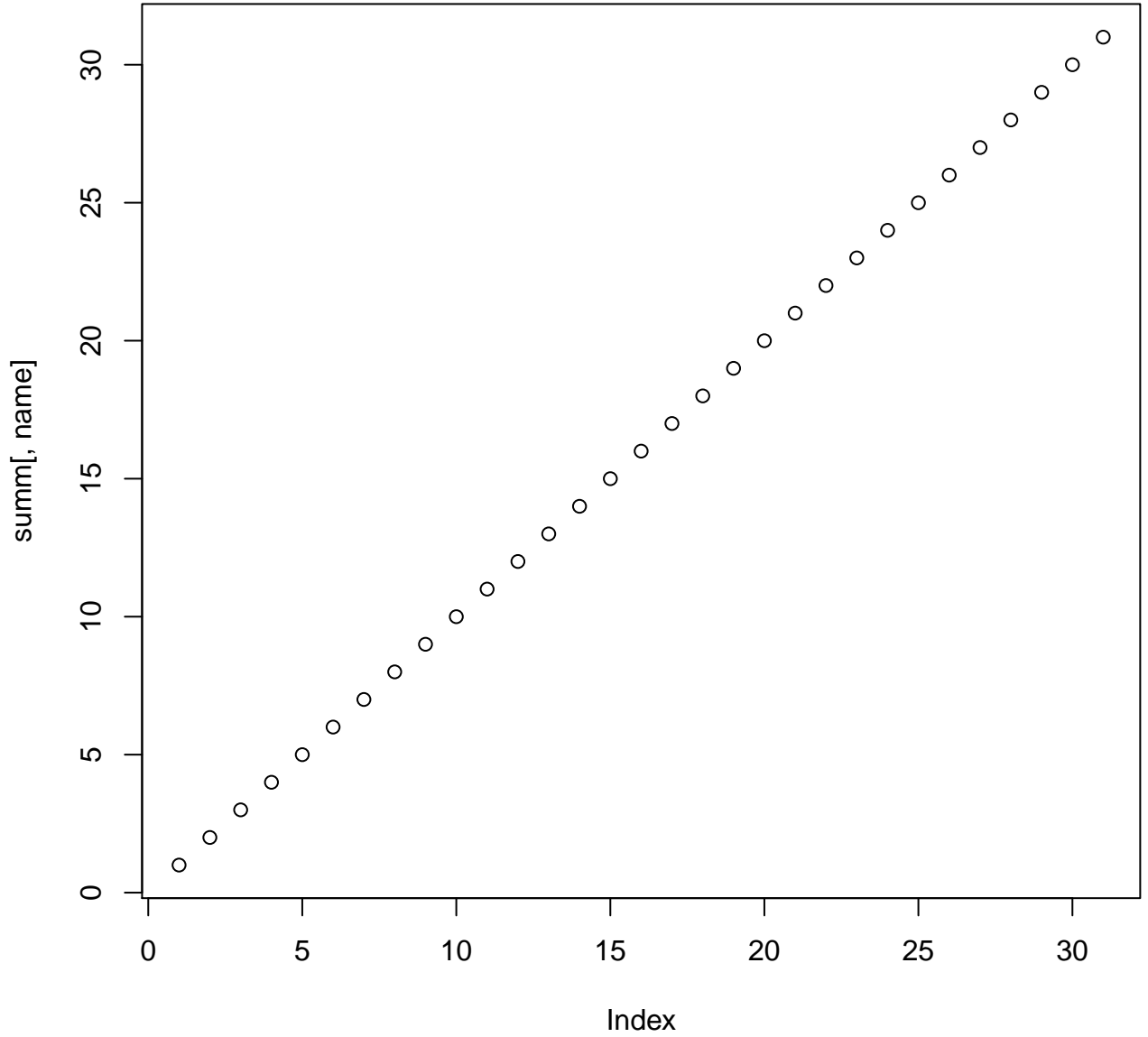
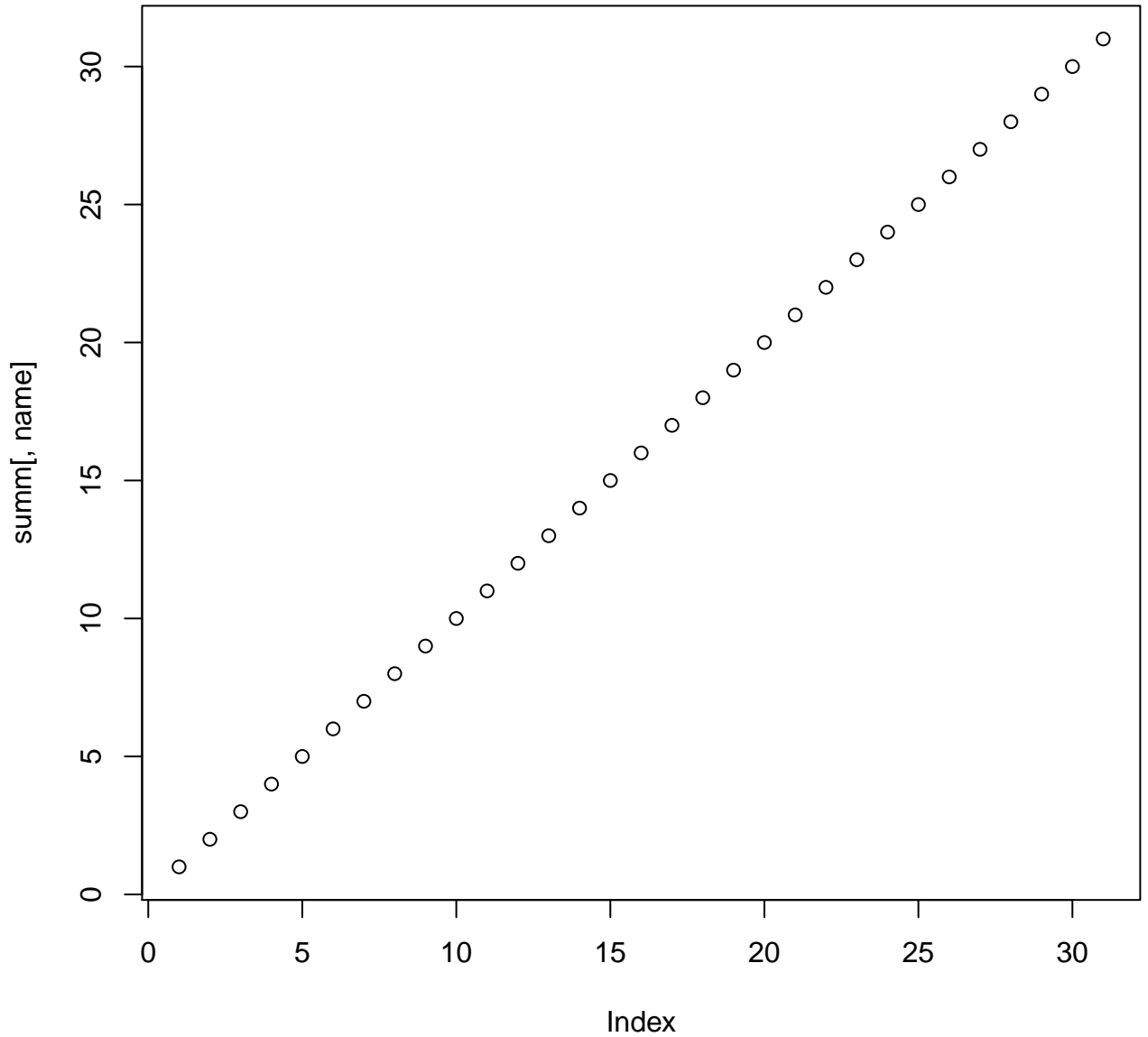


**X**

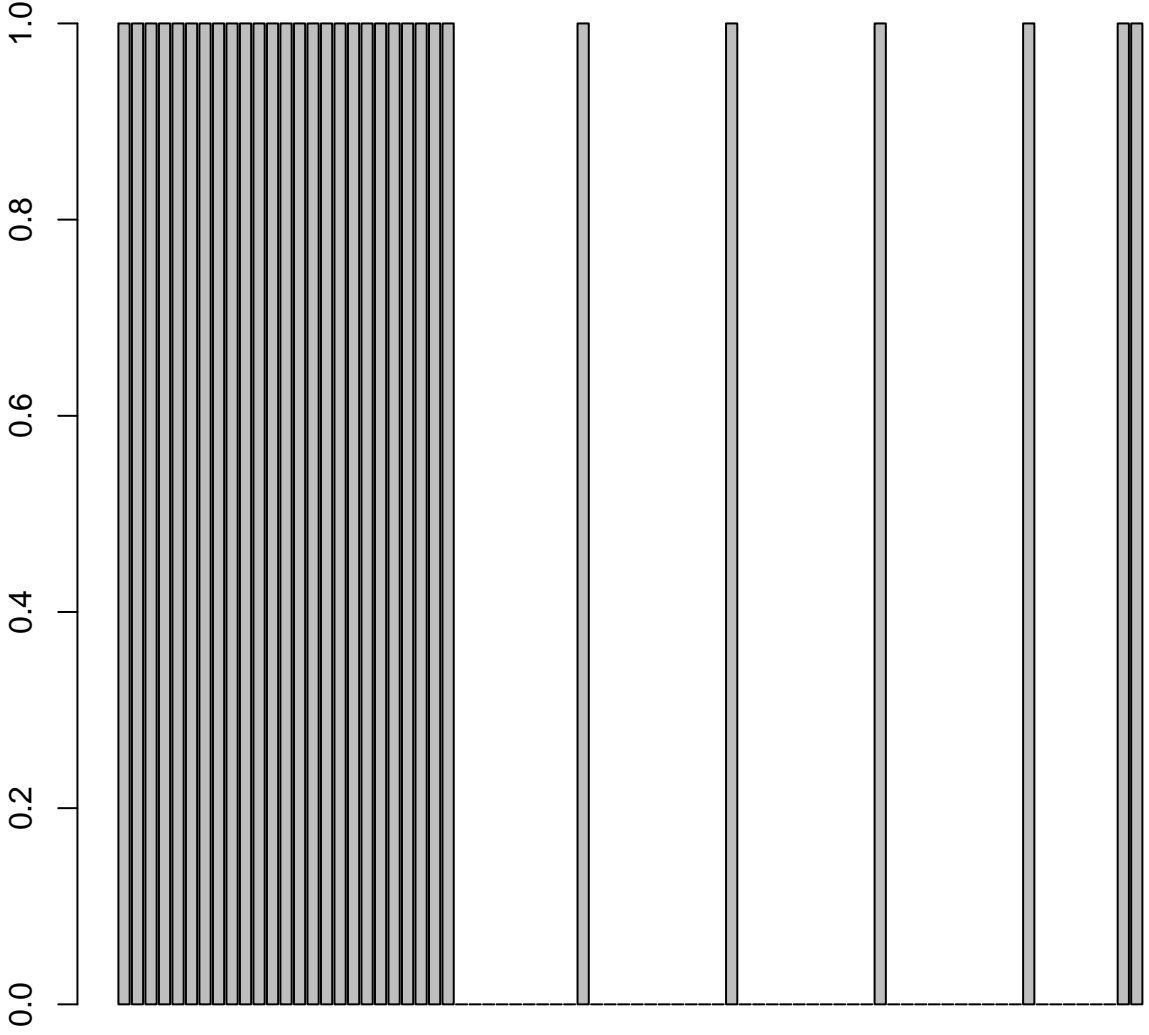


X.1





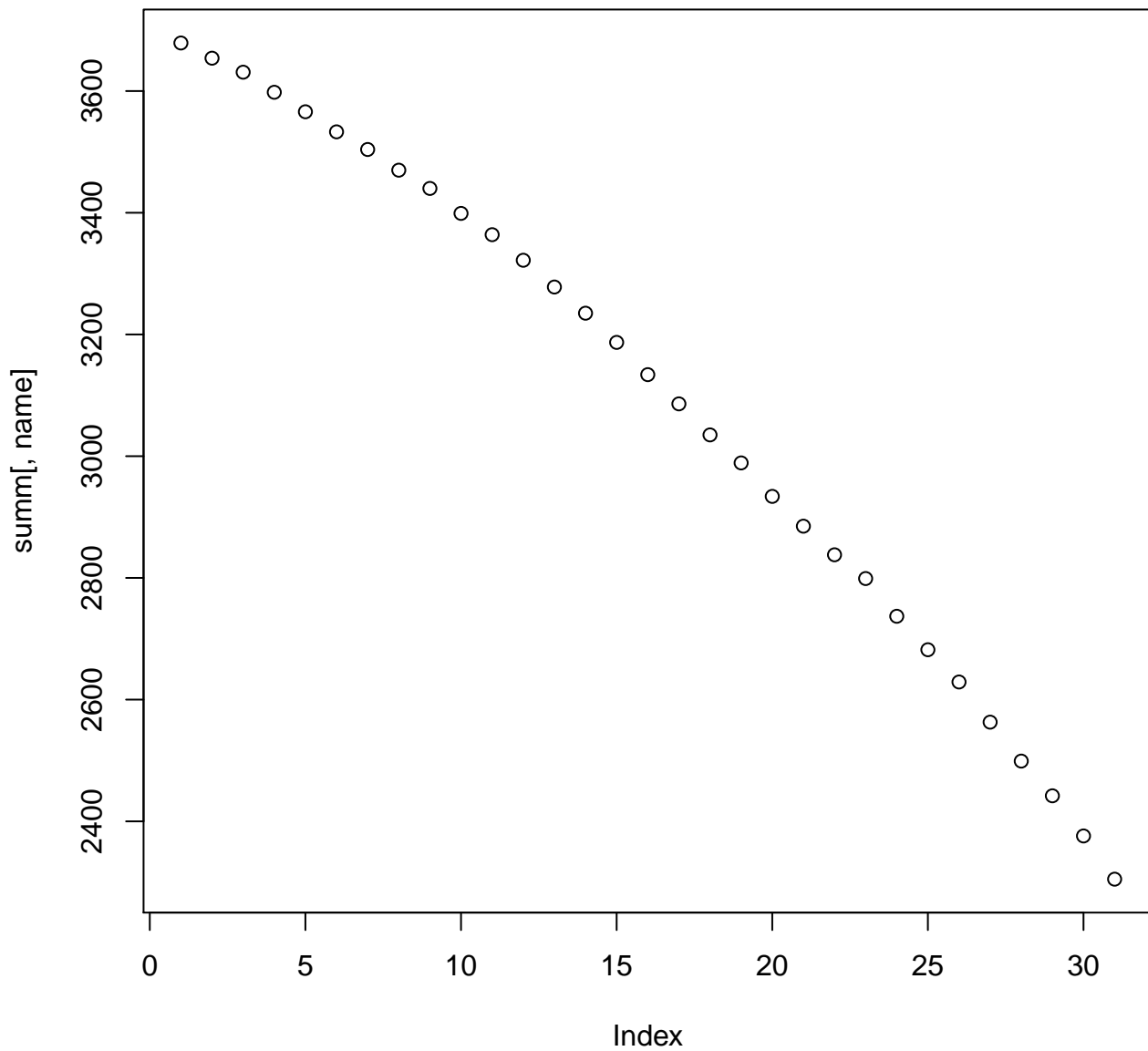
# summary



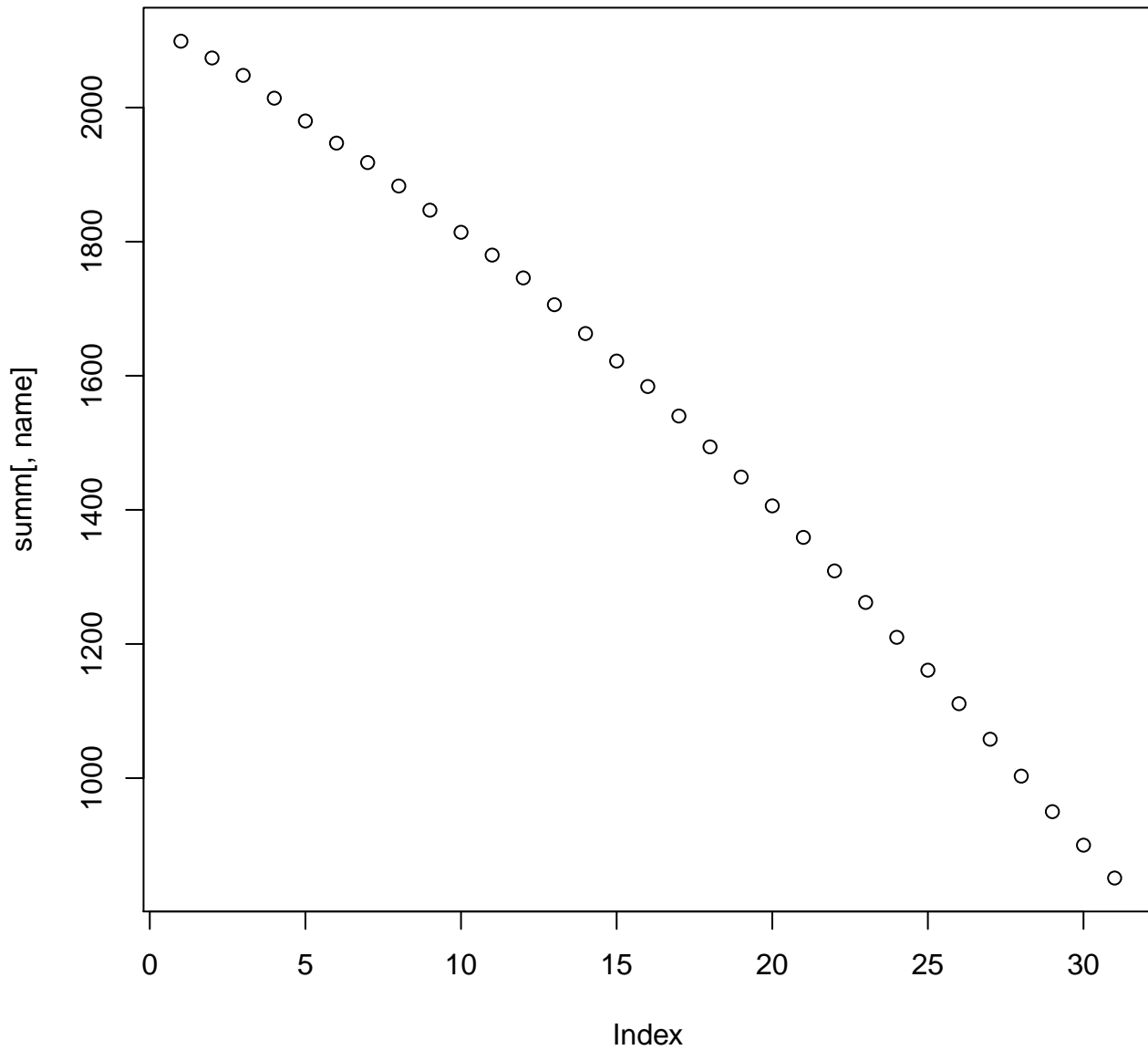
2/deweylab/nathanae/real/cc/cc\_0

2/deweylab/nathanae/real/cc/cc\_48

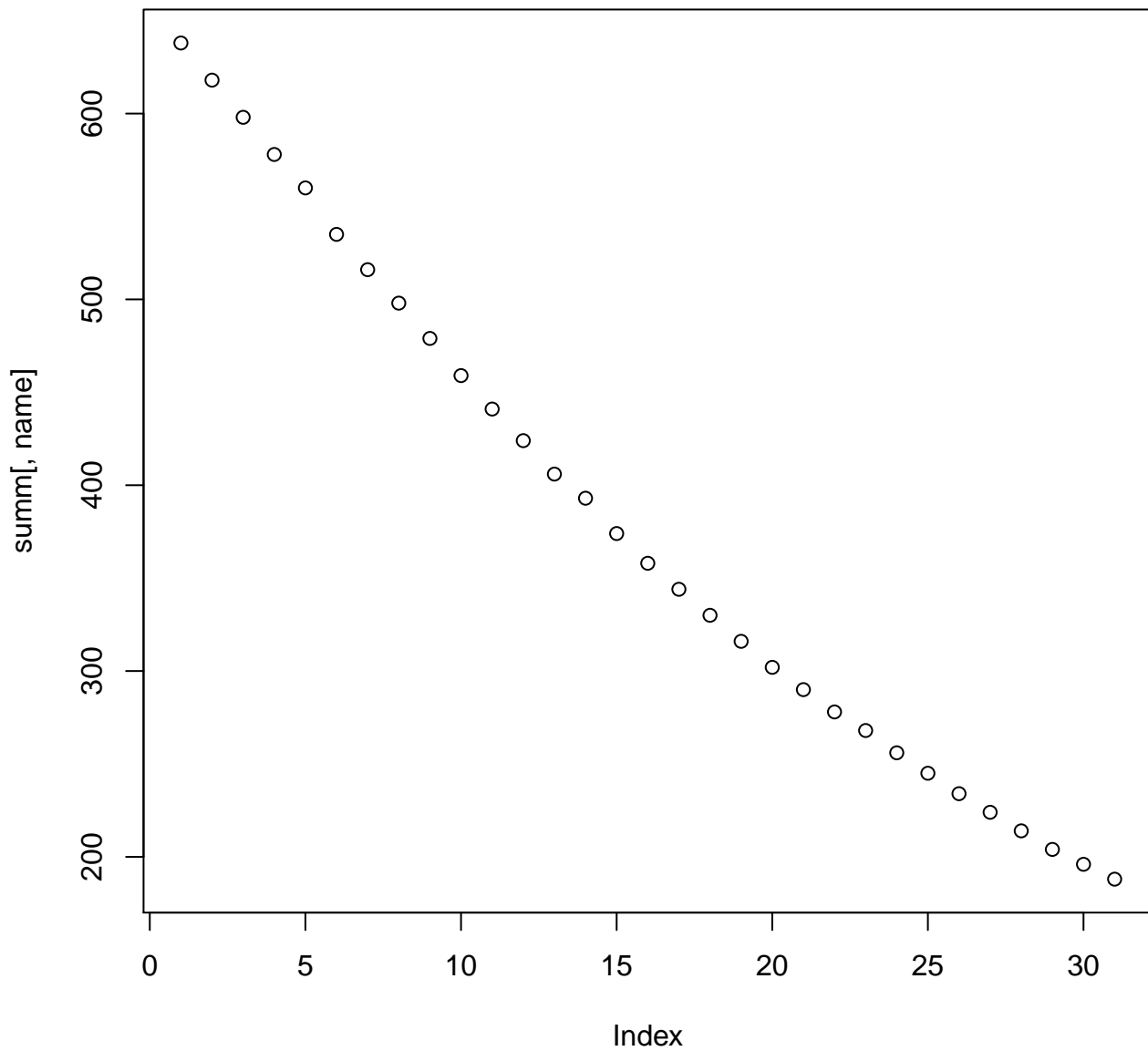
# assembly.N25



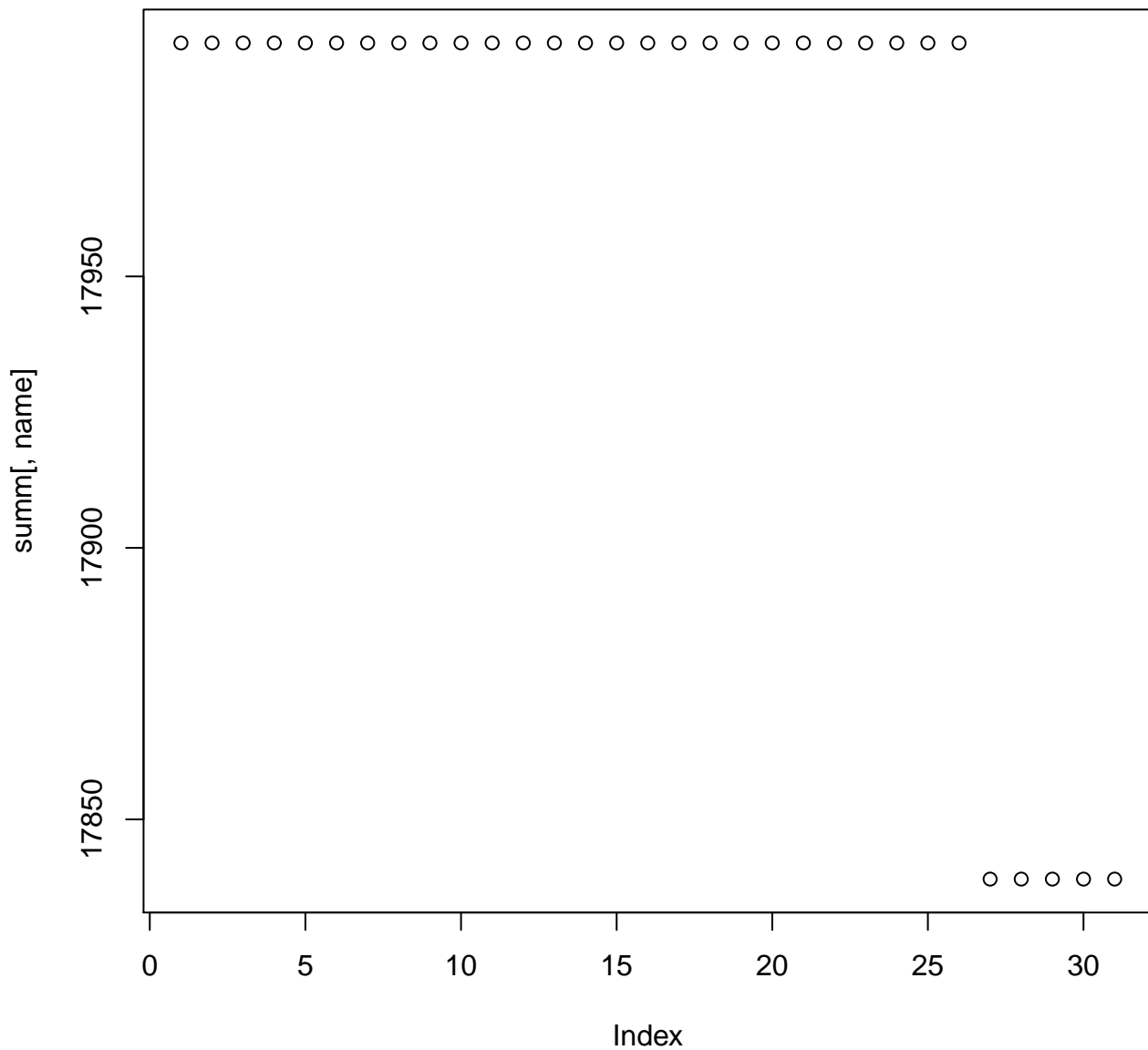
# assembly.N50



# assembly.N75

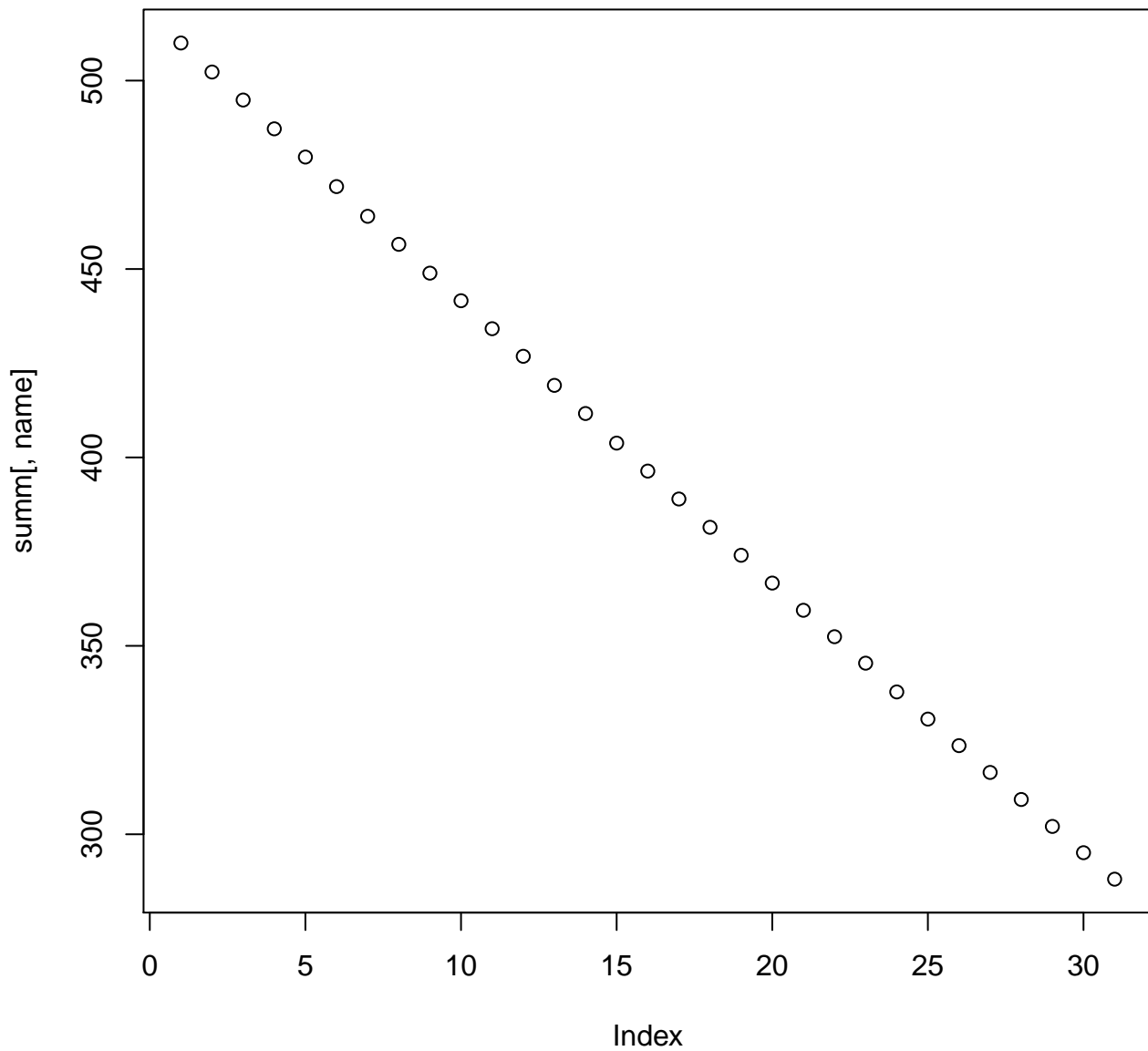


# assembly.longest

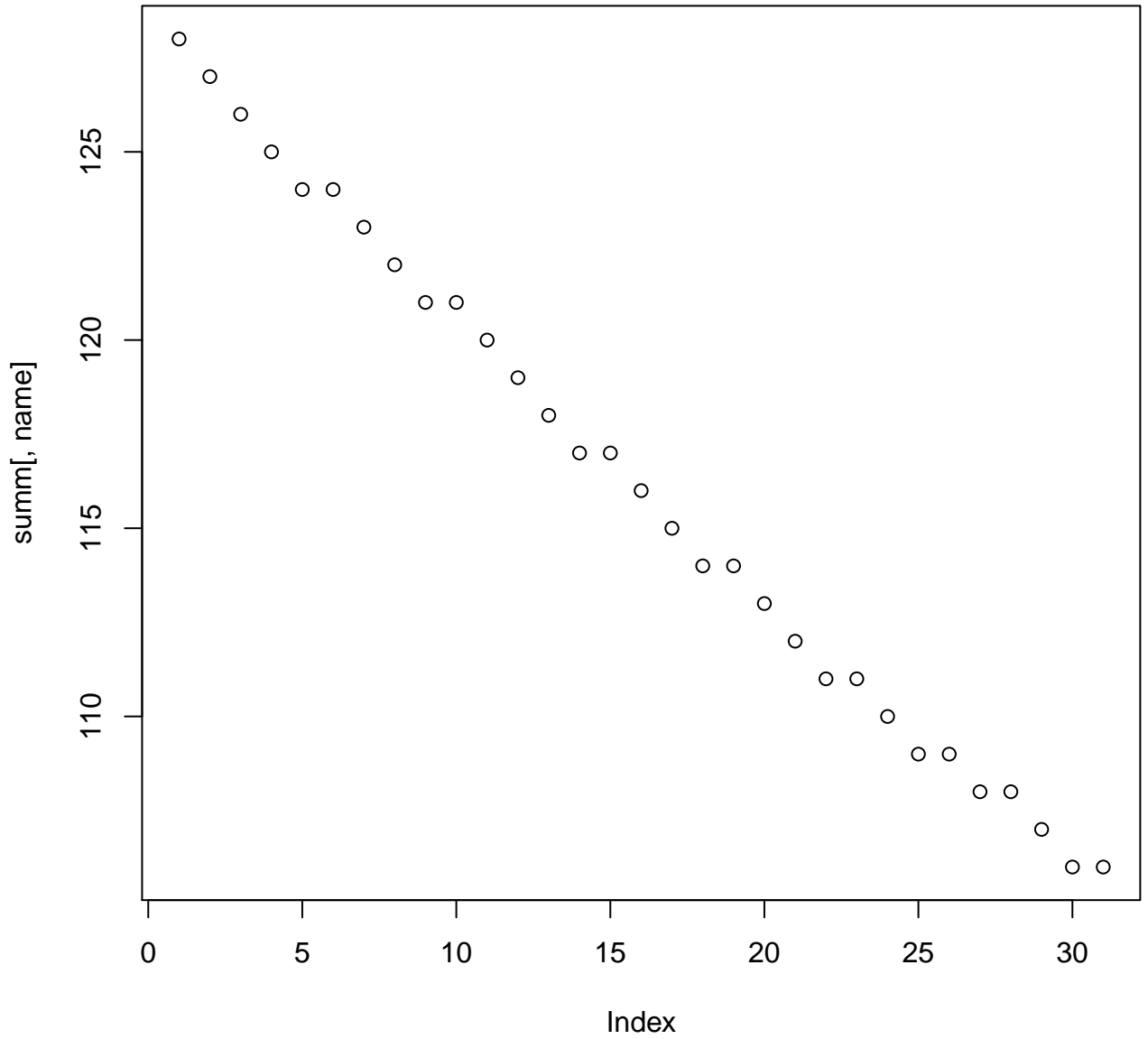




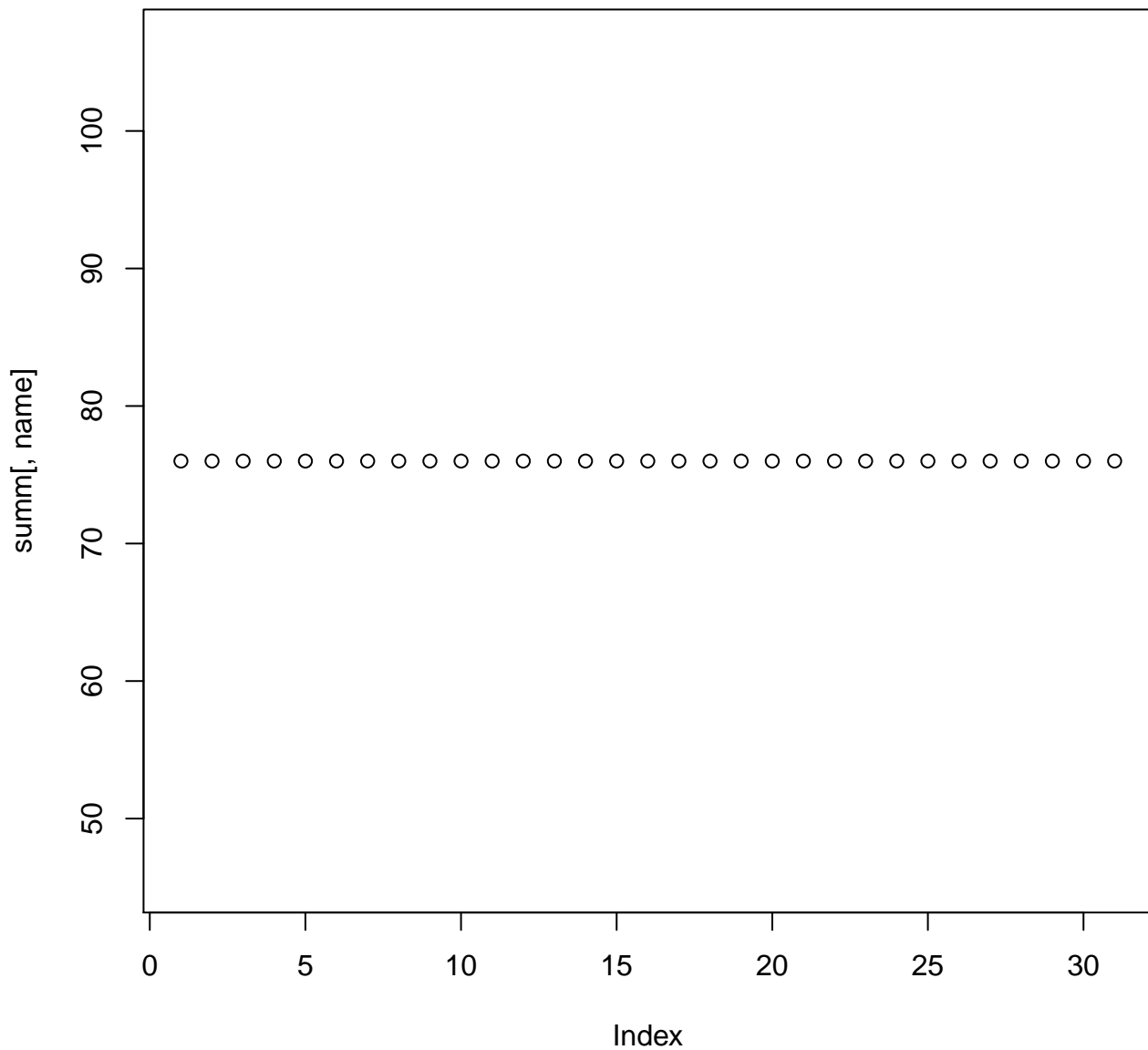
# assembly.mean



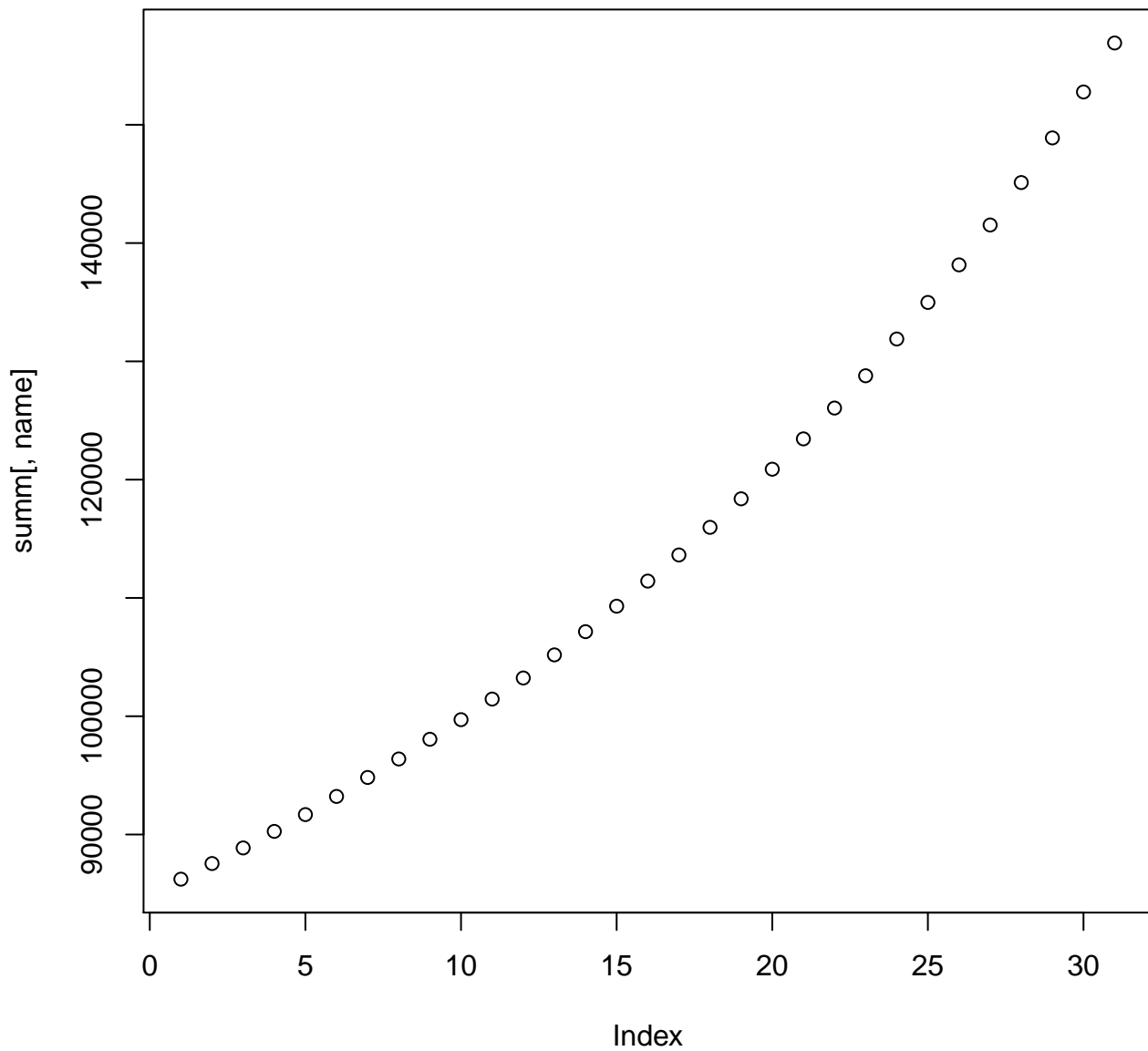
# assembly.median



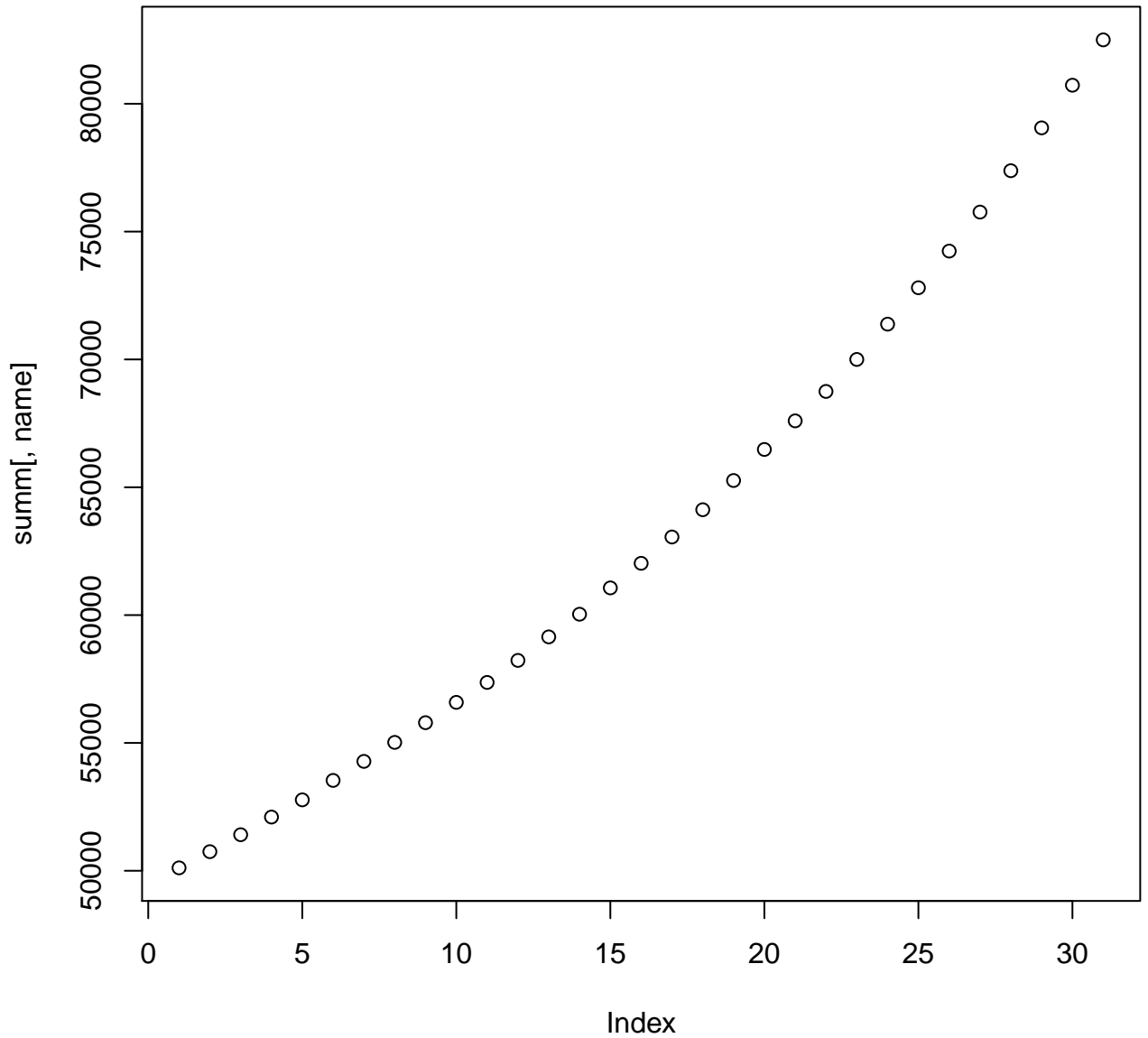
# assembly.shortest



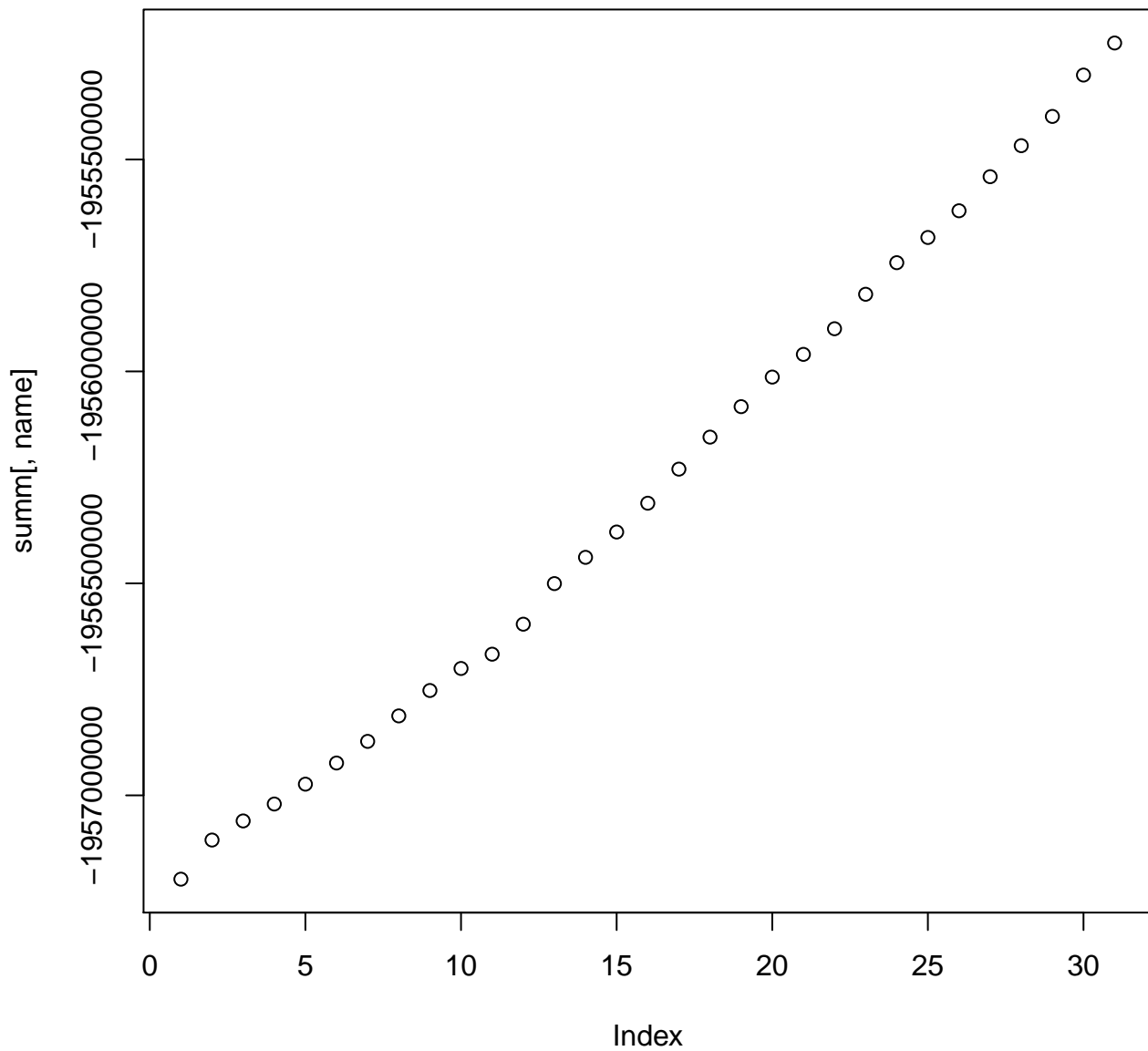
# assembly.num.contigs



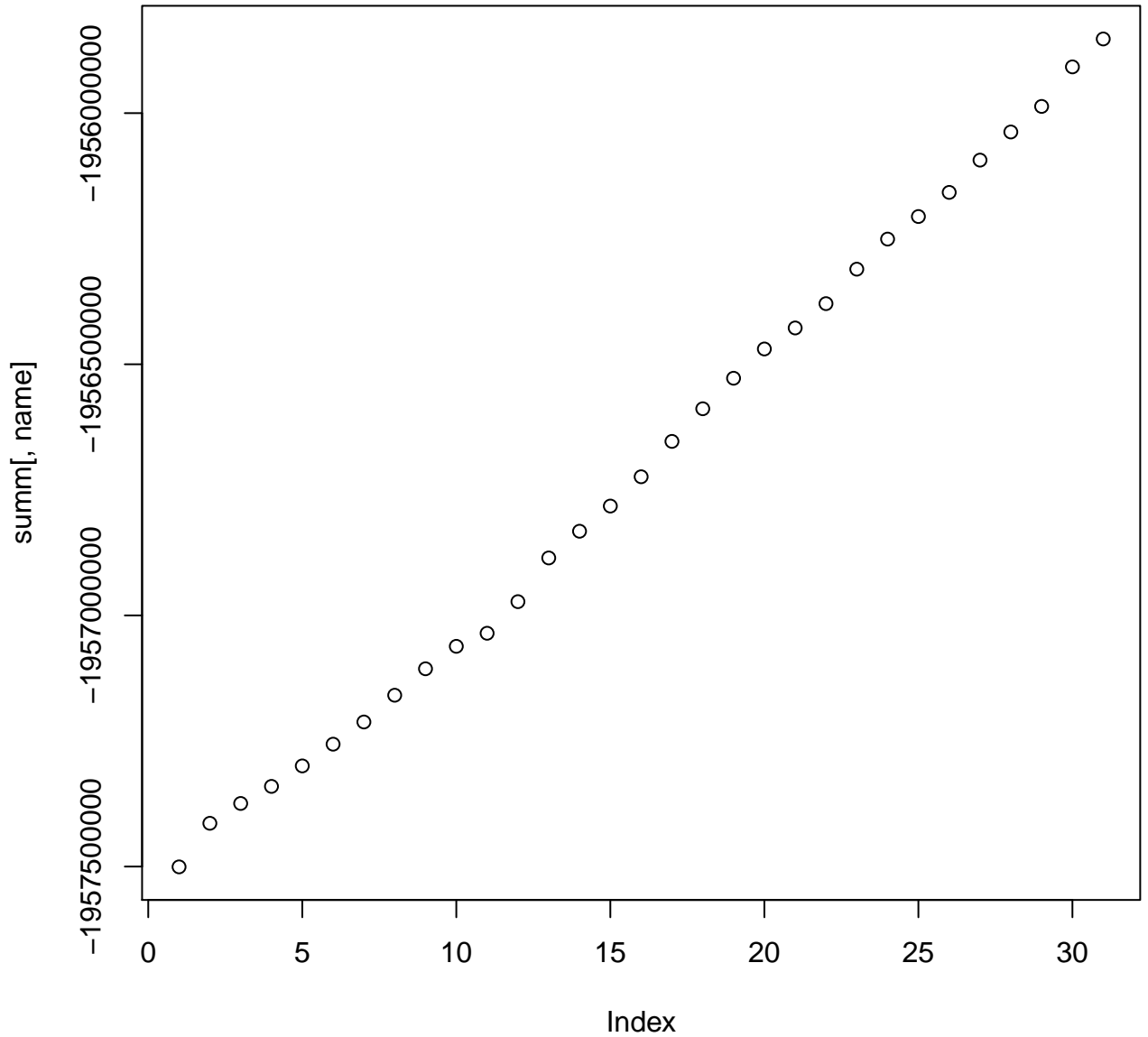
# assembly.num.contigs.longer.than.100.bp



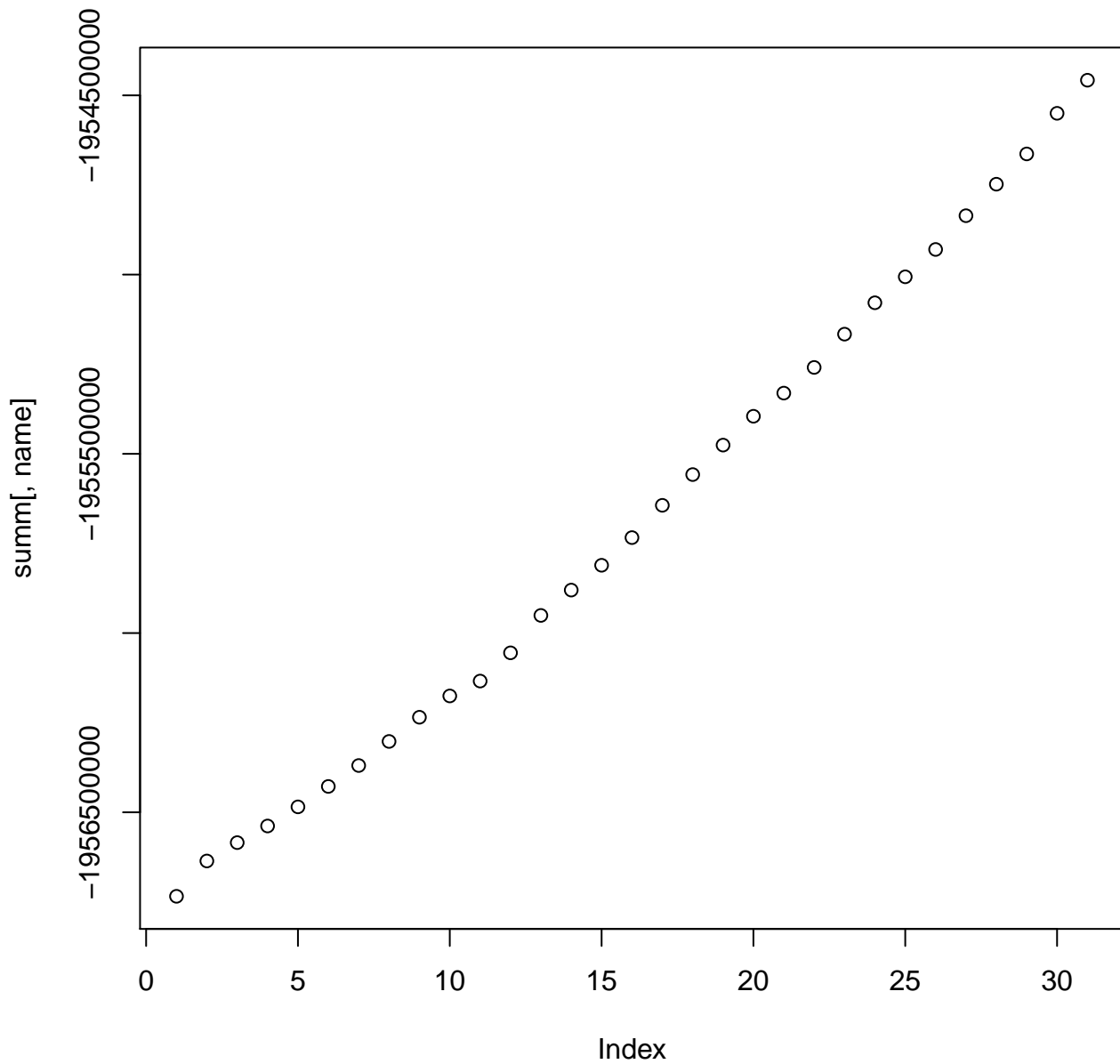
# rsem.approx.approx



# rsem.approx.bic

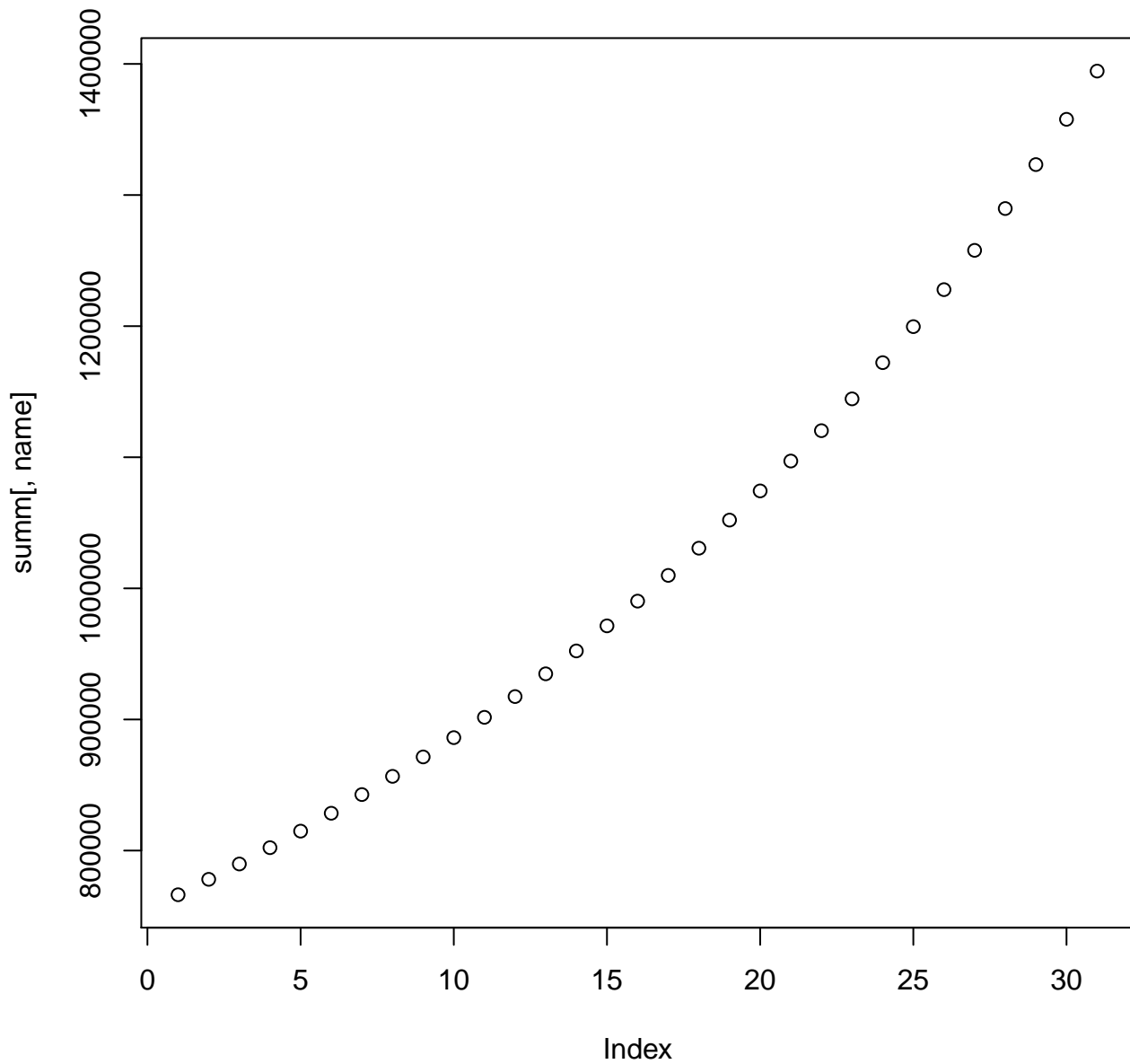


# rsem.approx.loglikelihood

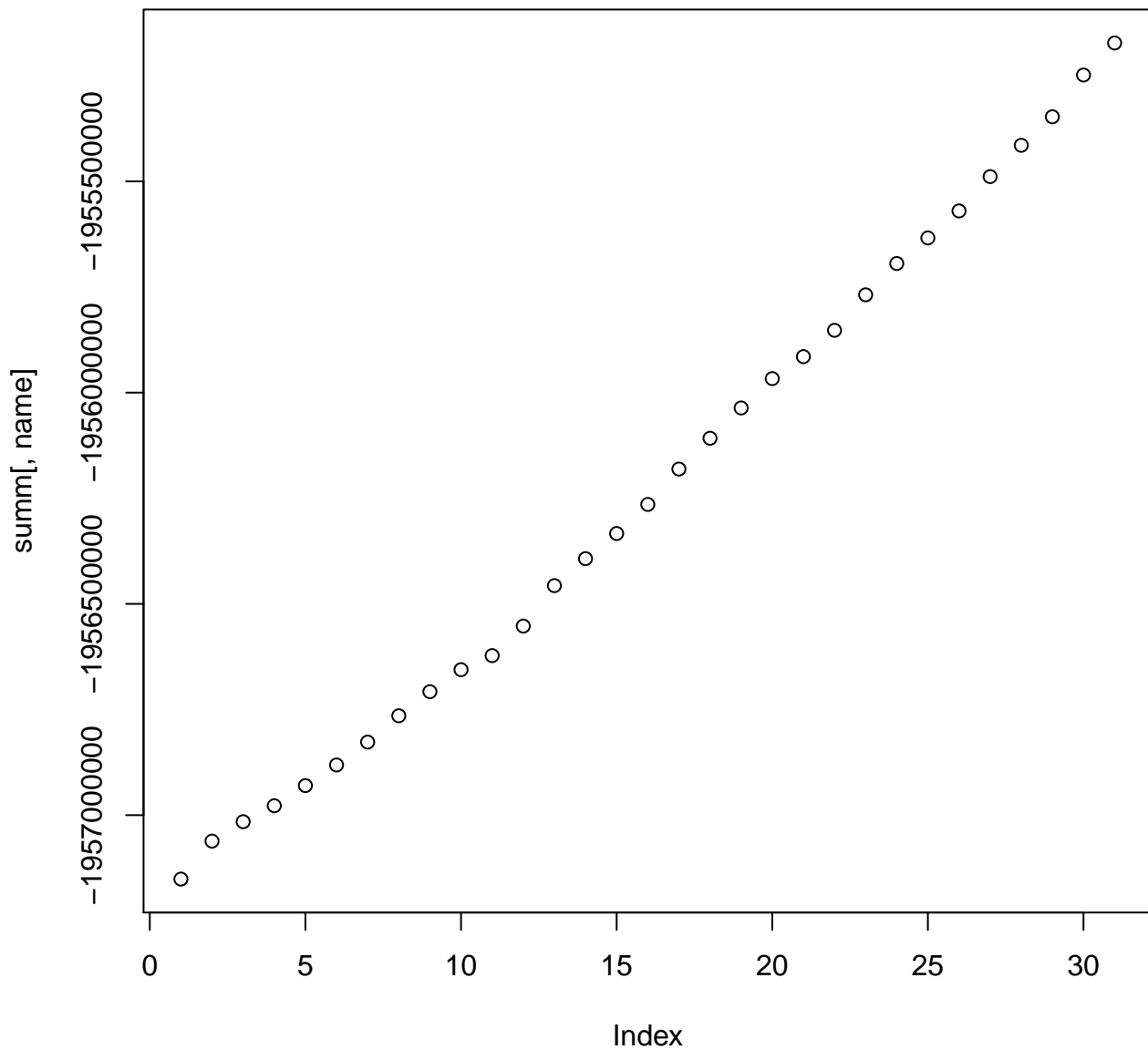




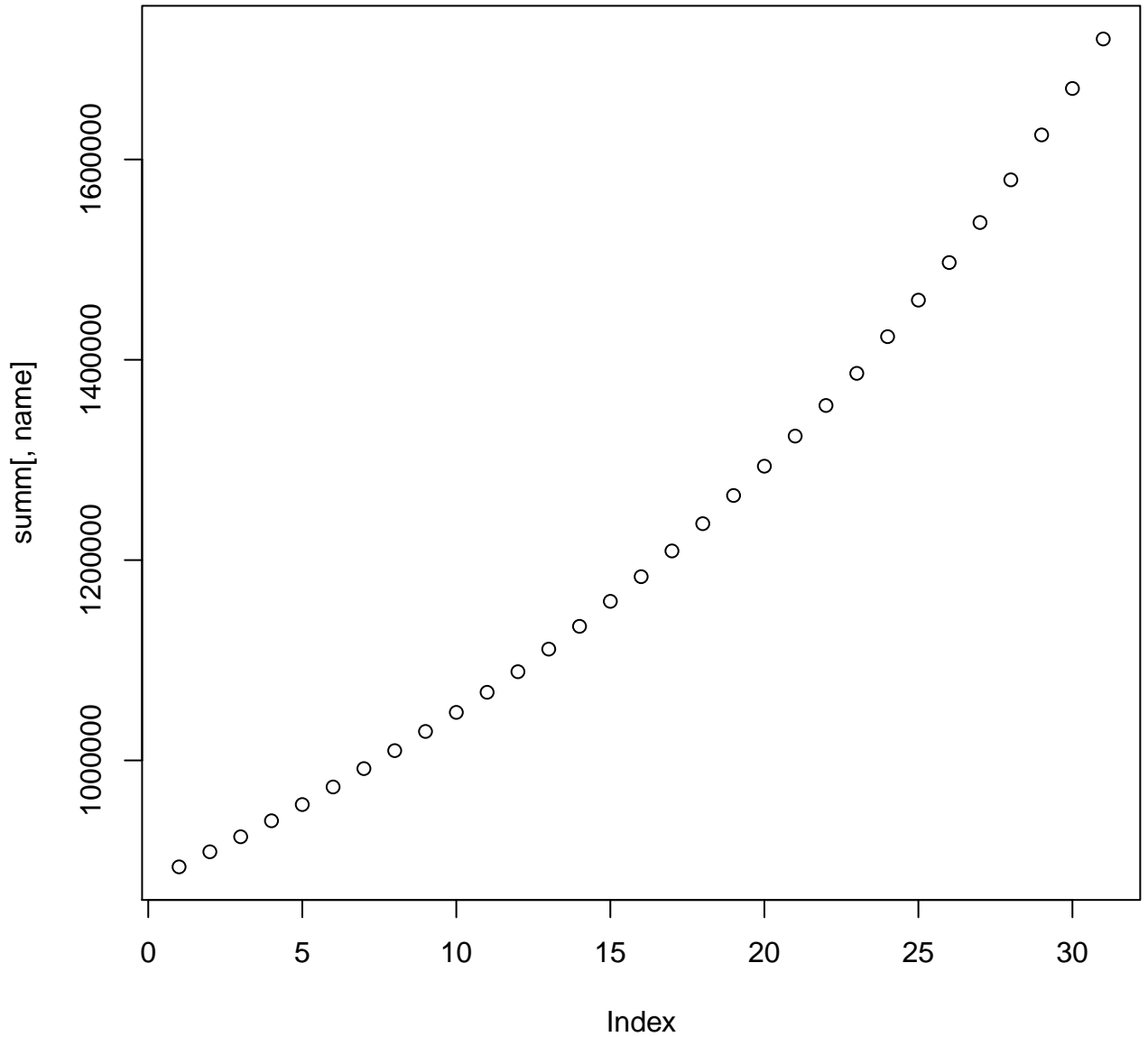
# rsem.approx.loglikelihood.penalty



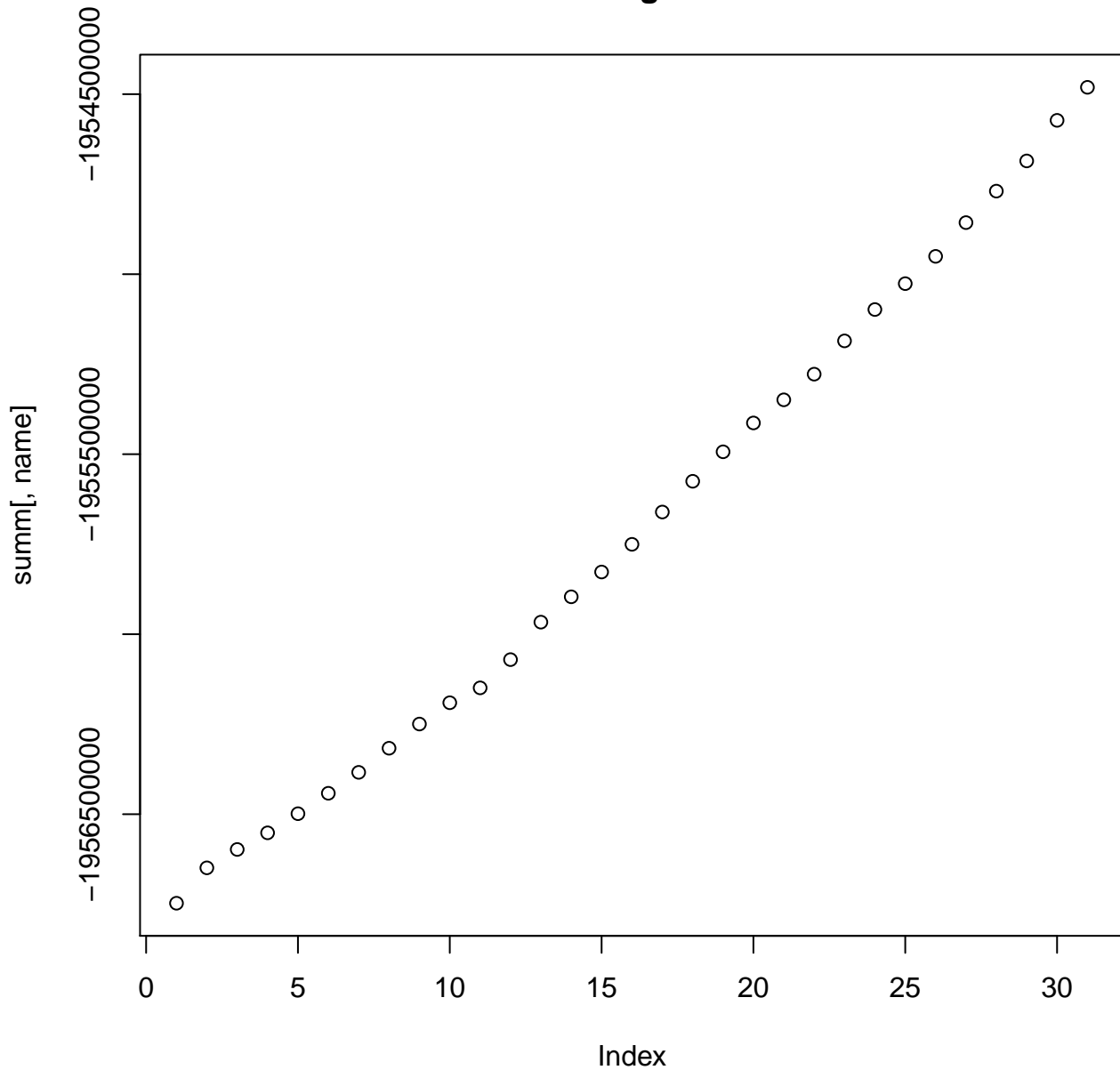
# rsem.eval.lognumer.minus.logdenom



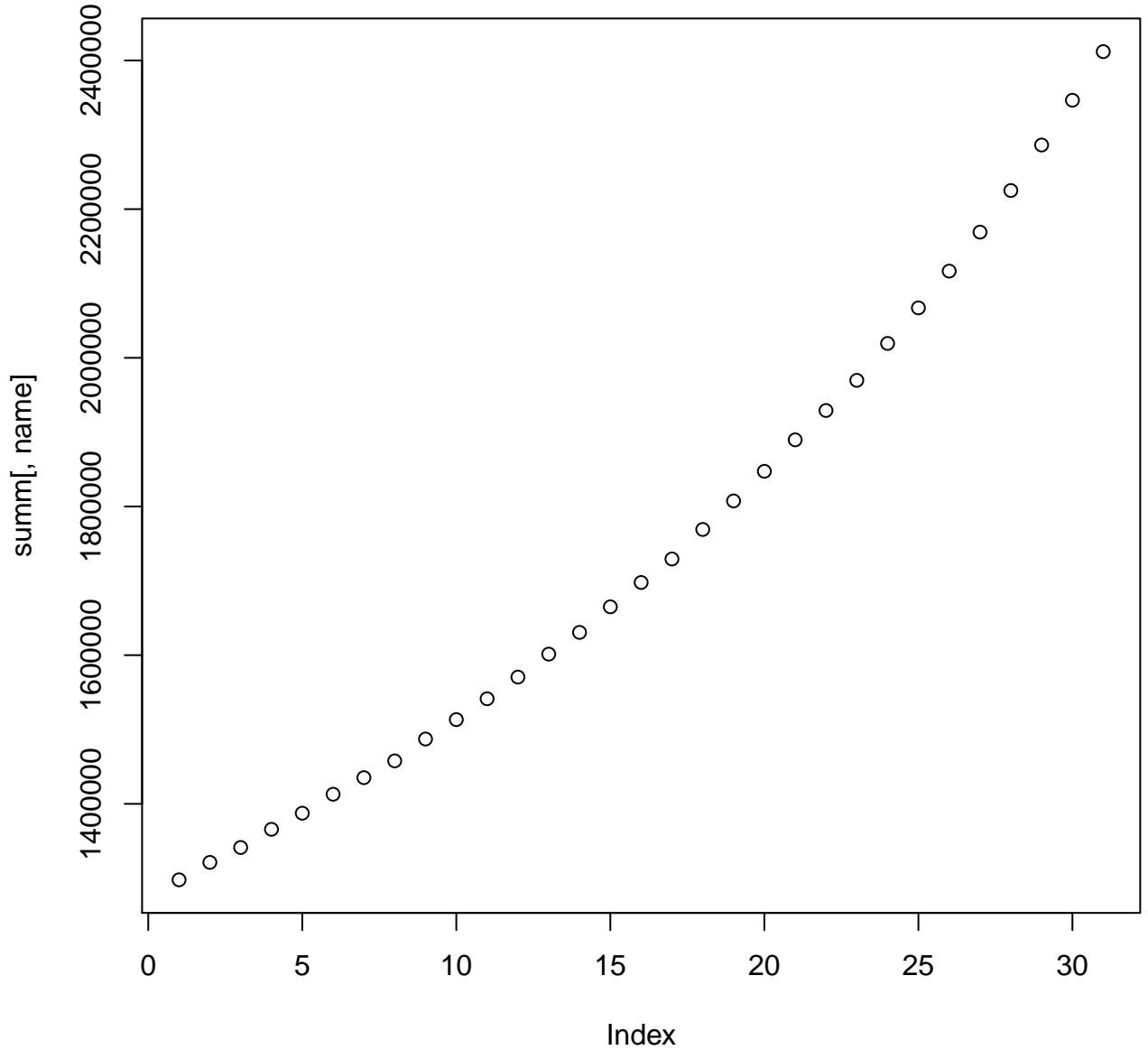
# rsem.eval.logprior



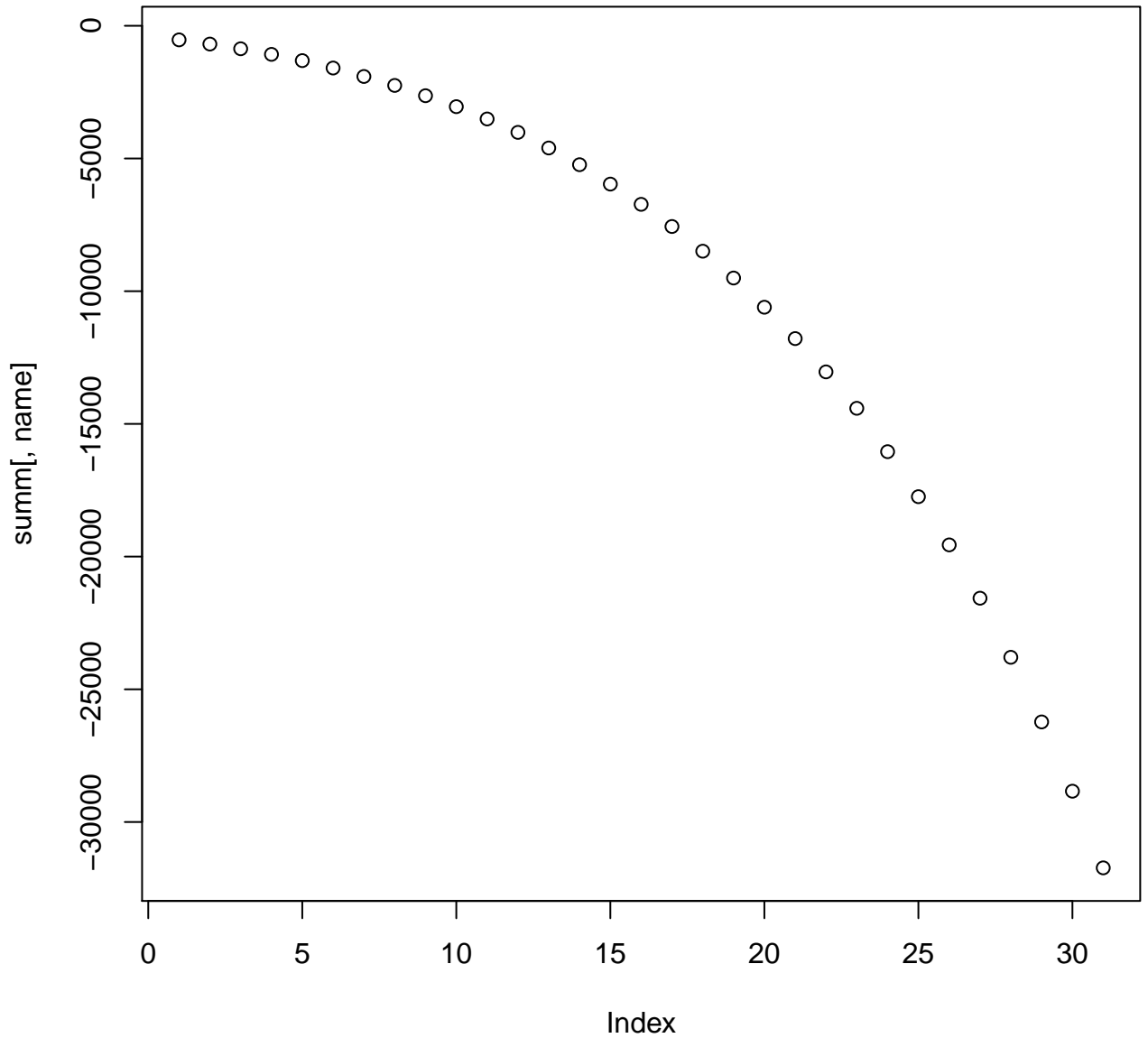
# rsem.eval.loglikelihood



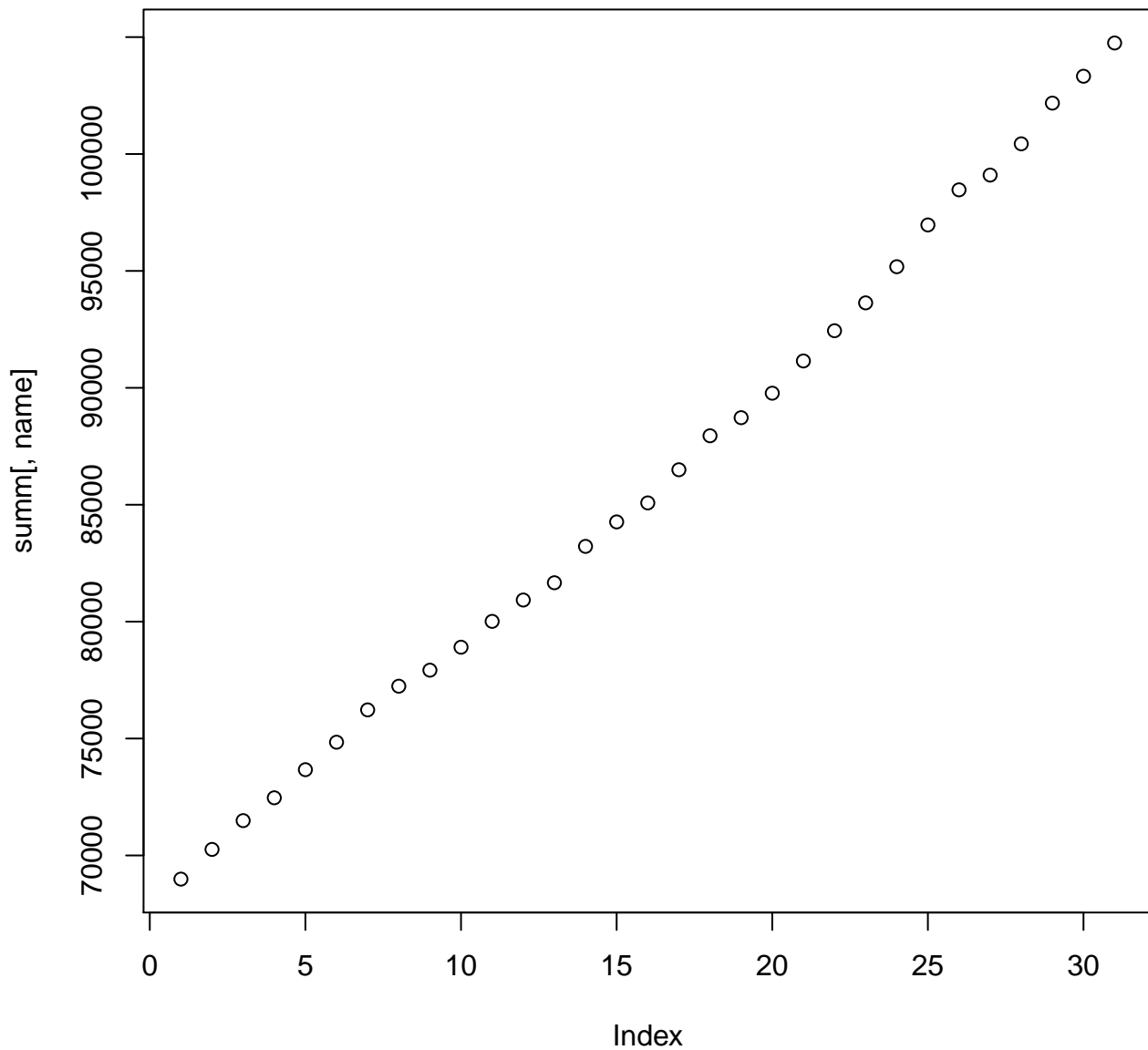
# rsem.eval.logdenom



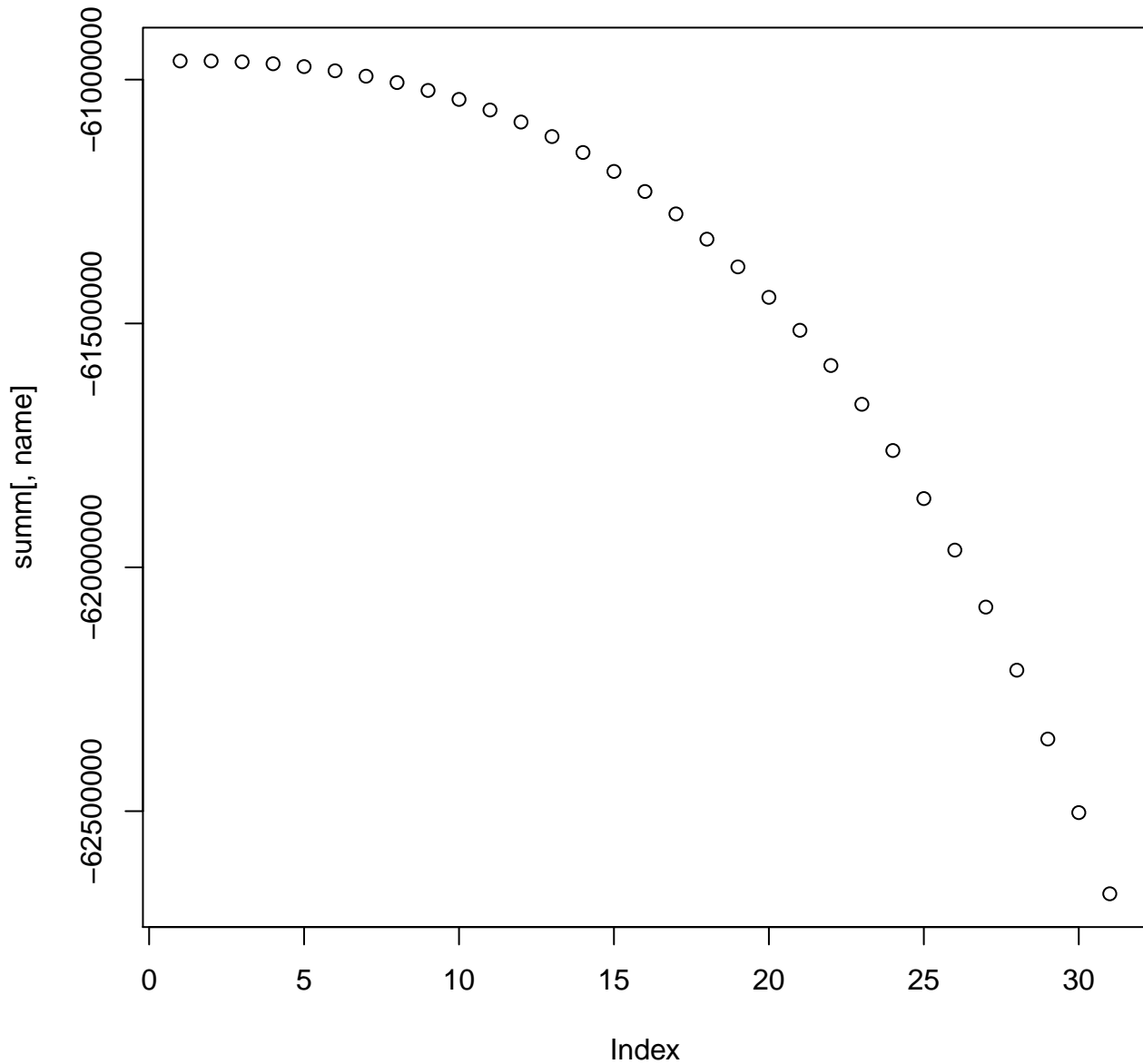
# rsem.prior.log.prob.M



# rsem.prior.log.prob.L.given.M

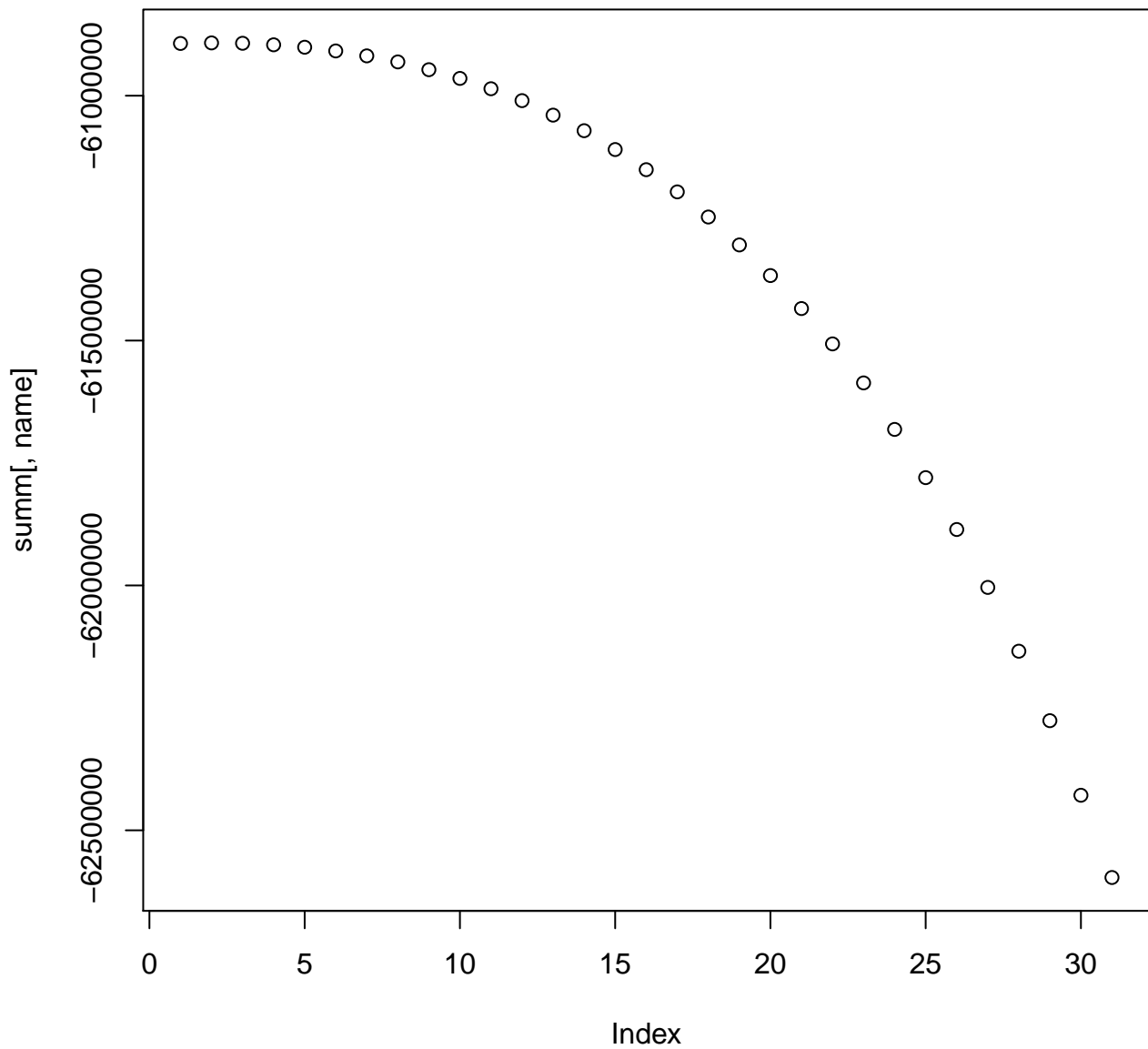


# rsem.prior.log.prob.Sequences.given.L.and.M

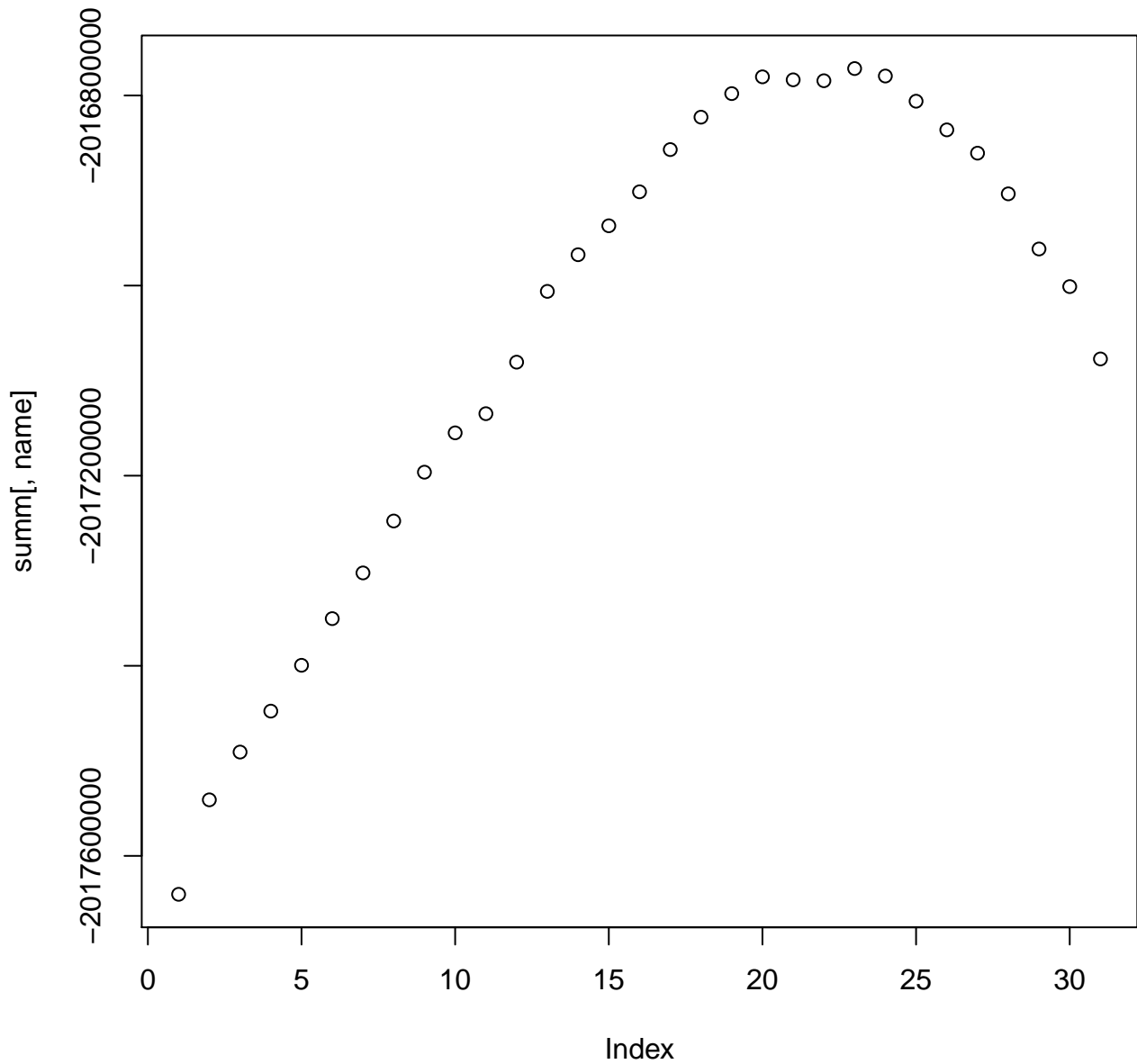




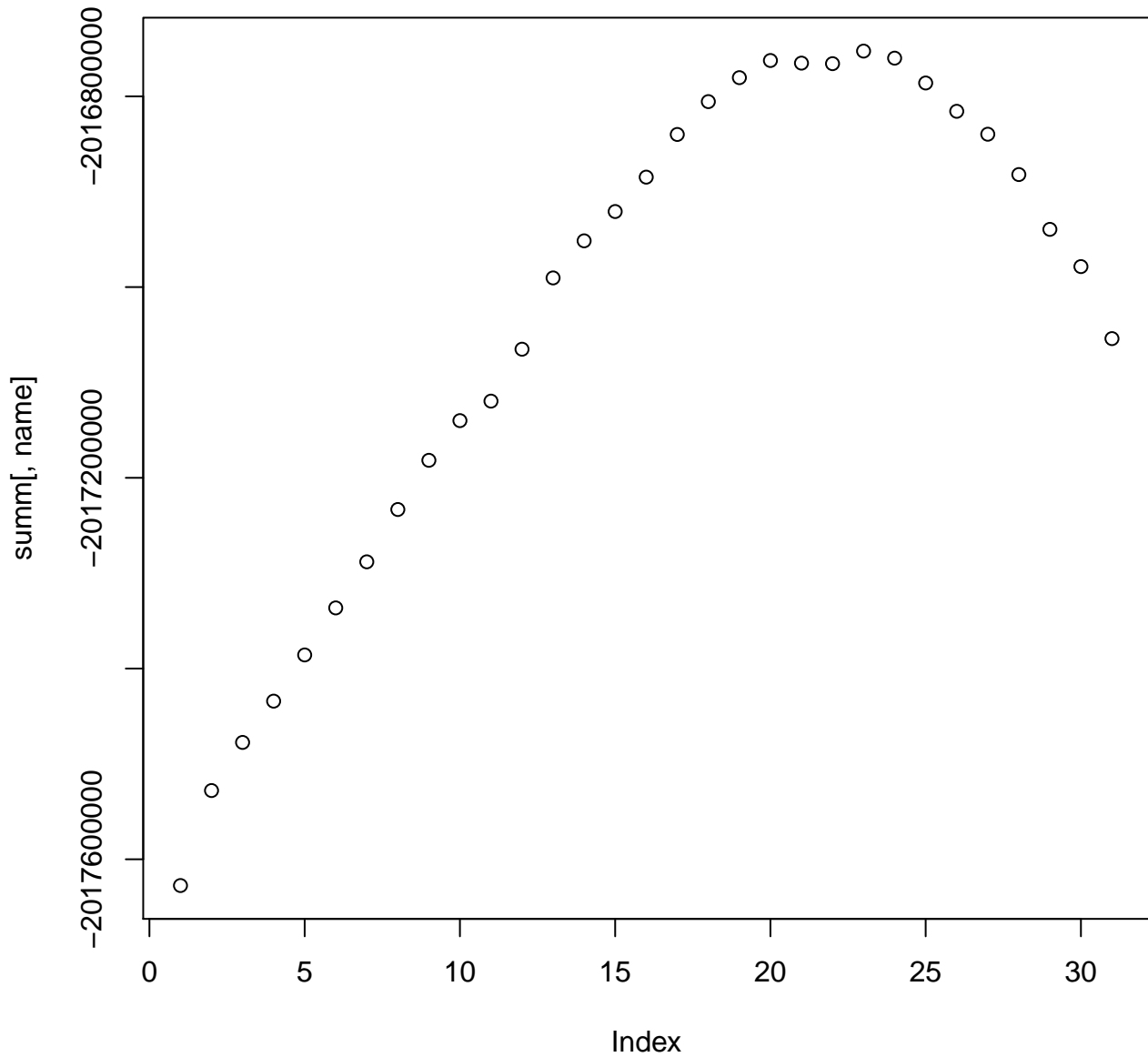
# rsem.prior.log.prob.A



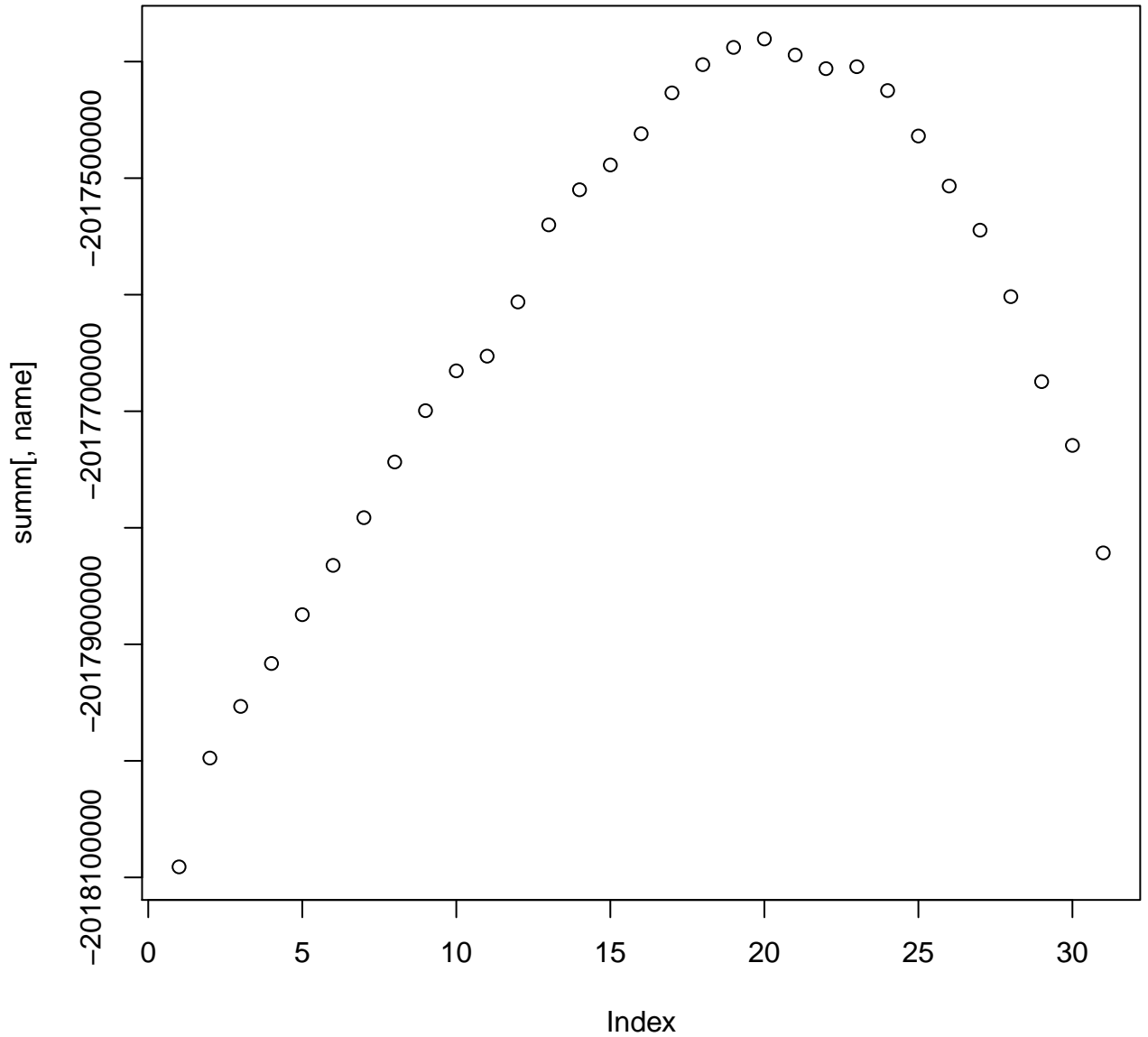
# rsem.eval.loglikelihood.plus.rsem.prior.log.prob.A



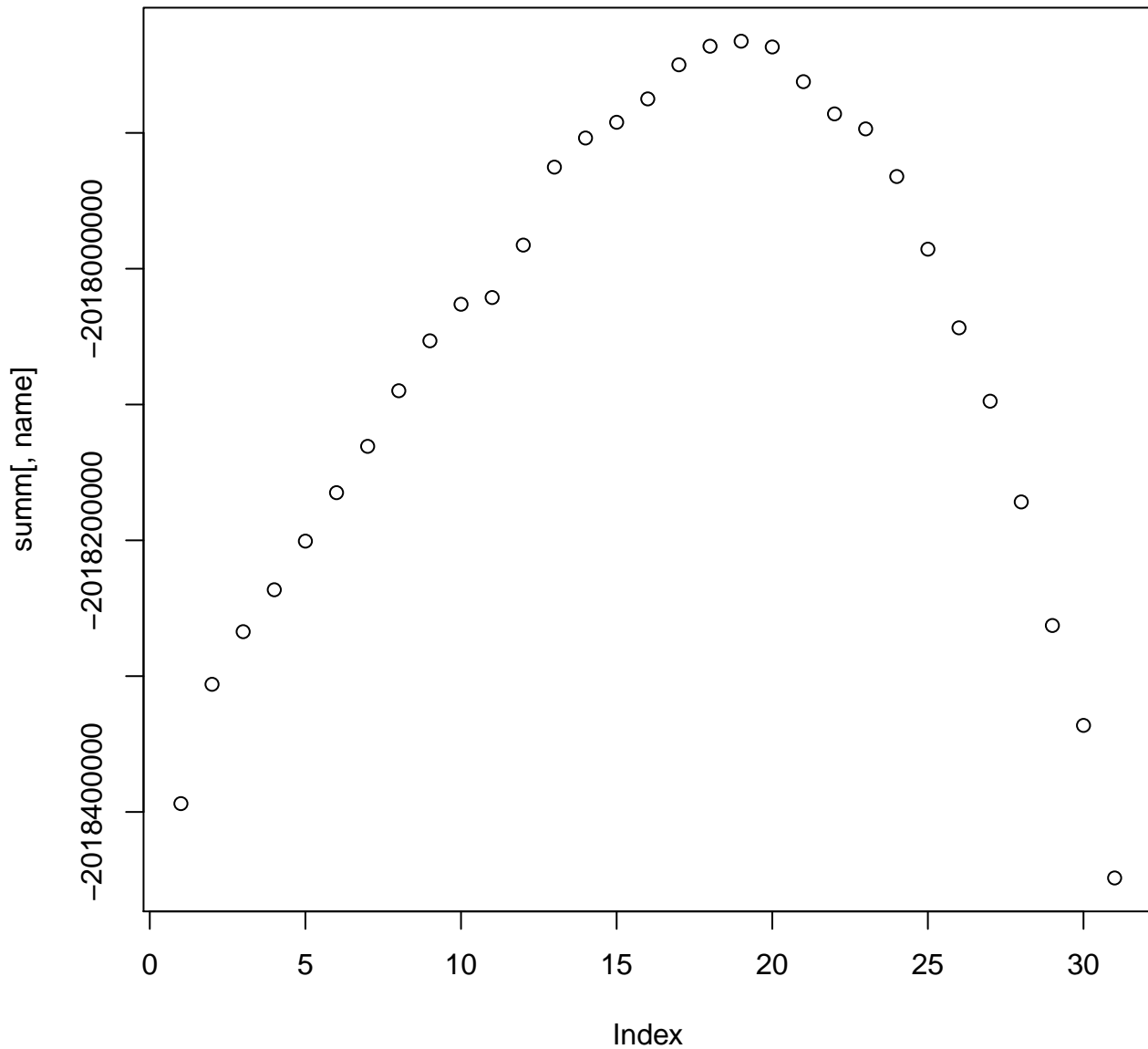
# rsem.approx.loglikelihood.plus.rsem.prior.log.prob.A



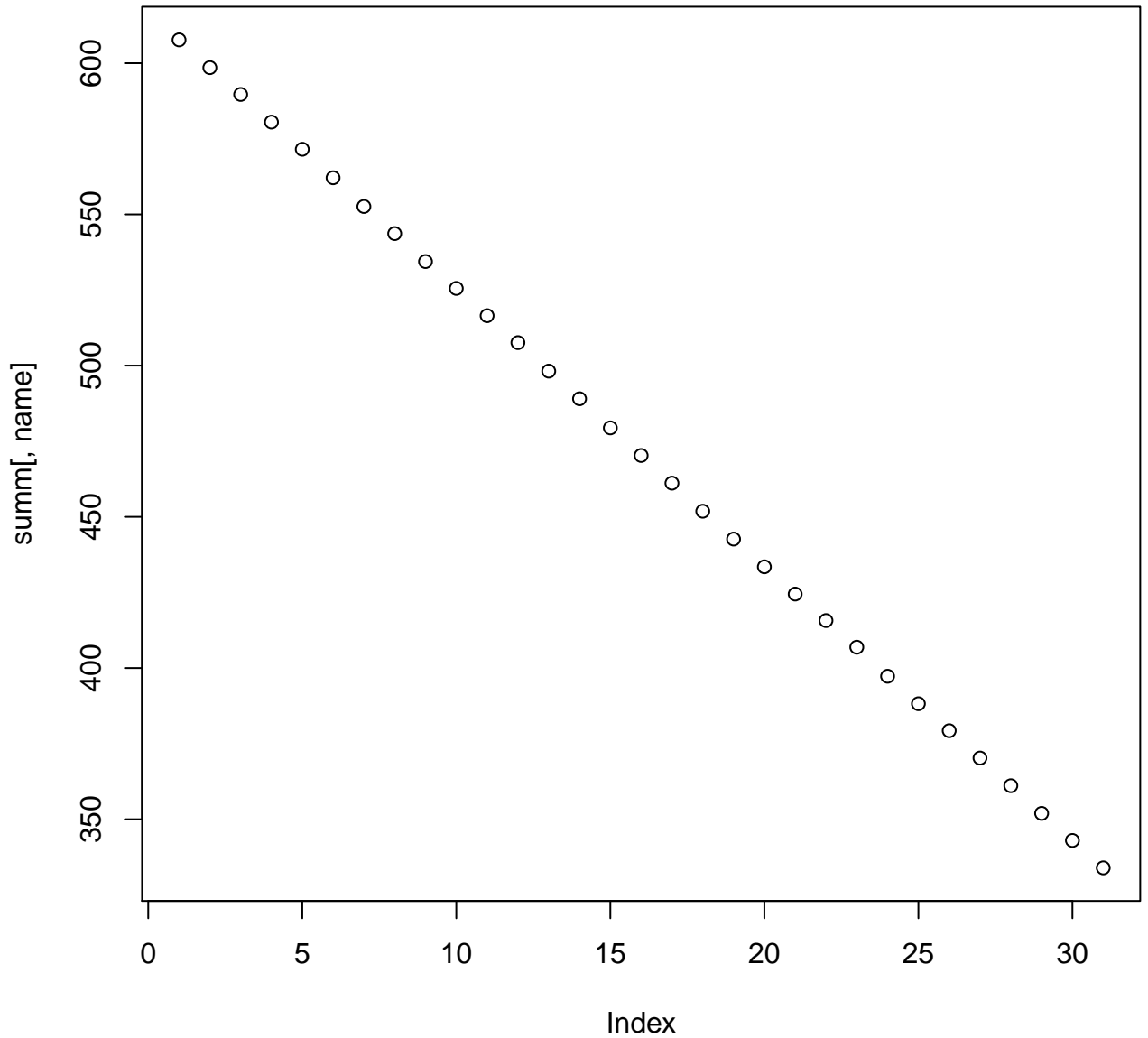
# rsem.approx.approx.plus.rsem.prior.log.prob.A



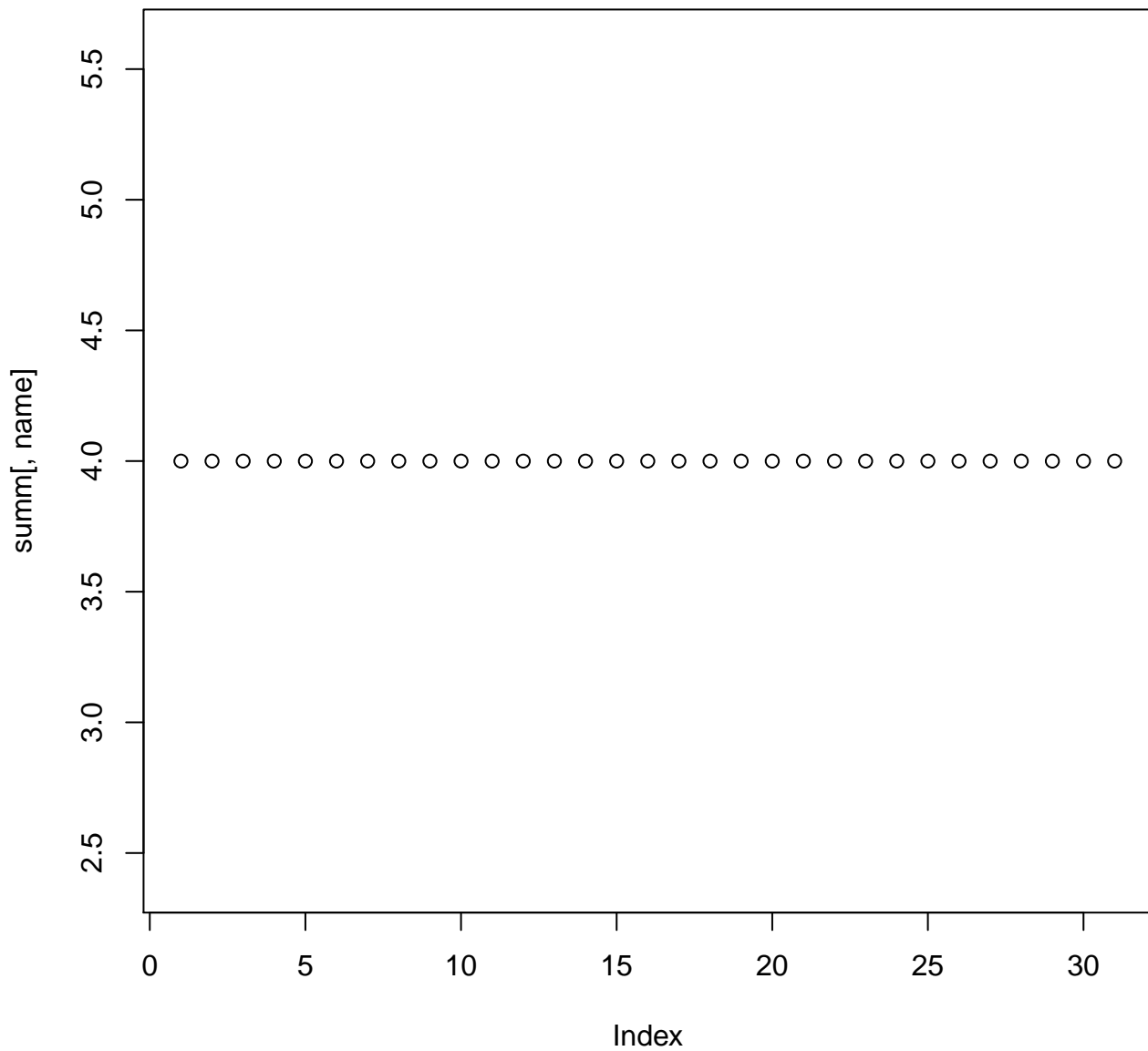
# rsem.approx.bic.plus.rsem.prior.log.prob.A



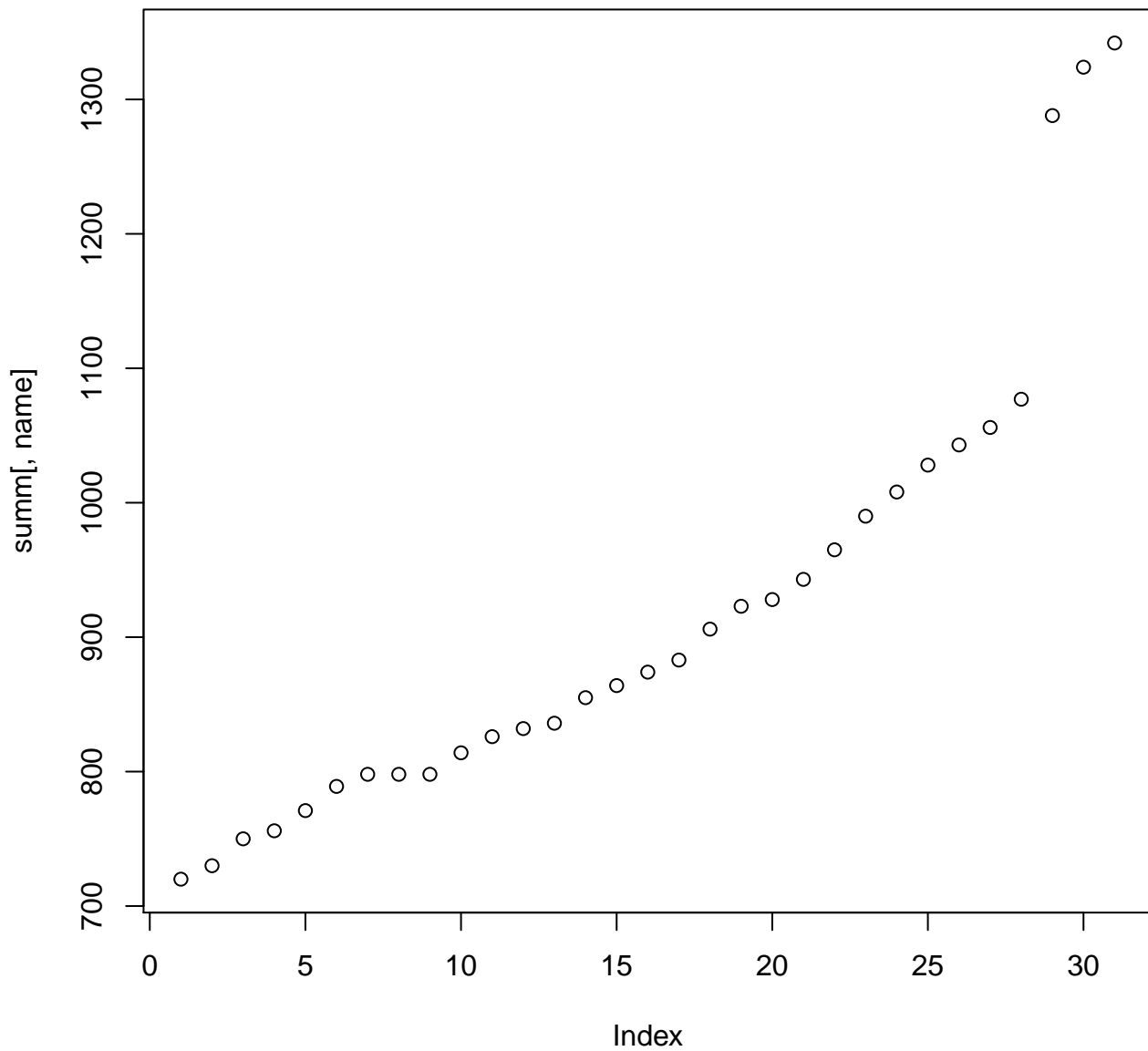
# rsem.ss.mean.num.reads.per.transcript



# rsem.ss.median.num.reads.per.transcript

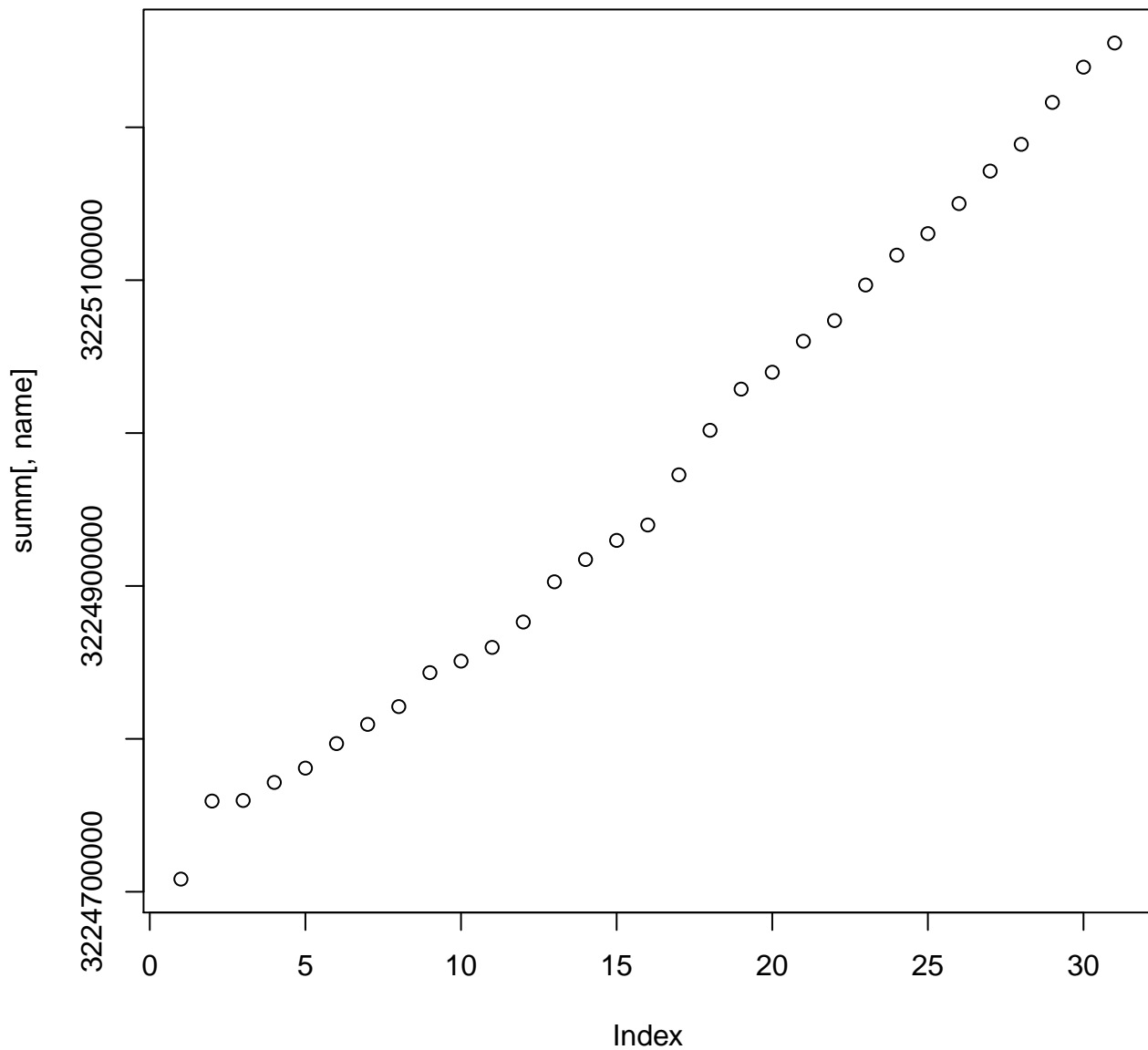


# rsem.ss.num.transcripts.with.zero.reads





# rsem.ss.num.matching.bases



# rsem.ss.num.mismatching.bases

