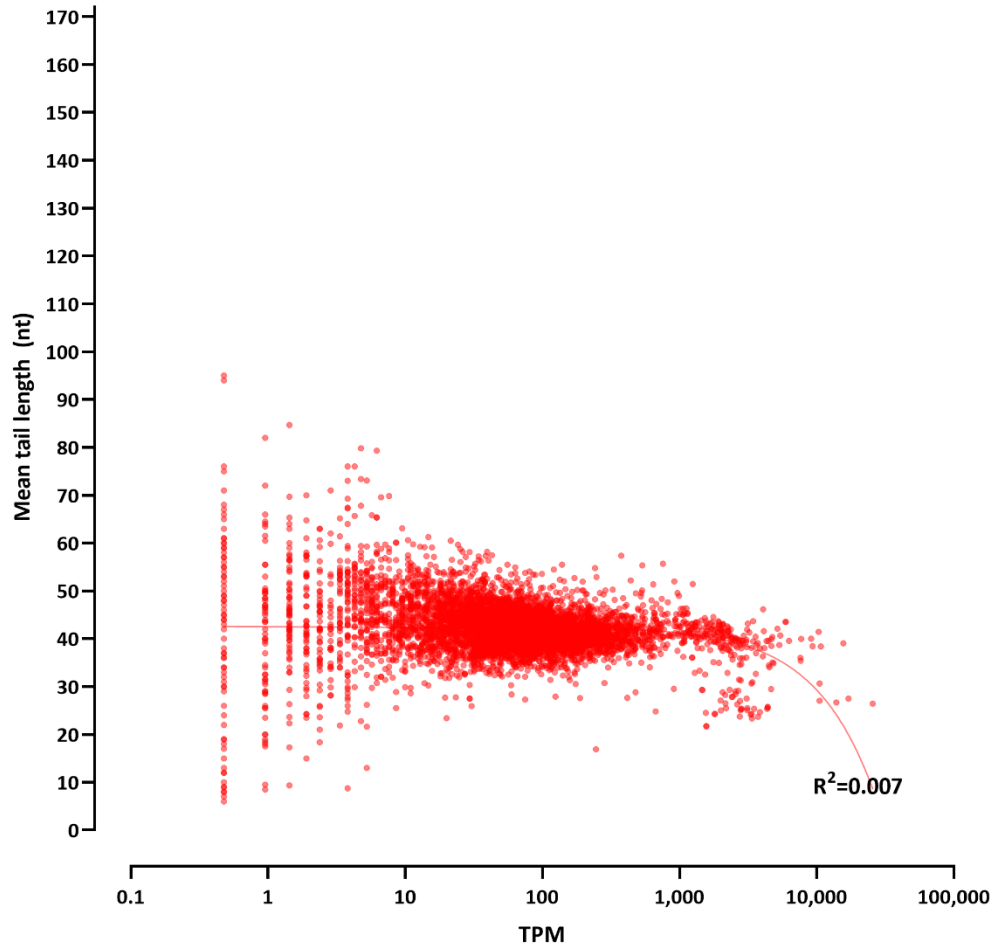


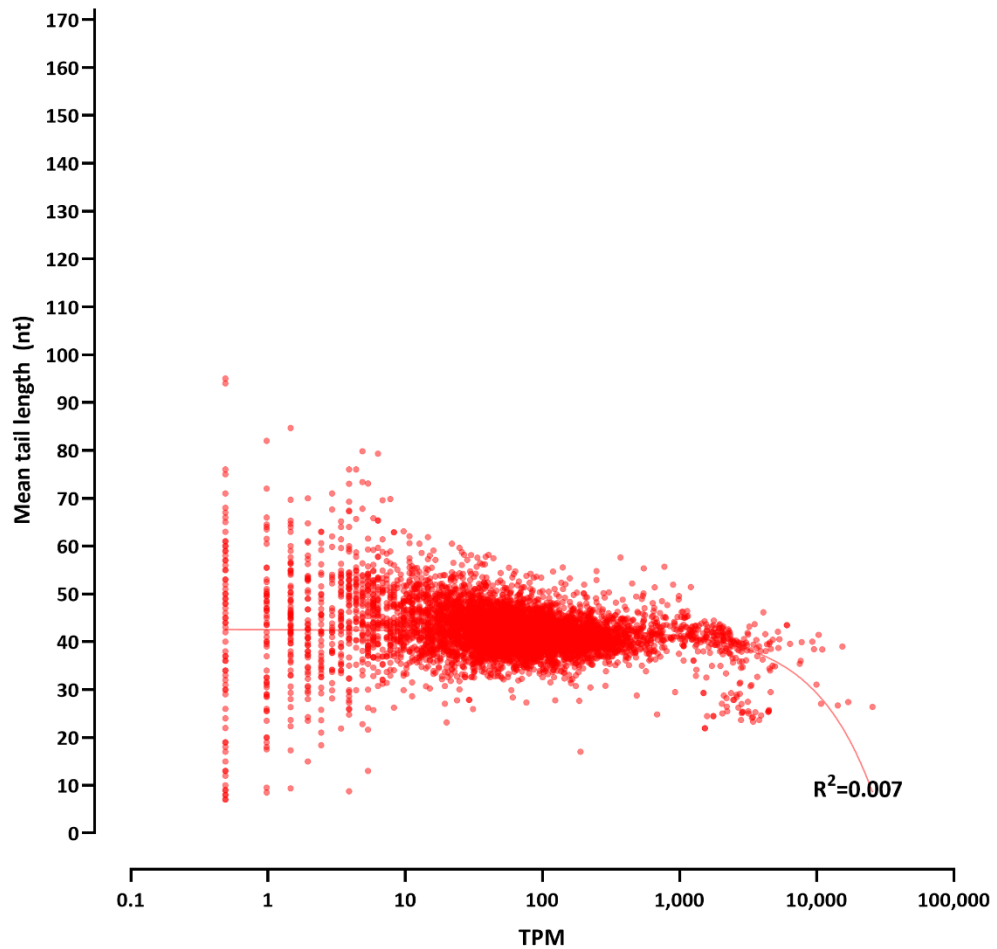
# READ LENGTH DISTRIBUTION



250228\_exp240718\_MS798\_B2



250228\_exp240718\_MS798\_B2 reads >= 100bp



# READ COVERAGE

Coverage of genes; >1000bp, sacCer3

