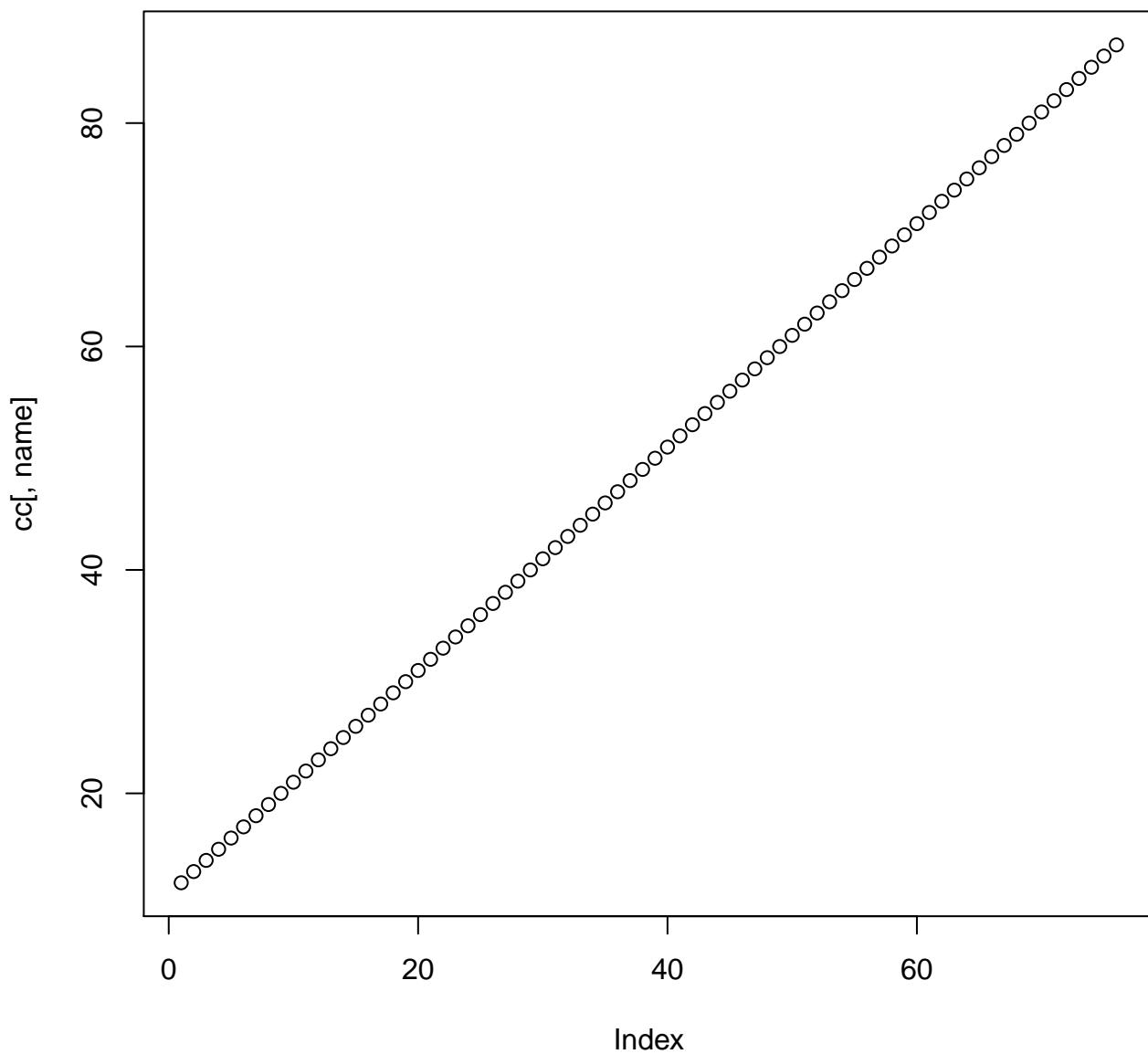
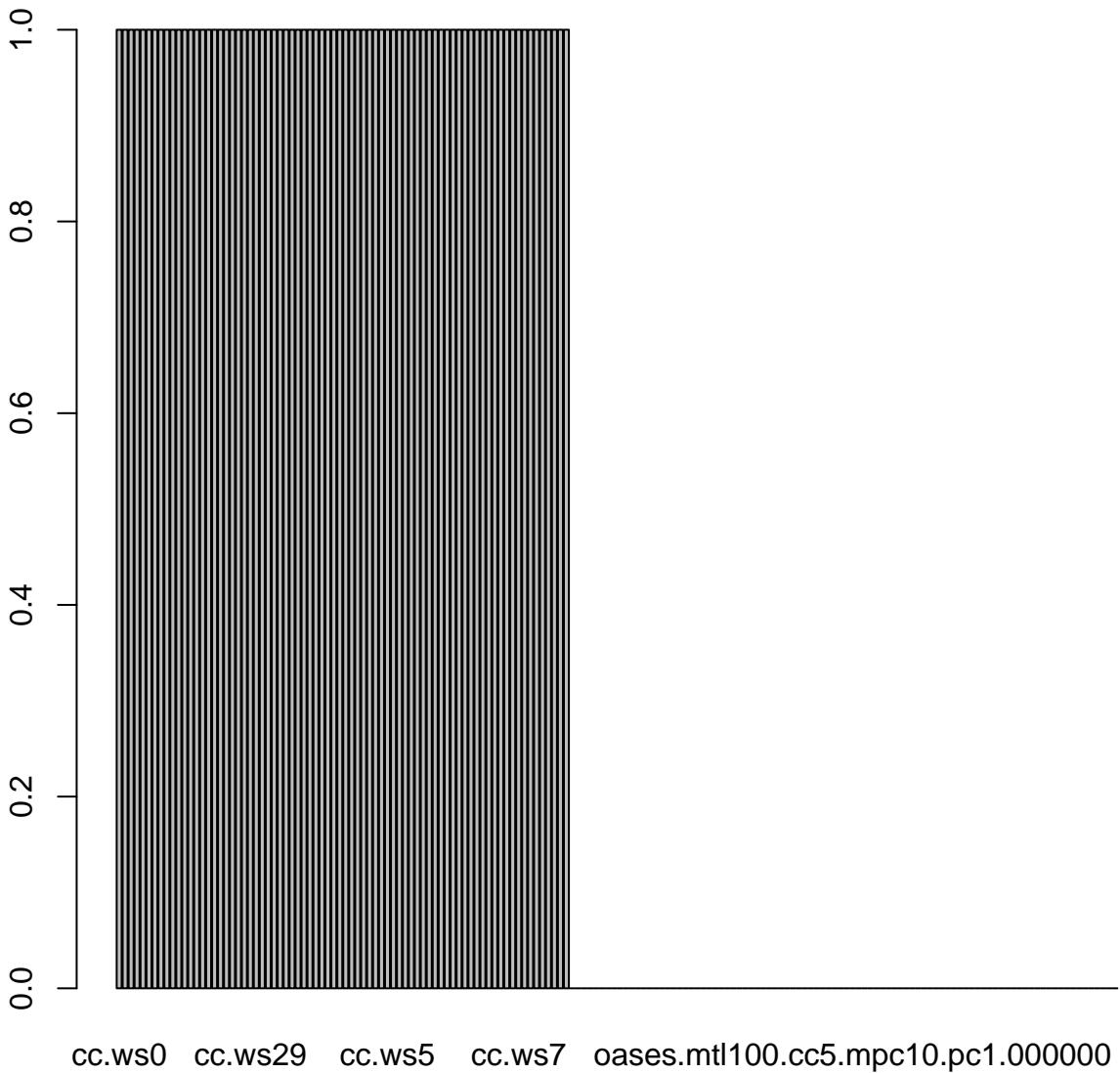


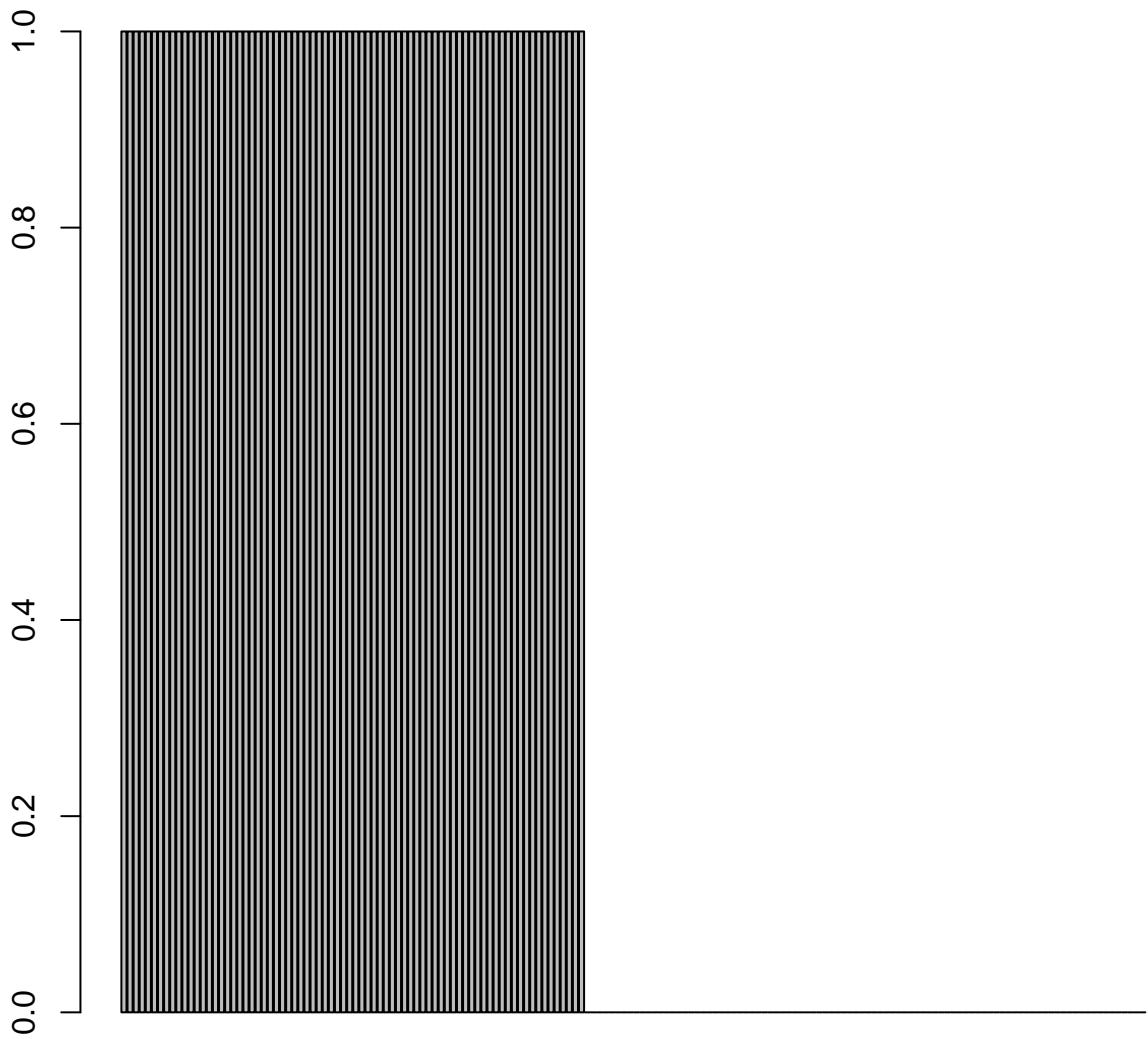
**X**



**name**

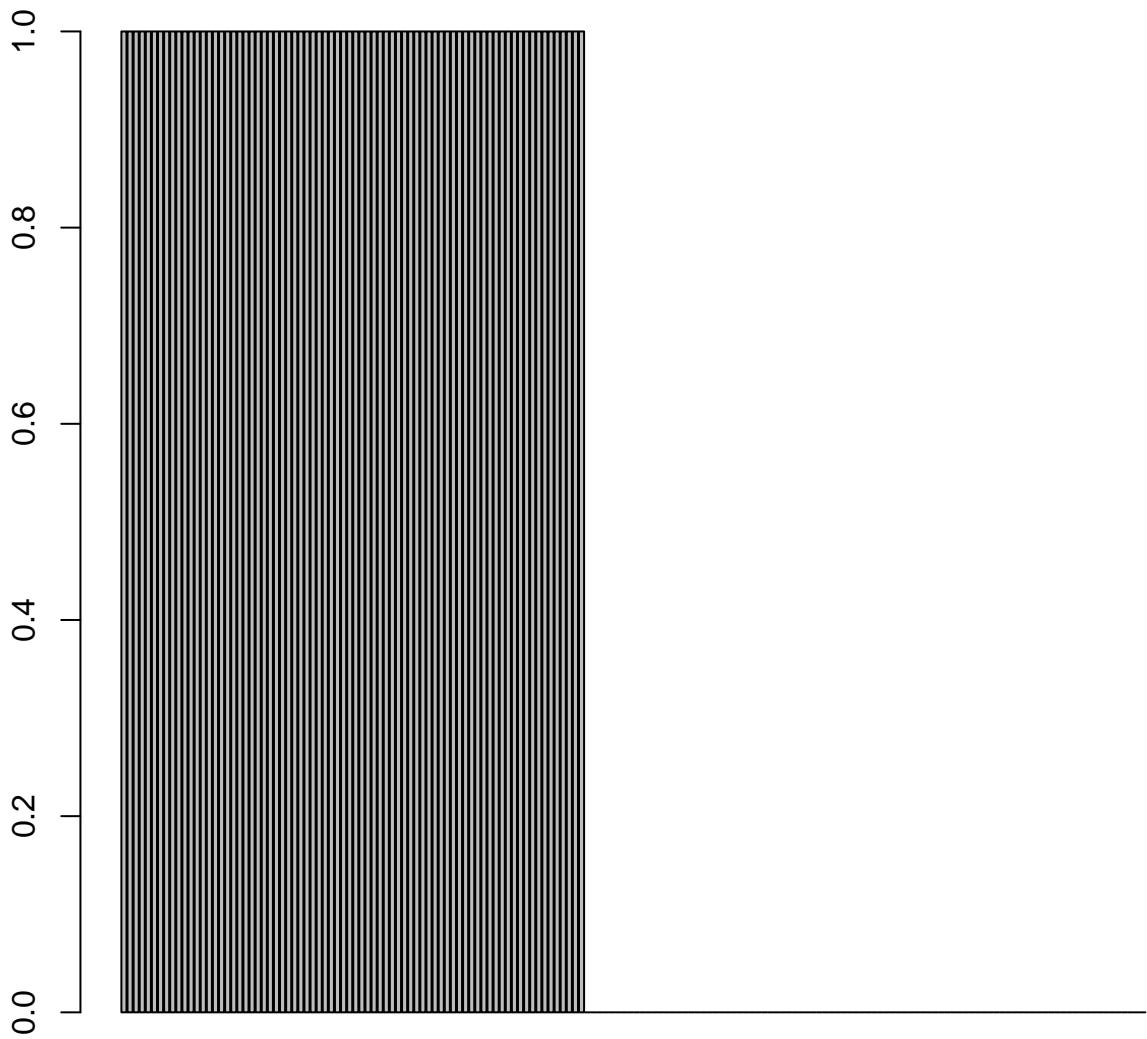


# assembly



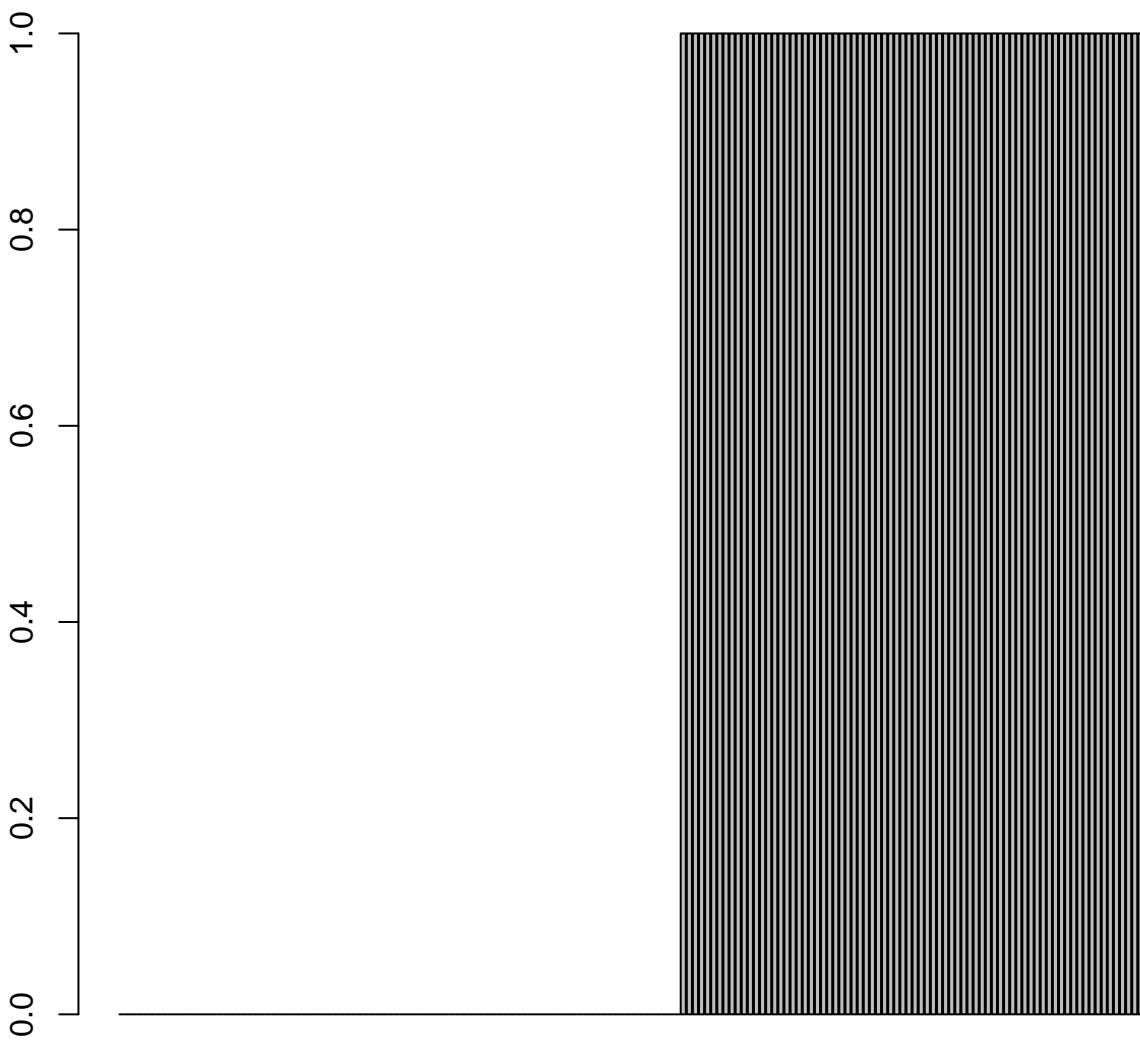
'data-oct2.single-full/gridsearch-collectContigs/rsem\_oracleset.ws0.fa

# summary

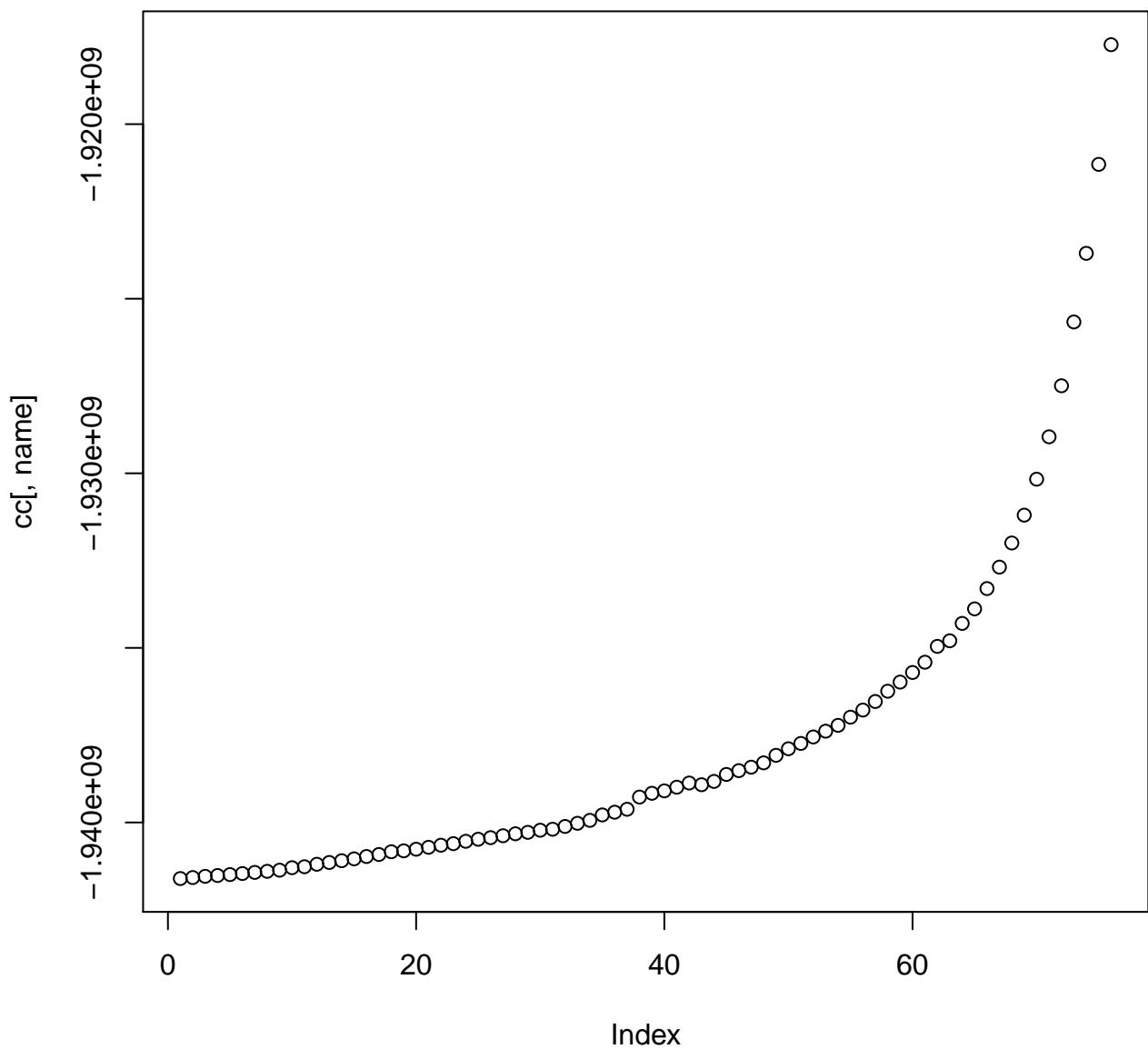


2.single–full//gridsearch–collectContigs/rsem\_oracleset–summary.ws0/summary

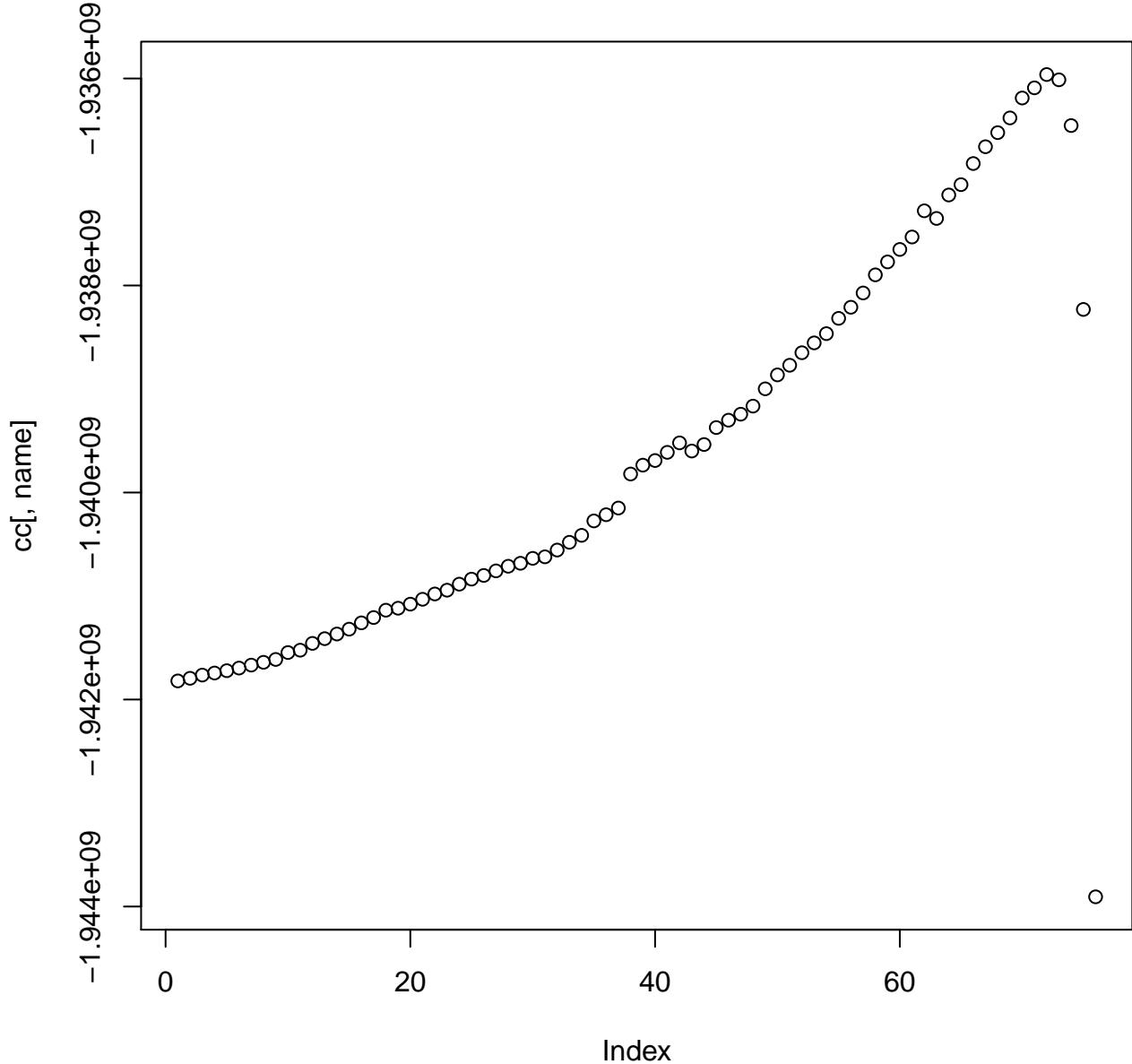
## **ssembly\_cmd**



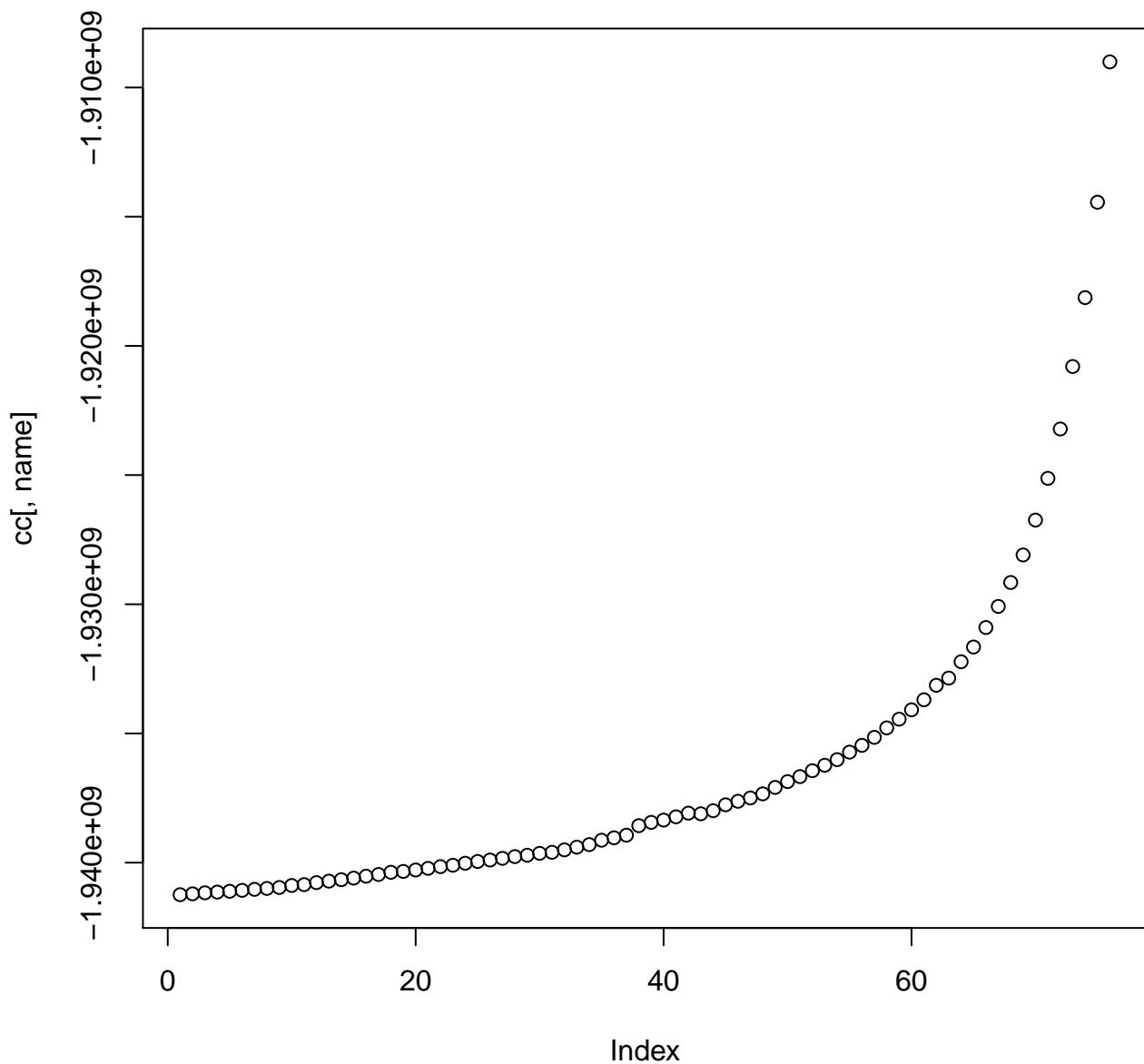
### rsem.approx.approx



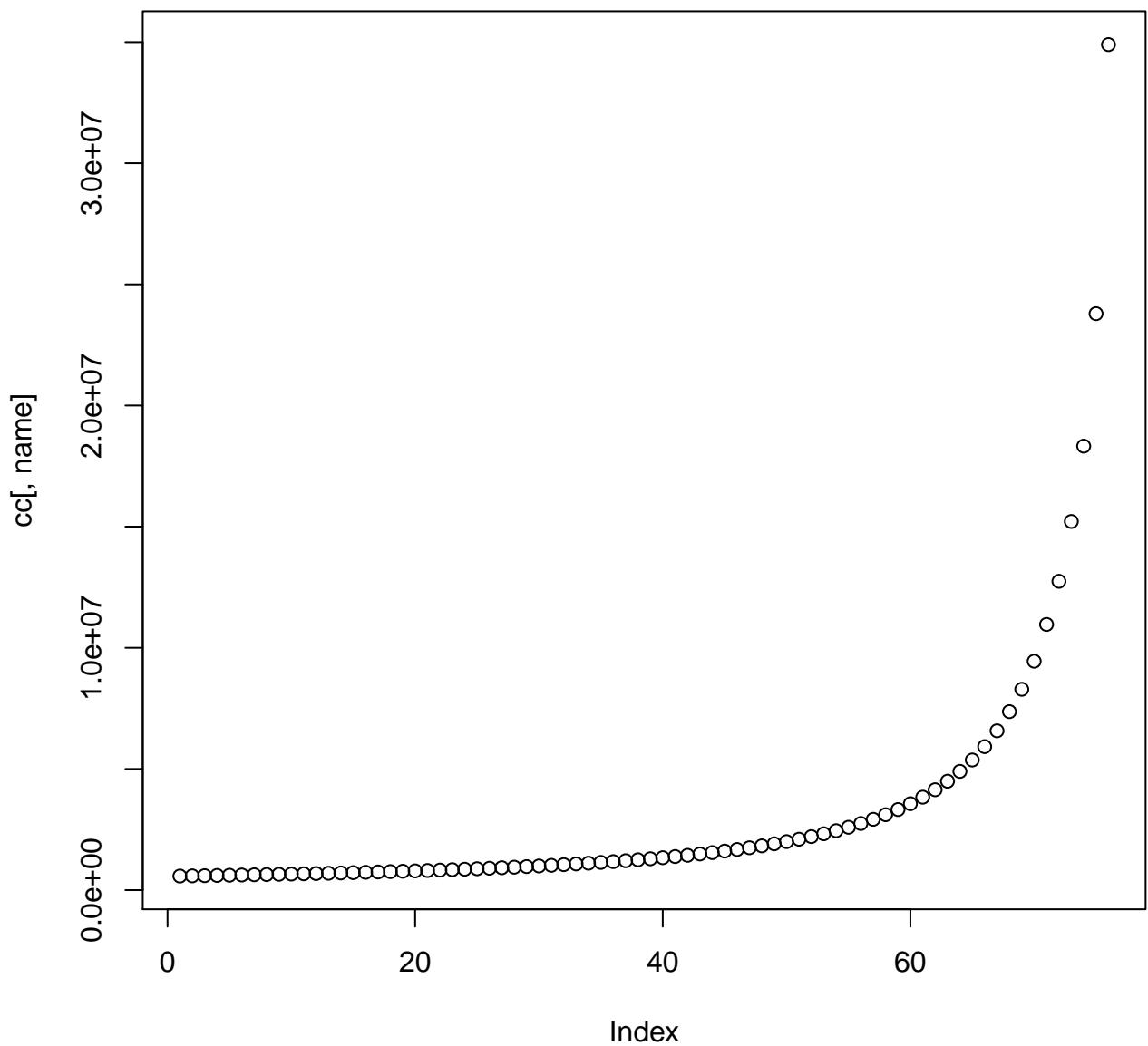
### rsem.approx.bic



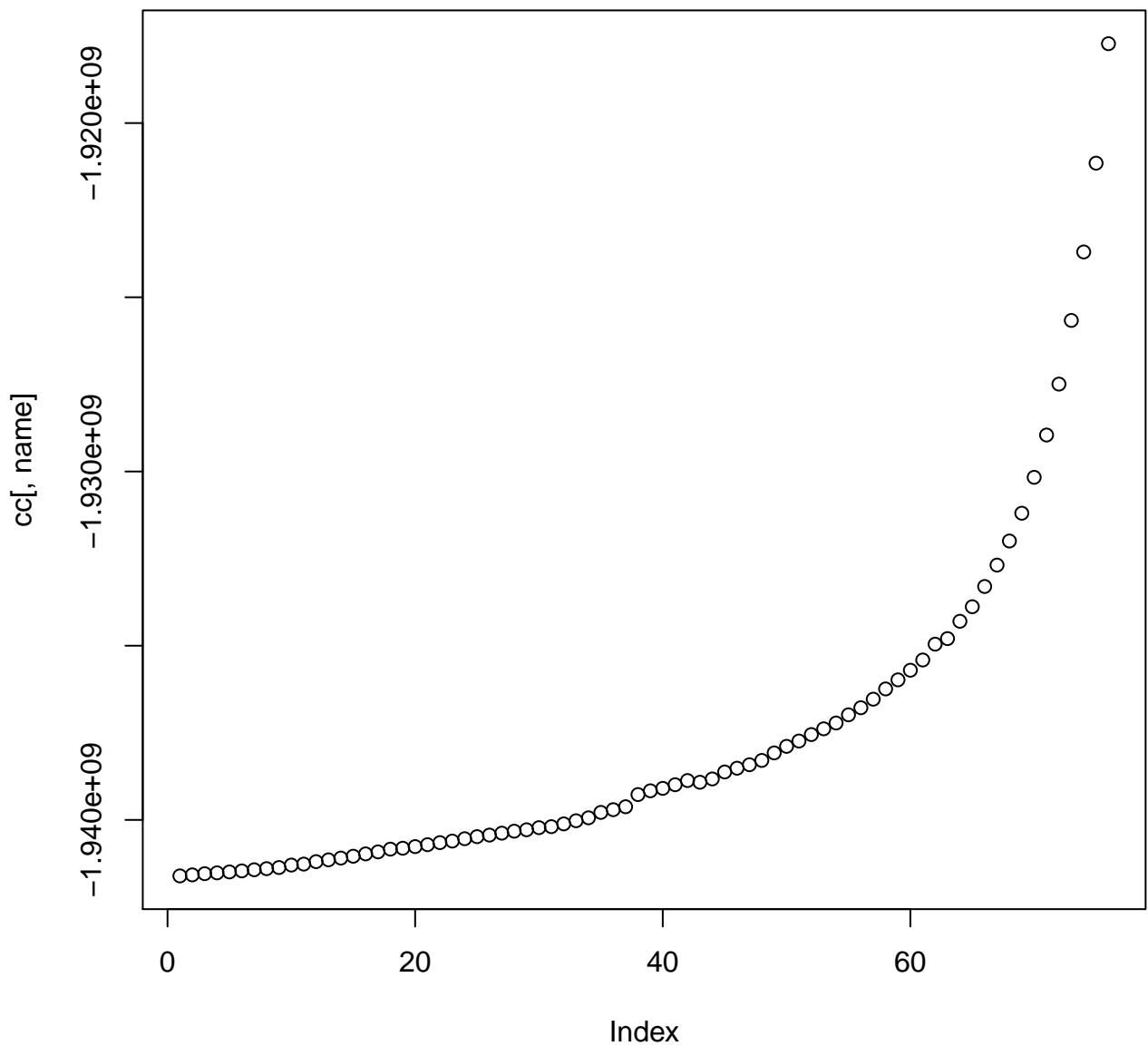
### rsem.approx.loglikelihood



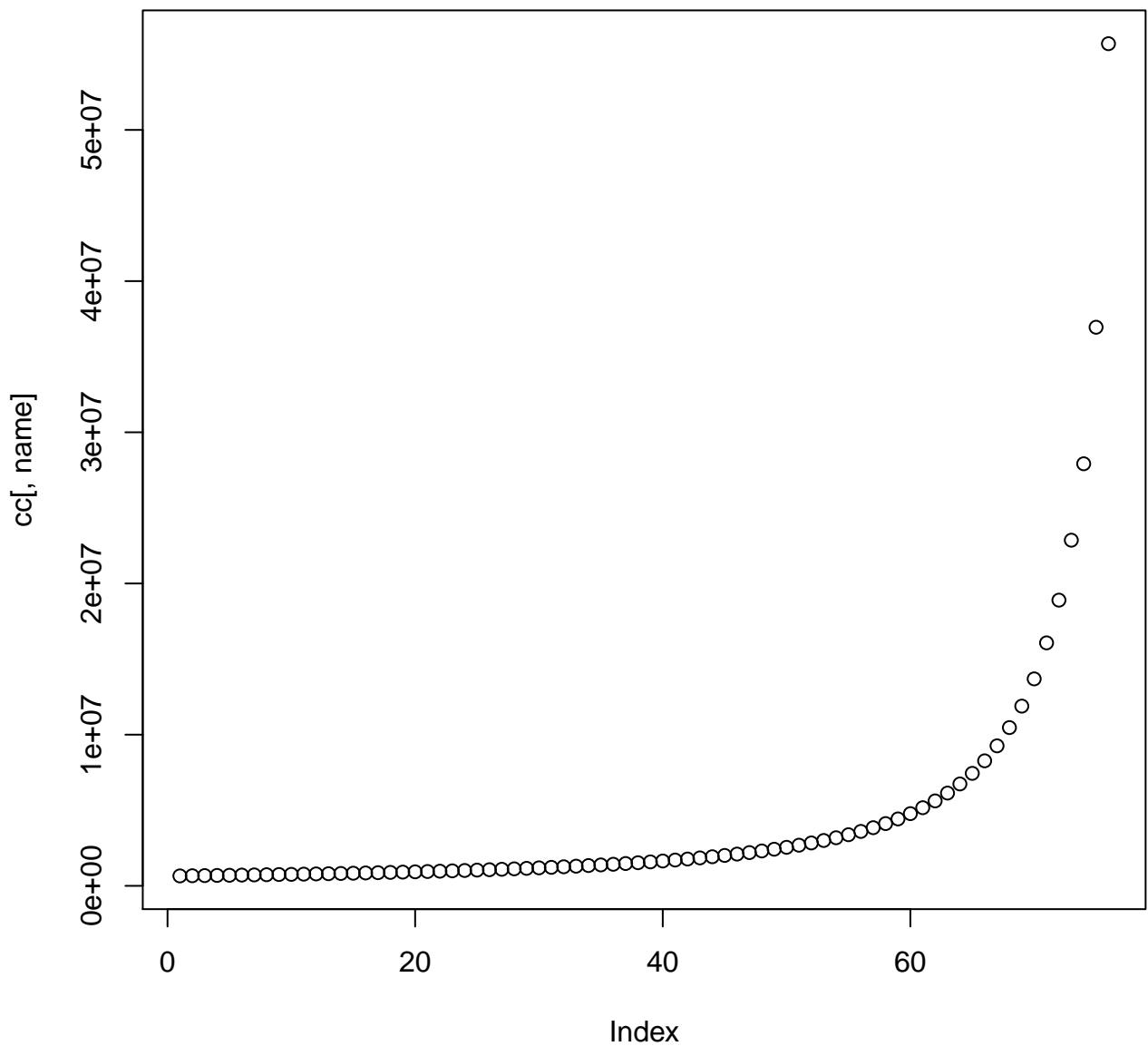
### rsem.approx.loglikelihood.penalty



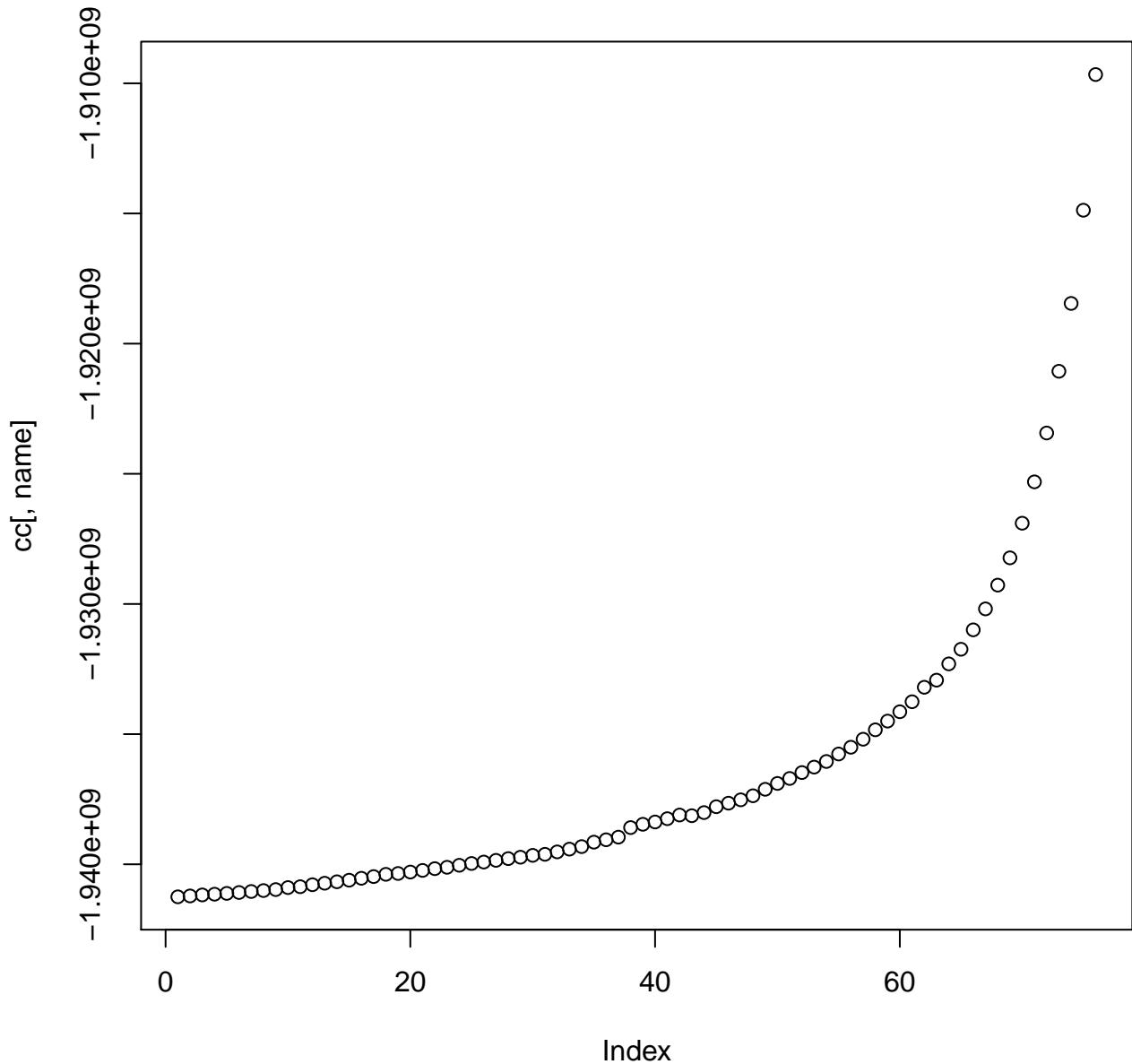
### rsem.eval.lognumer.minus.logdenom



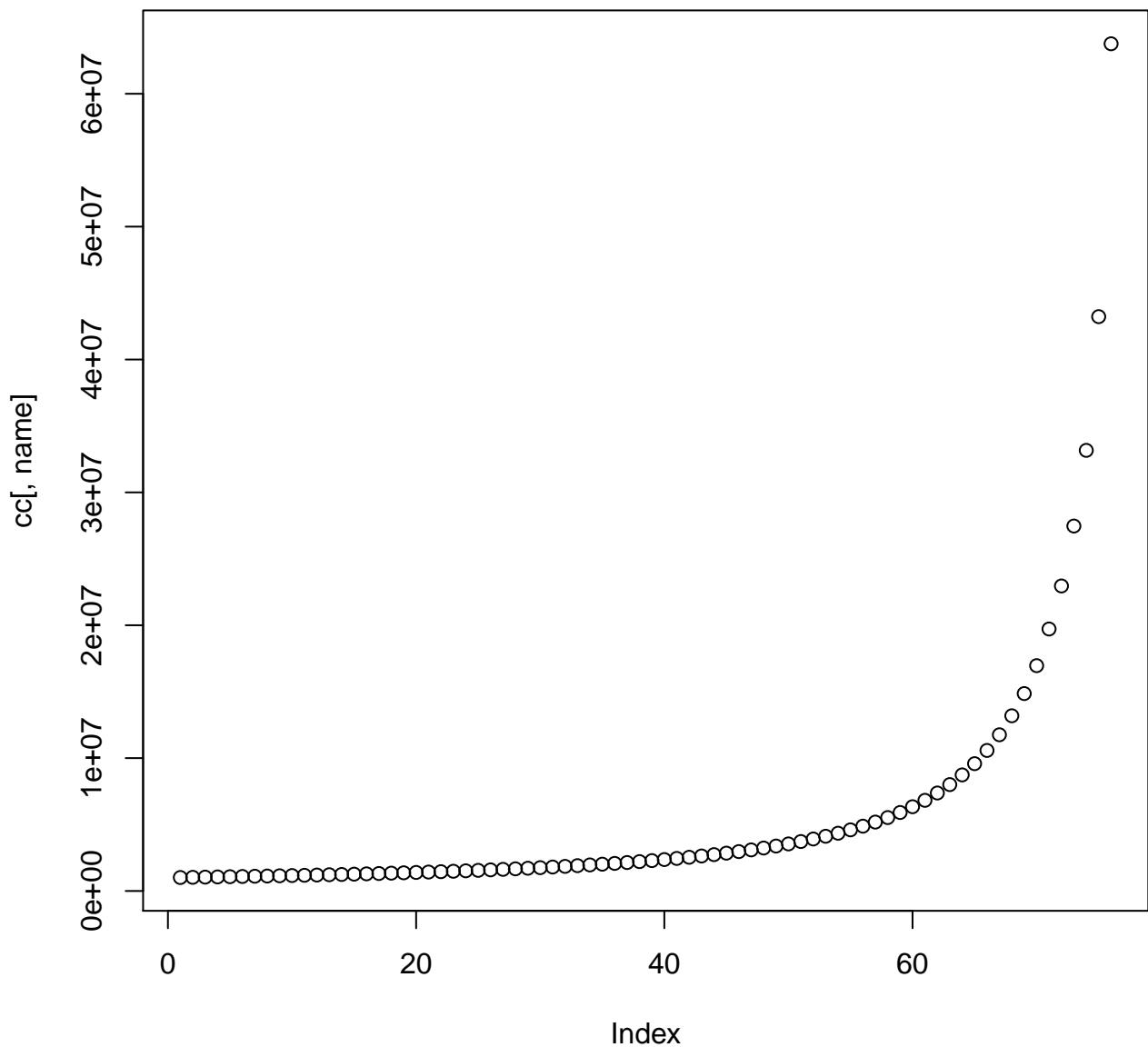
### rsem.eval.logprior



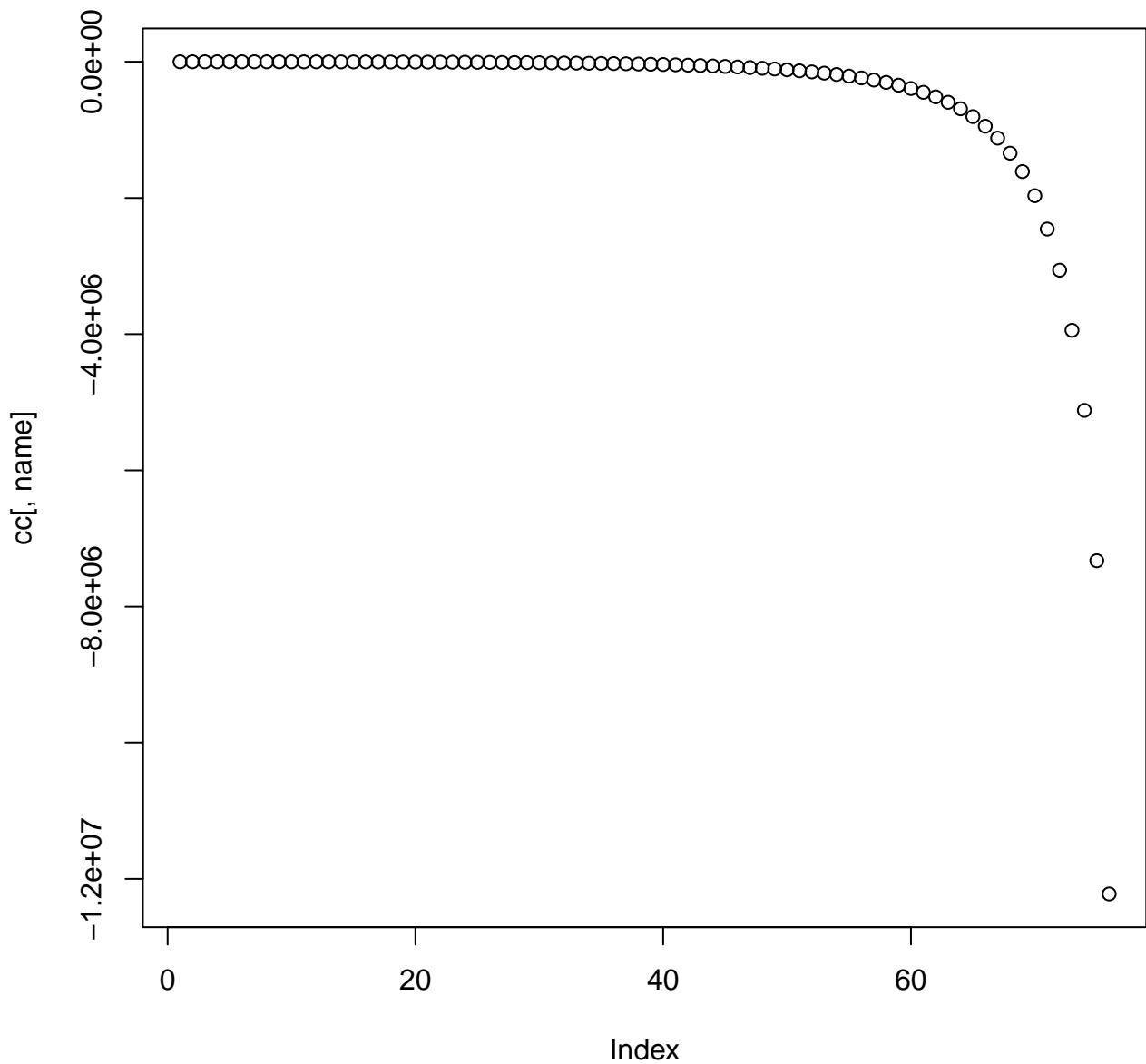
### rsem.eval.loglikelihood



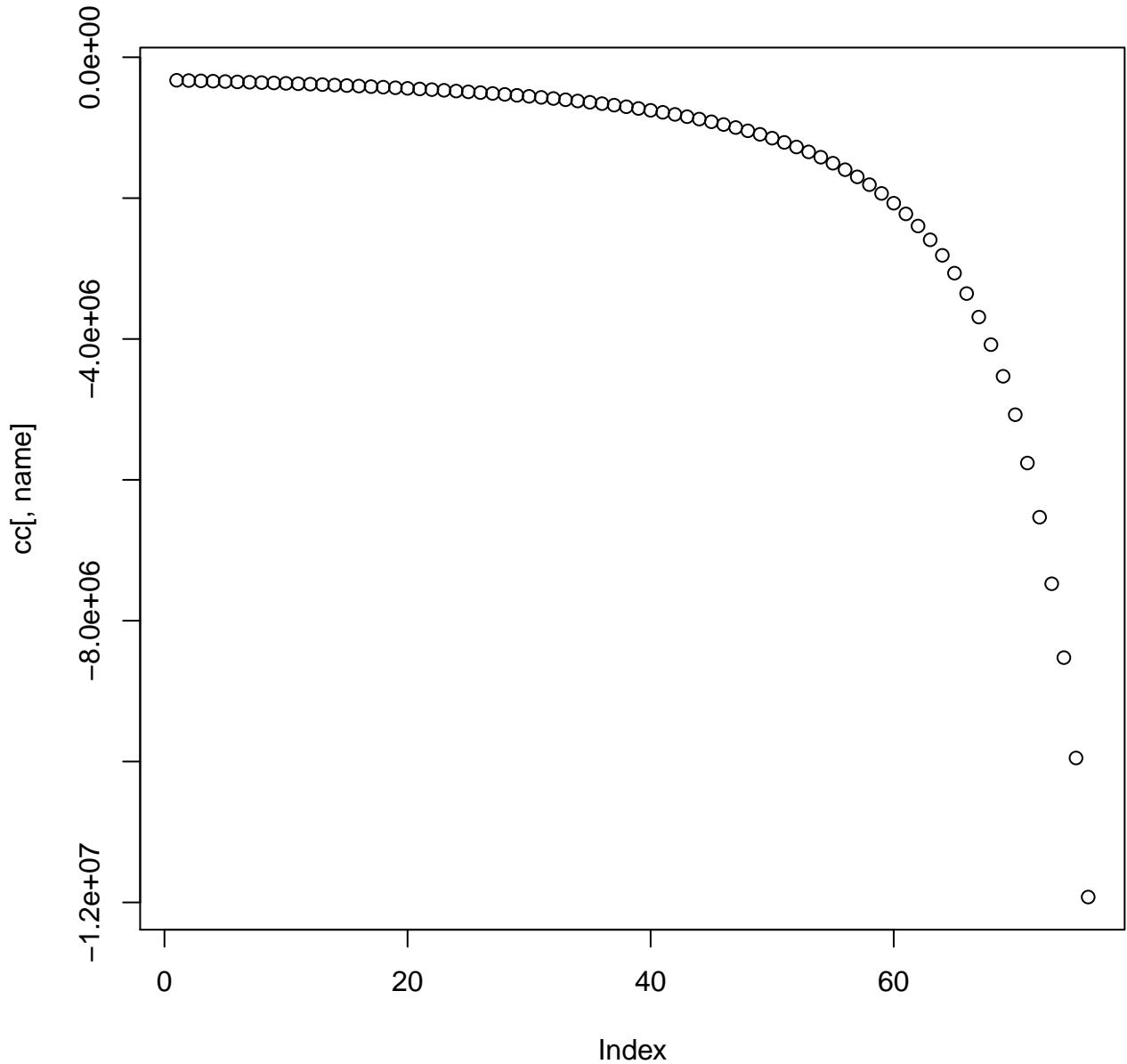
### rsem.eval.logdenom



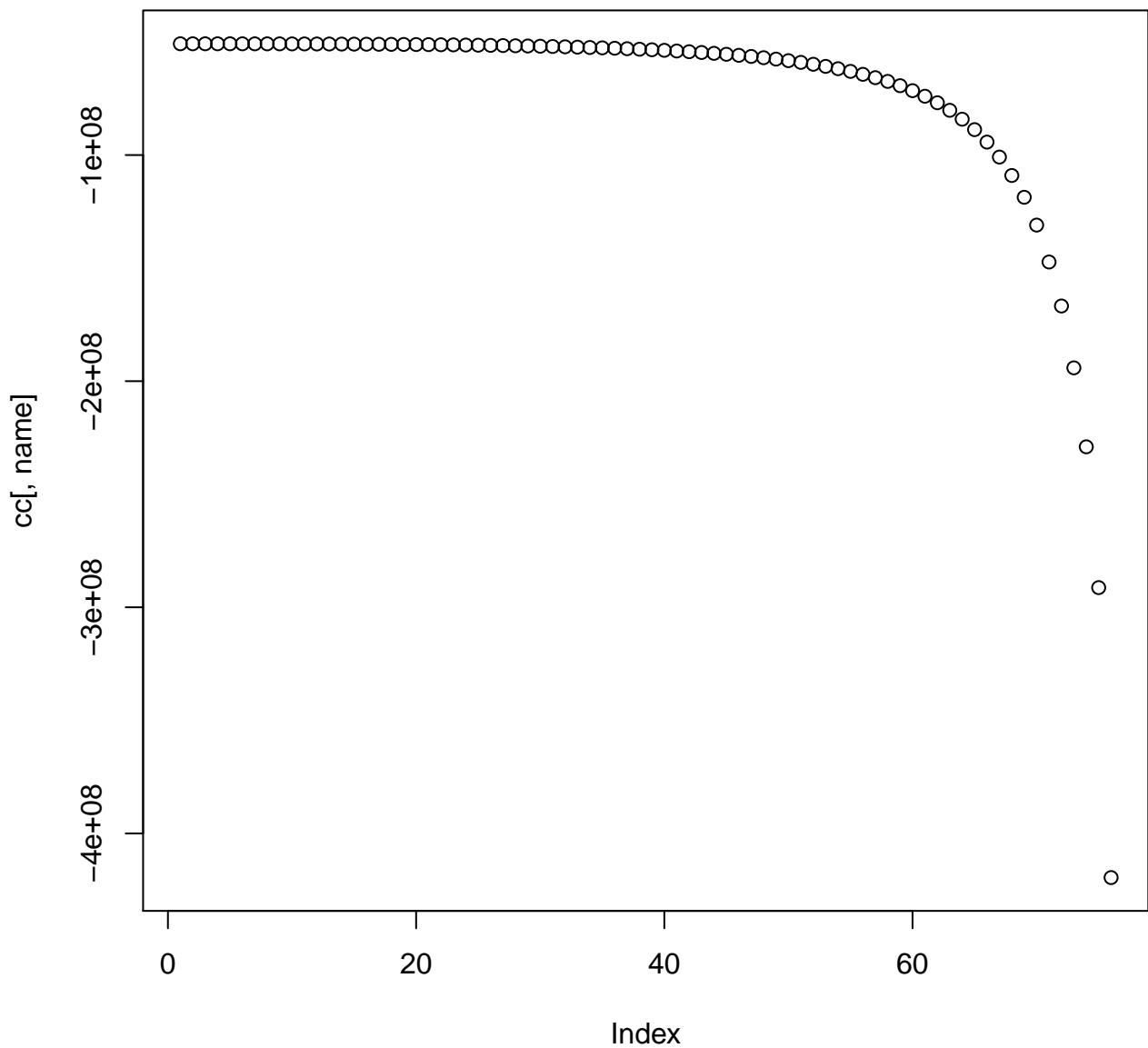
### rsem.prior.log.prob.M



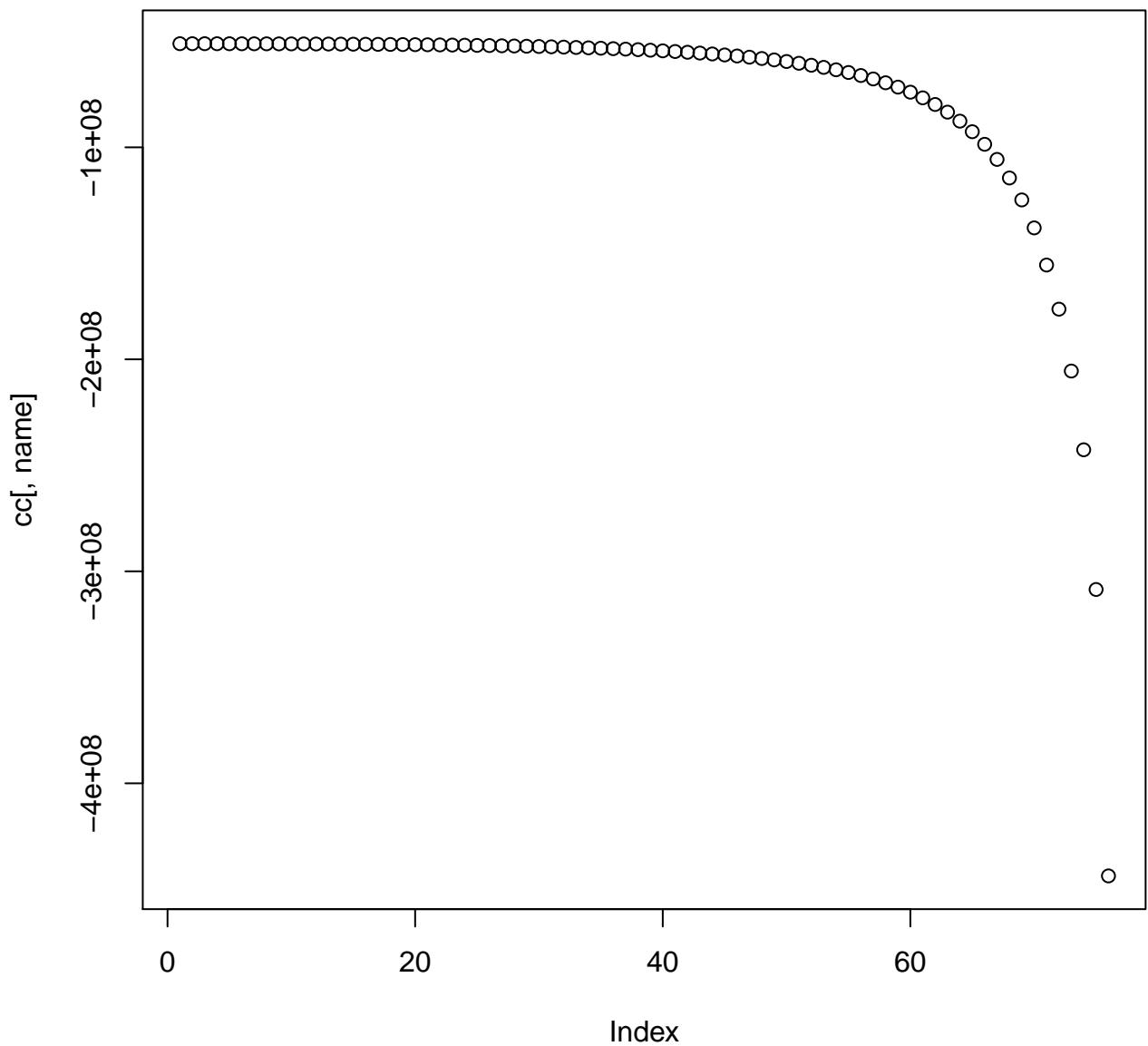
### rsem.prior.log.prob.L.given.M



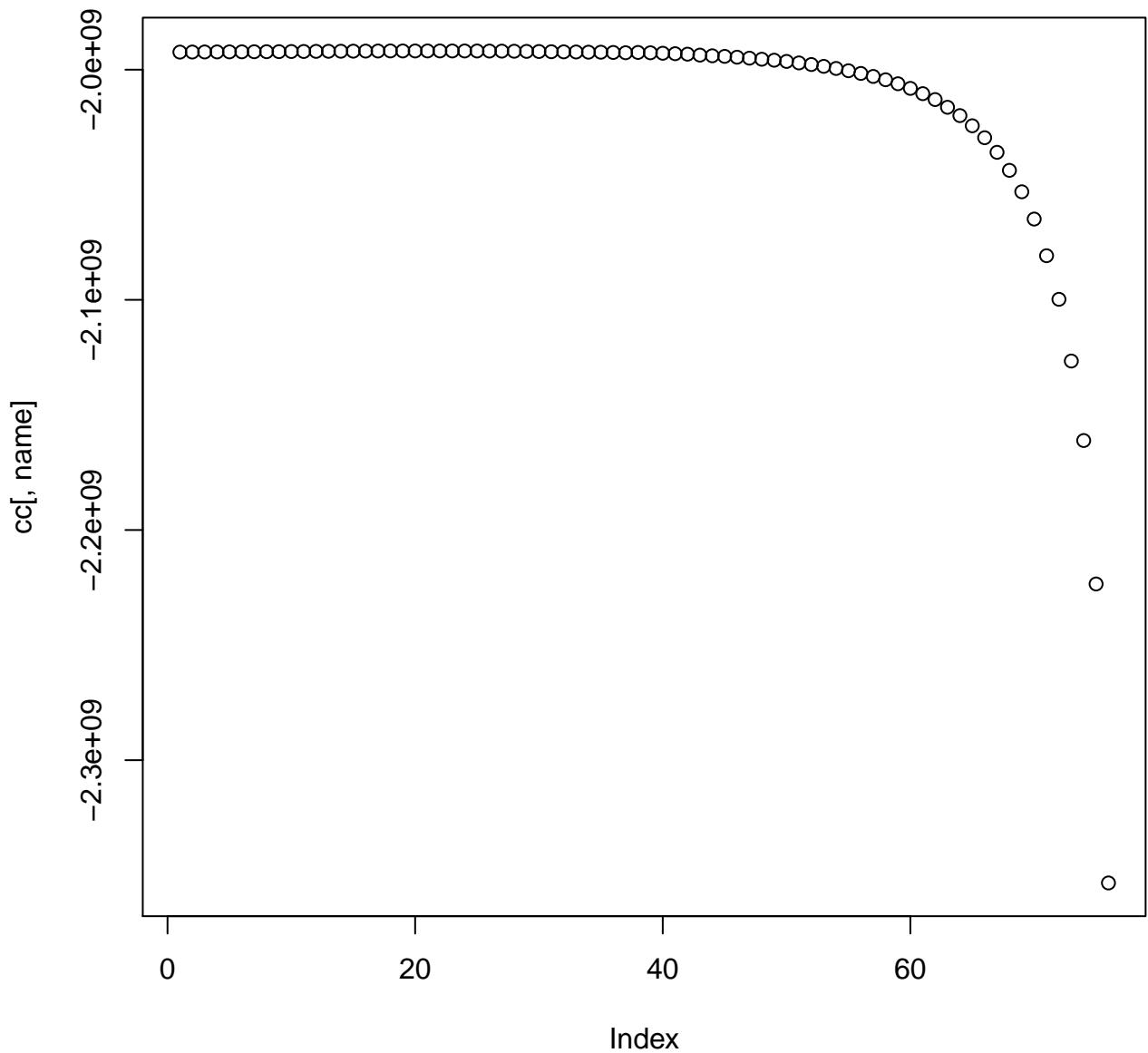
### rsem.prior.log.prob.Sequences.given.L.and.M



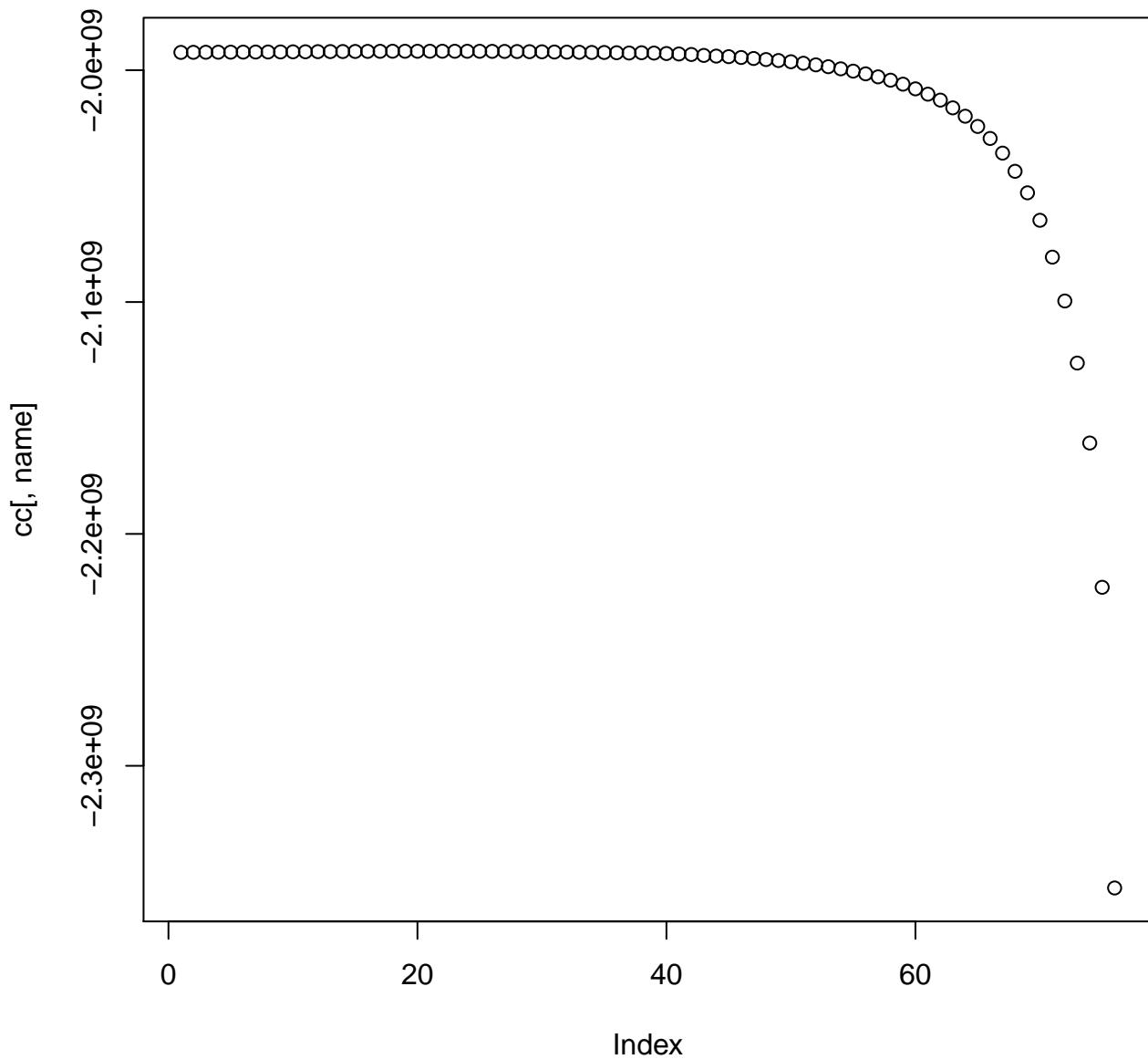
**rsem.prior.log.prob.A**



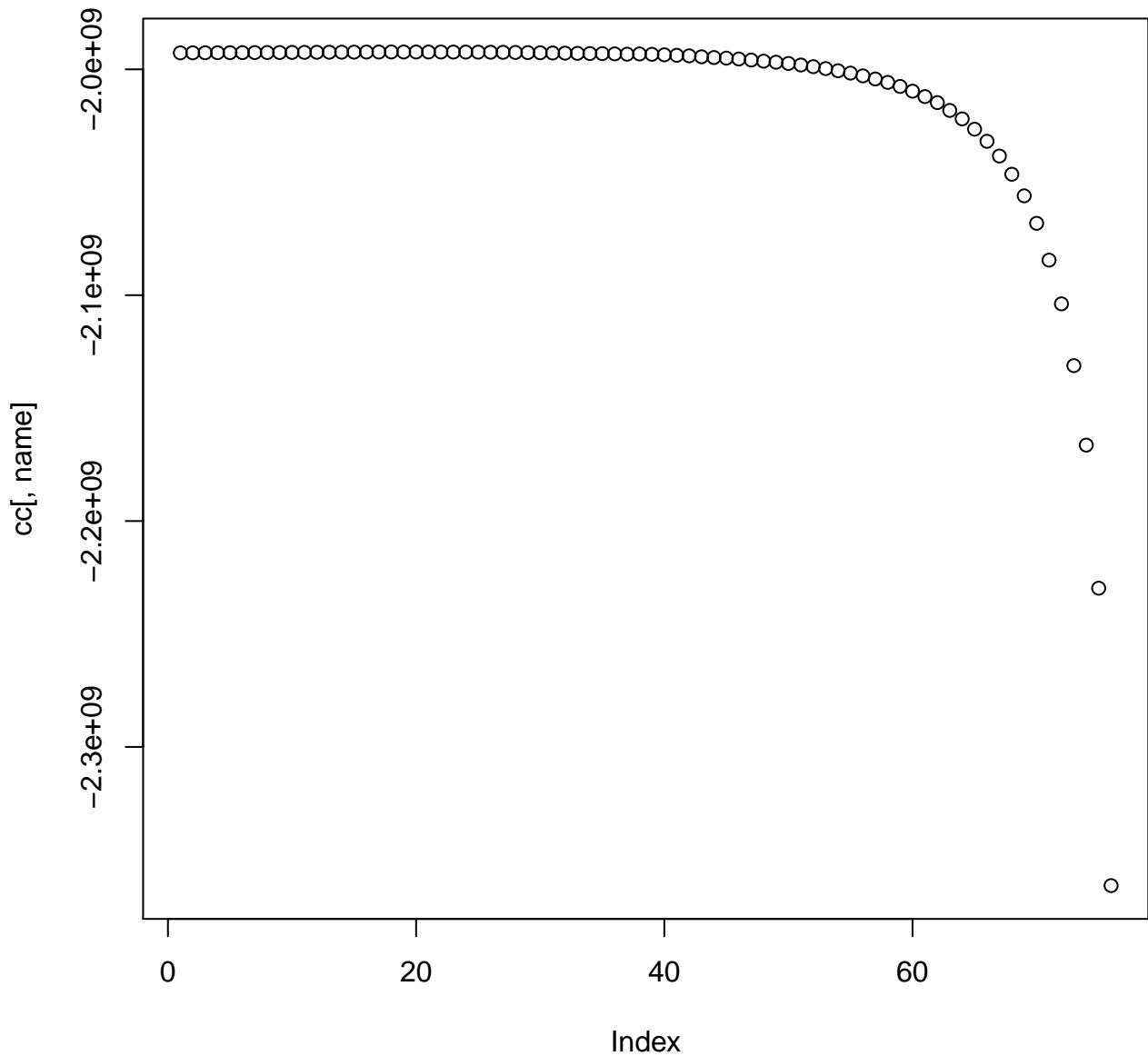
### rsem.eval.loglikelihood.plus.rsem.prior.log.prob.A



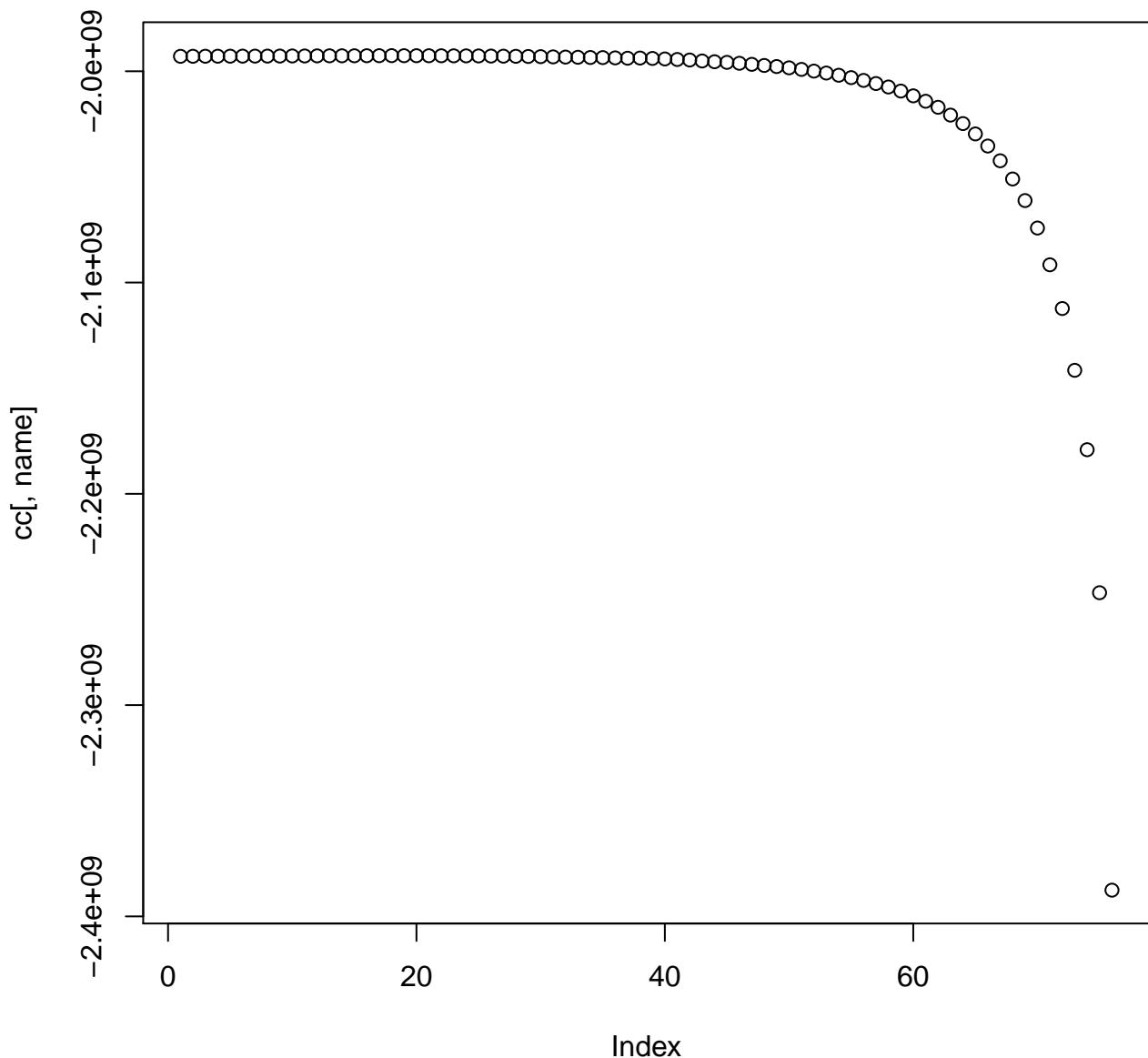
### rsem.approx.loglikelihood.plus.rsem.prior.log.prob.A



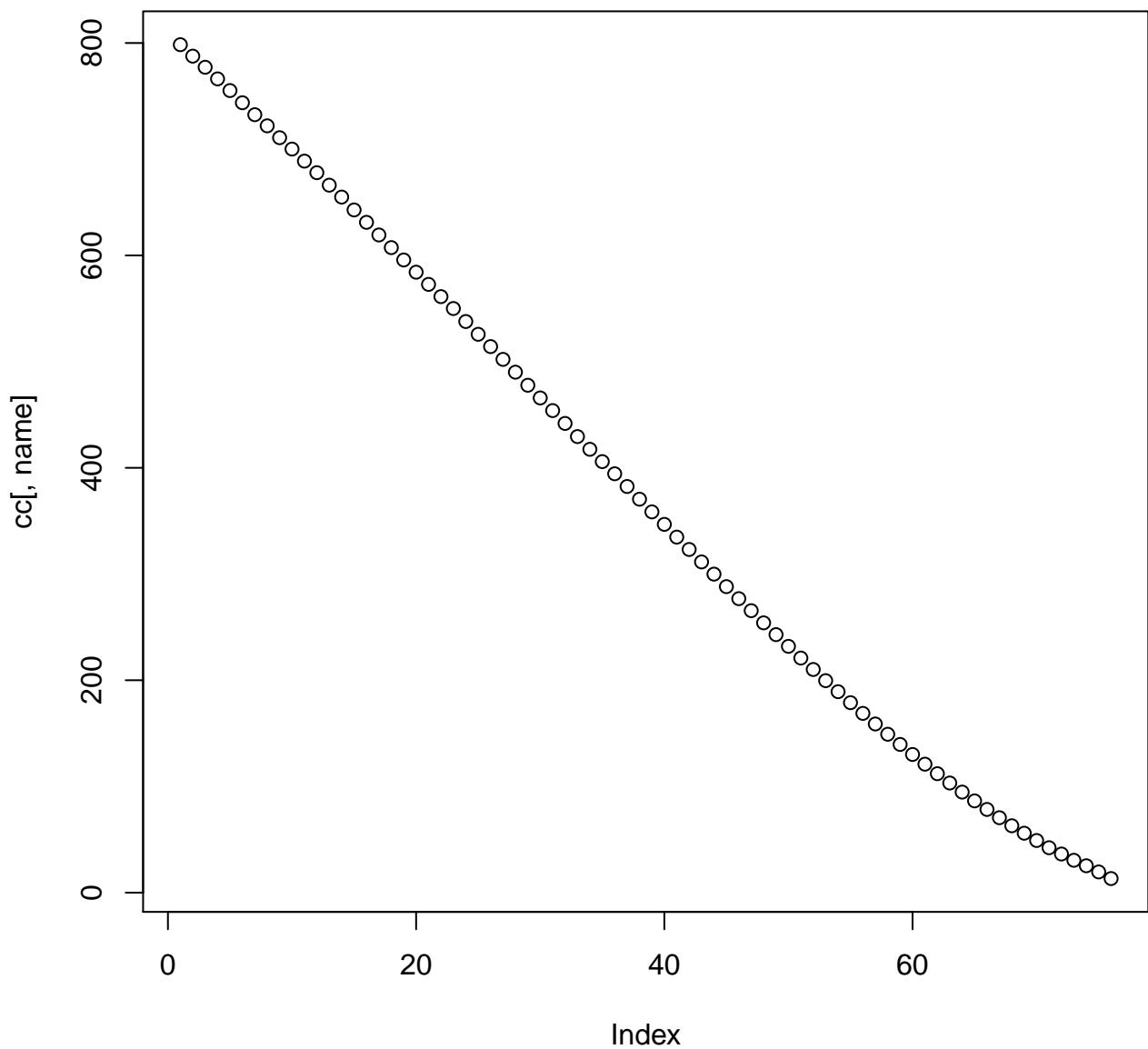
### rsem.approx.approx.plus.rsem.prior.log.prob.A



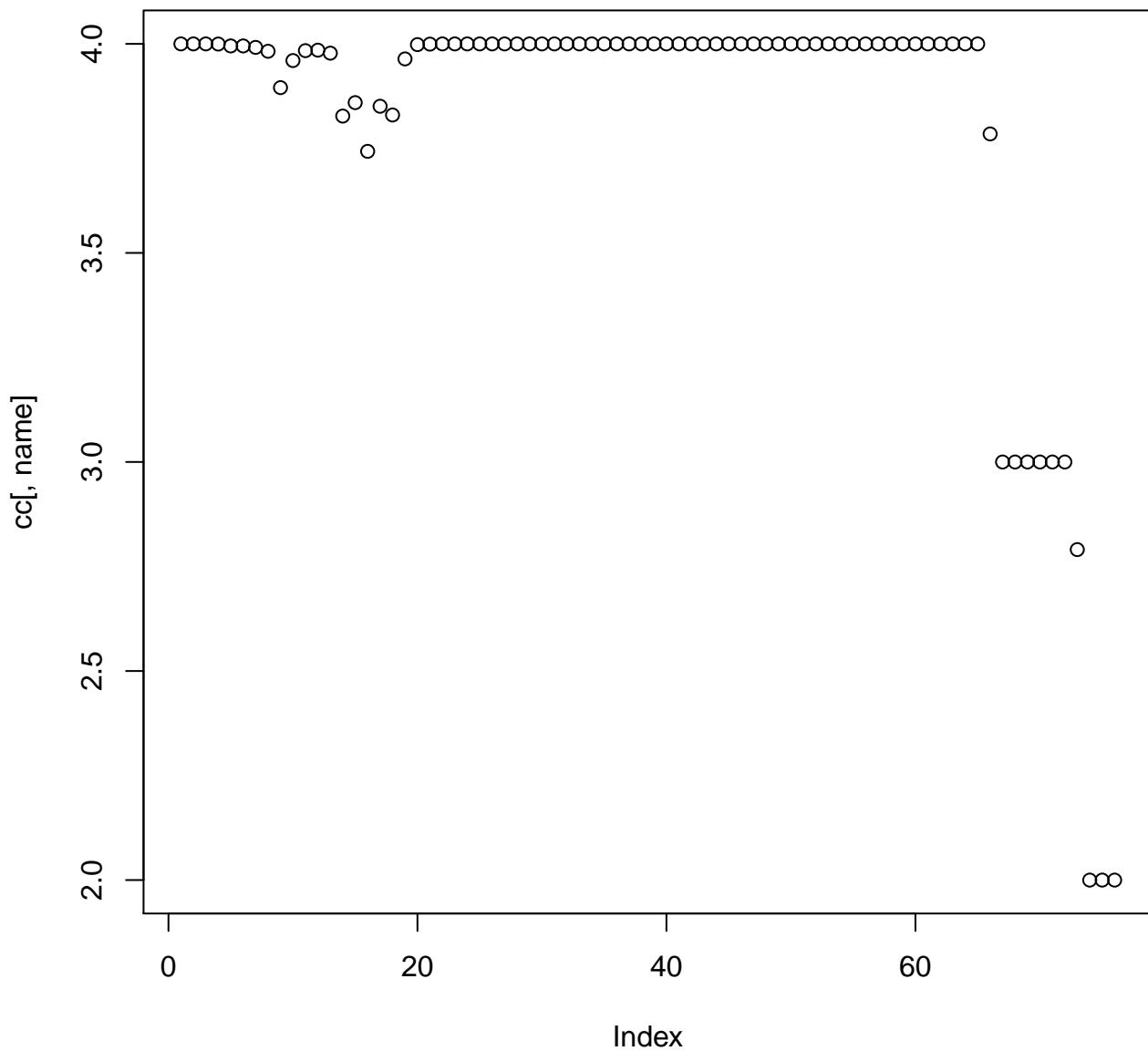
## rsem.approx.bic.plus.rsem.prior.log.prob.A



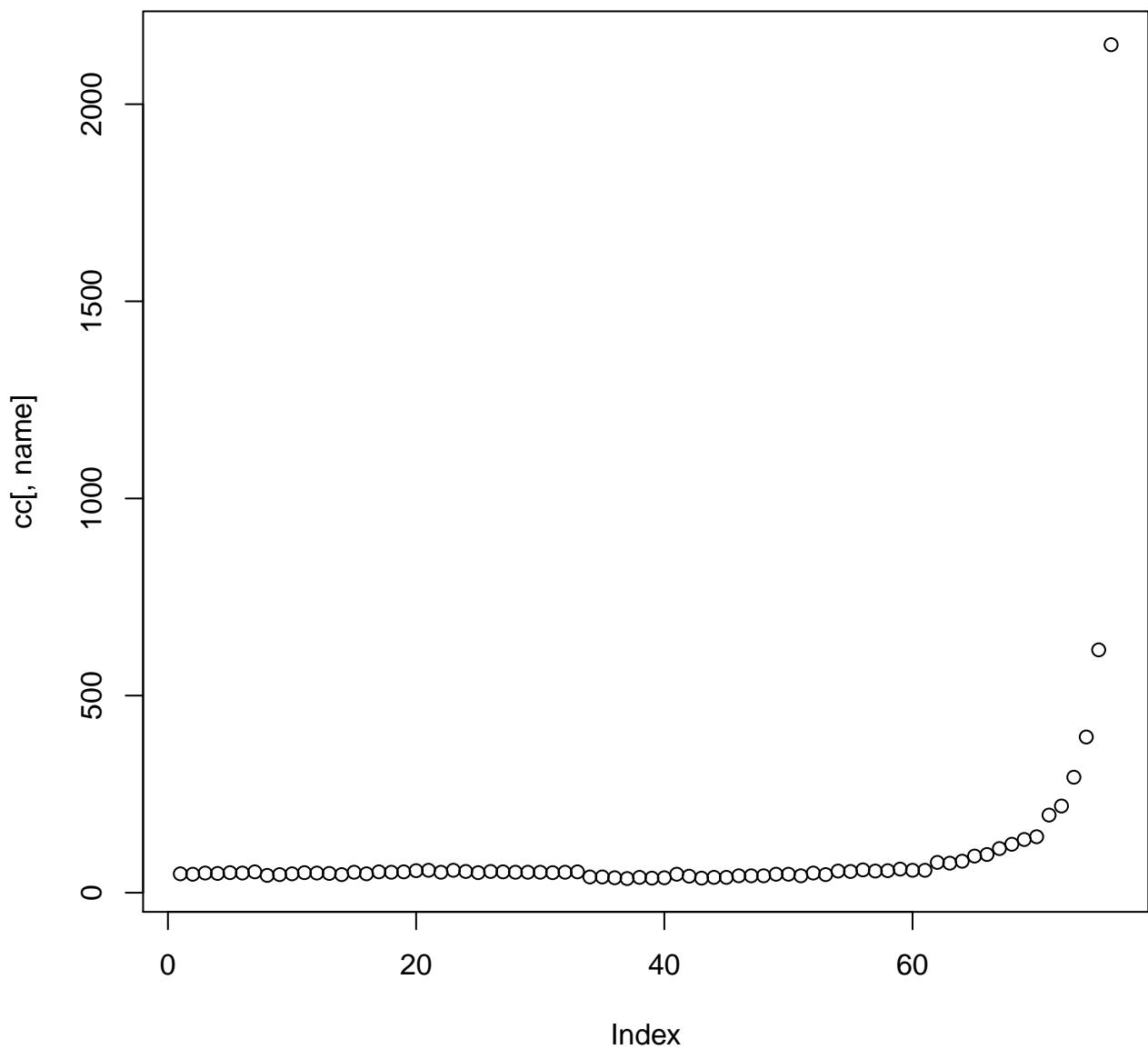
### rsem.ss.mean.num.reads.per.transcript



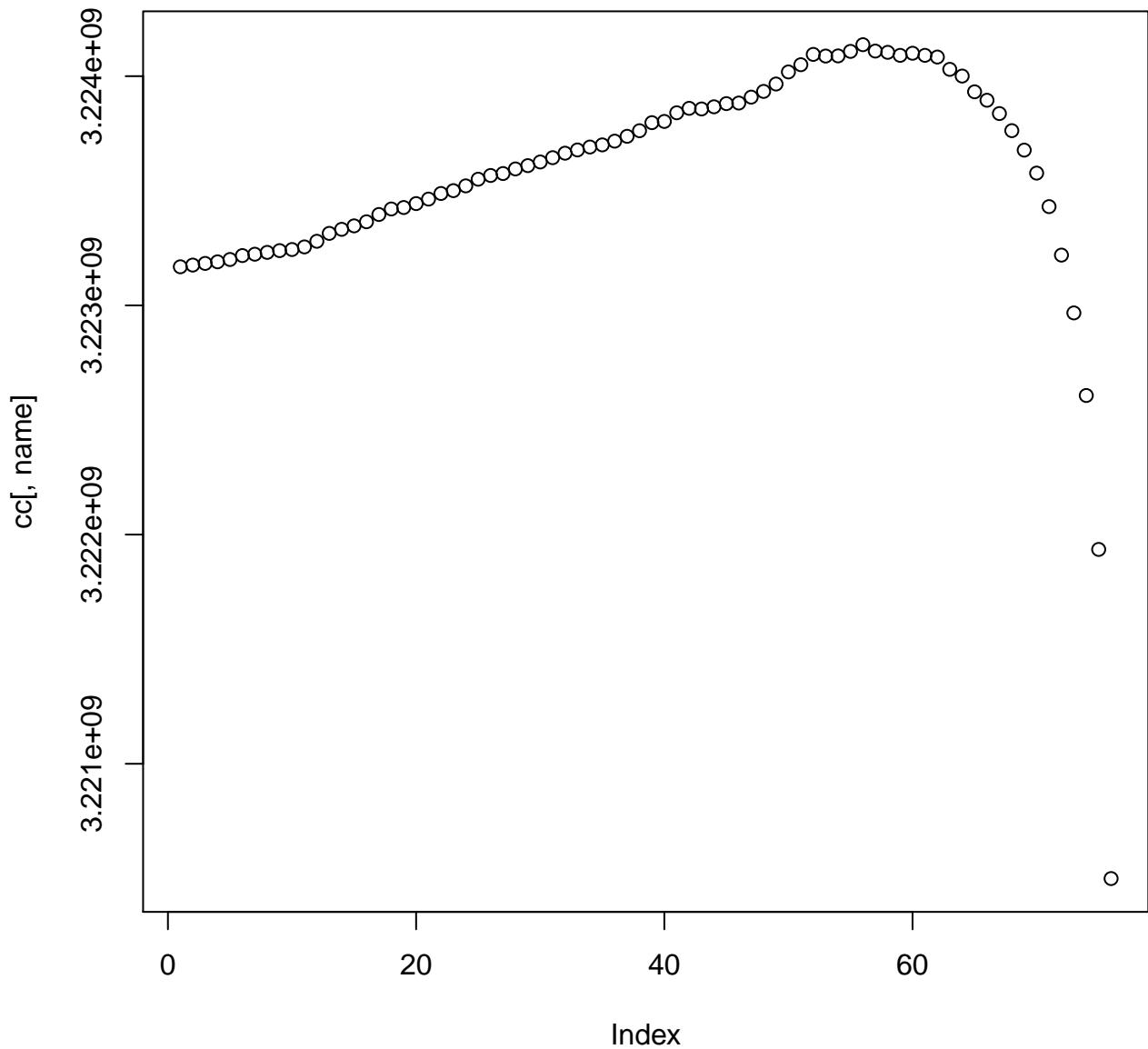
### rsem.ss.median.num.reads.per.transcript



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



### **rsem.ss.num.matching.bases**



### rsem.ss.num.mismatching.bases

