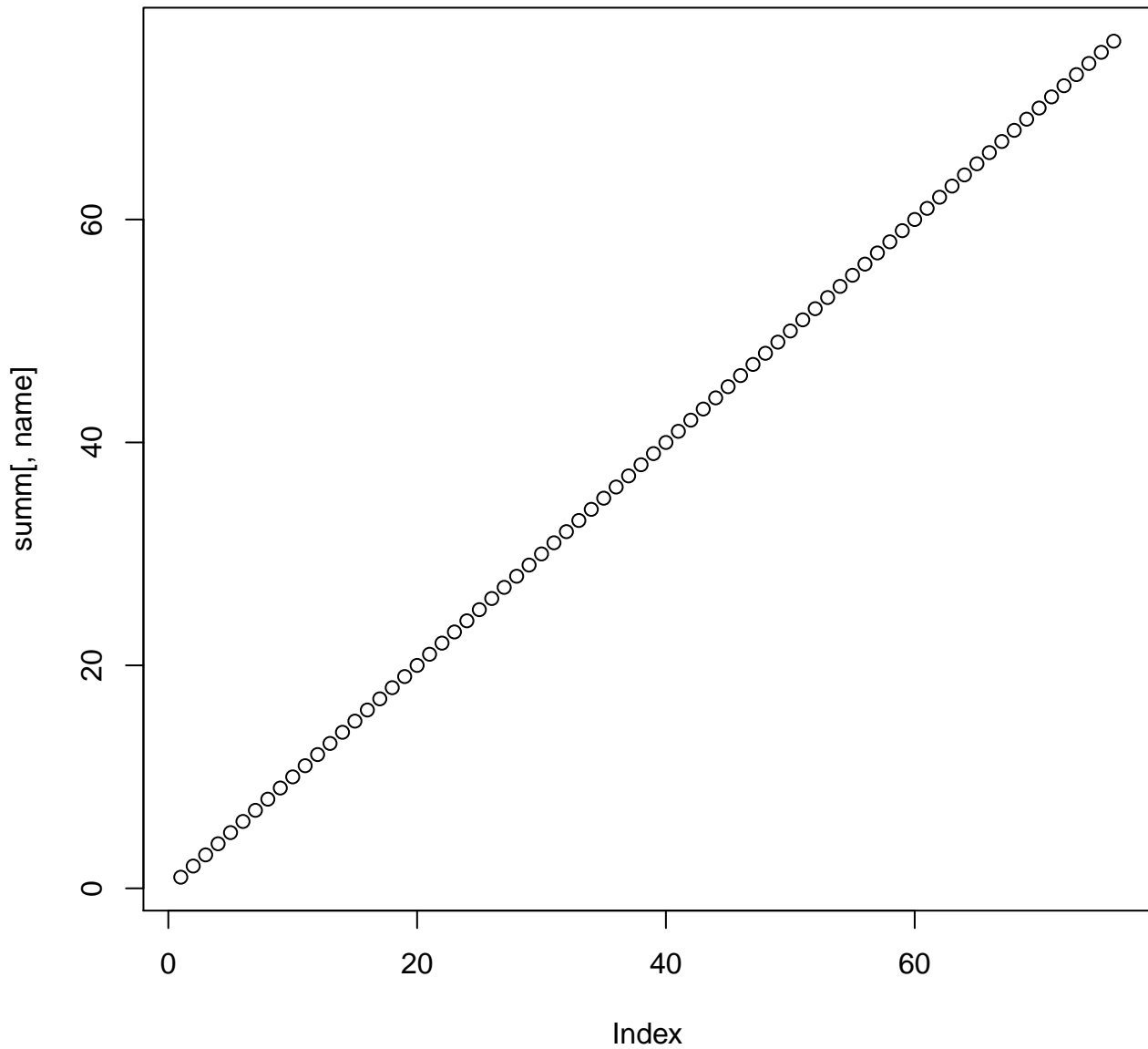
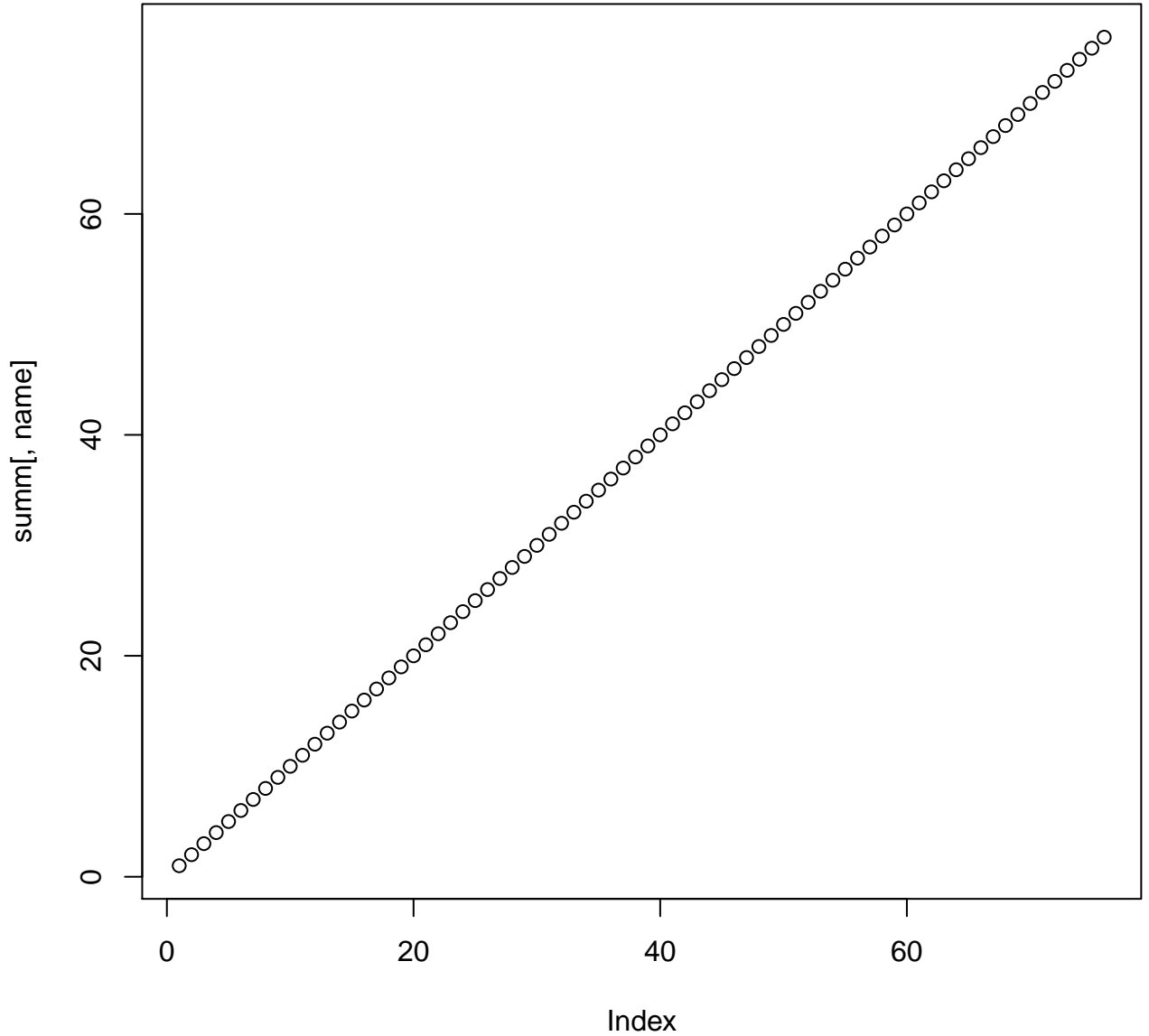


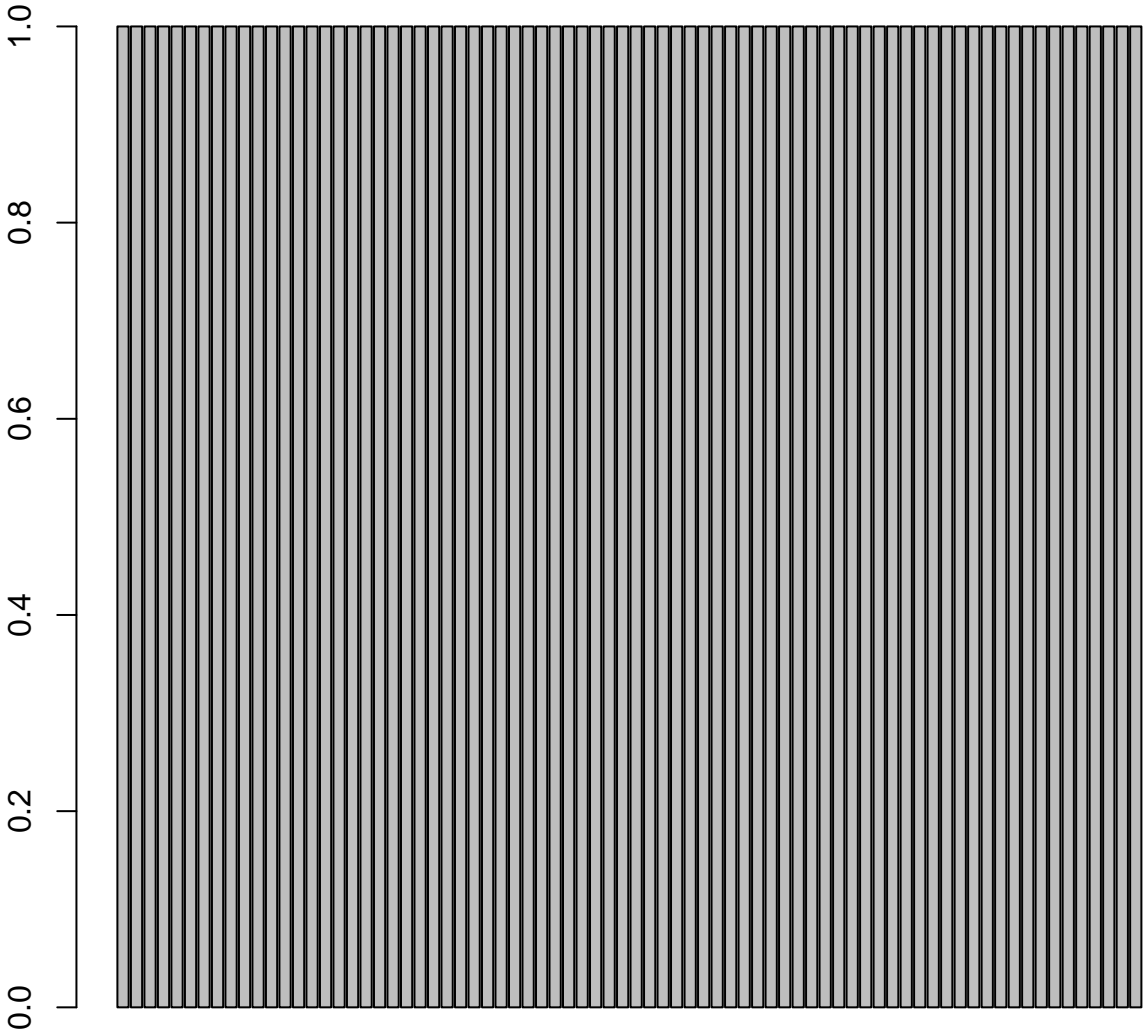
X



X.1

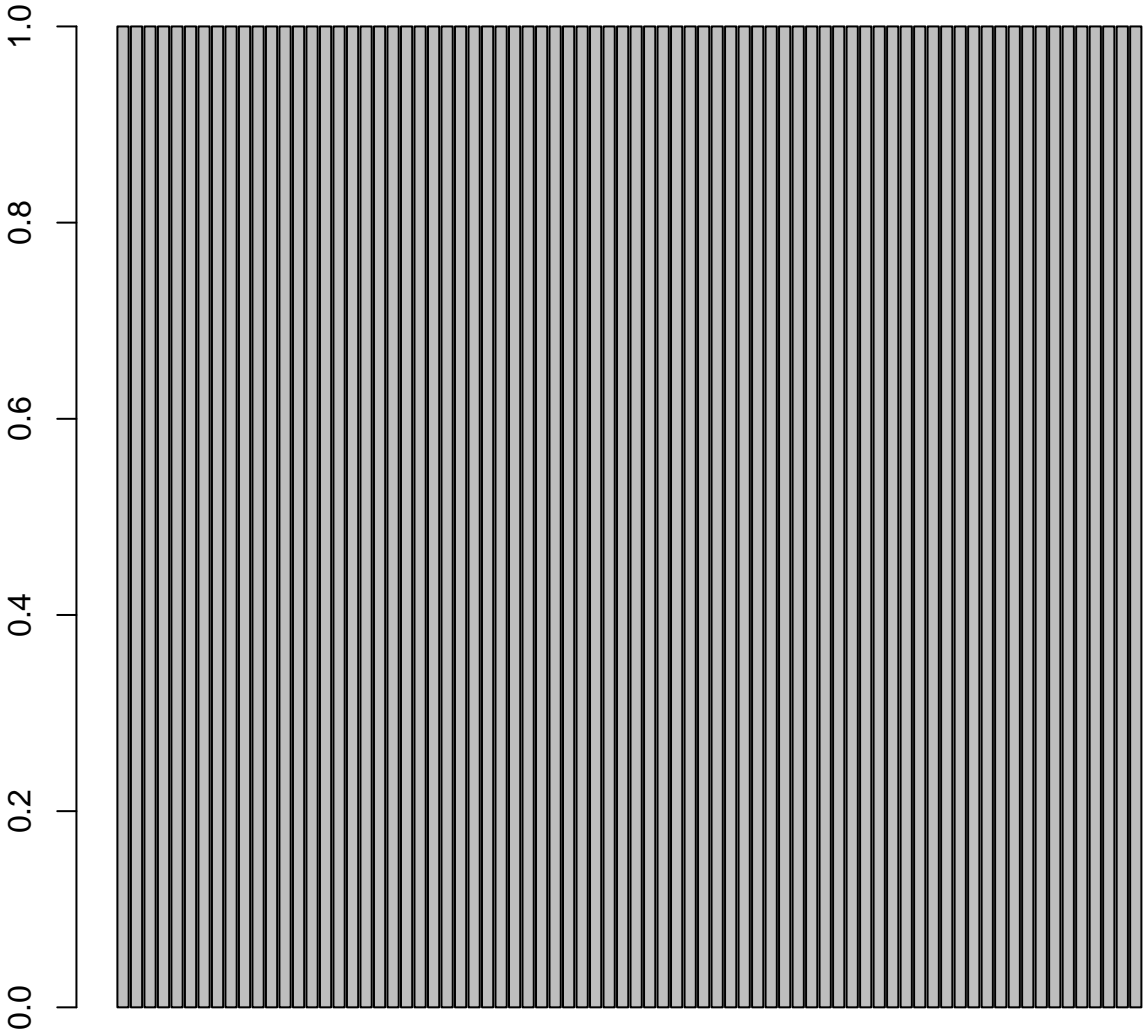


assembly



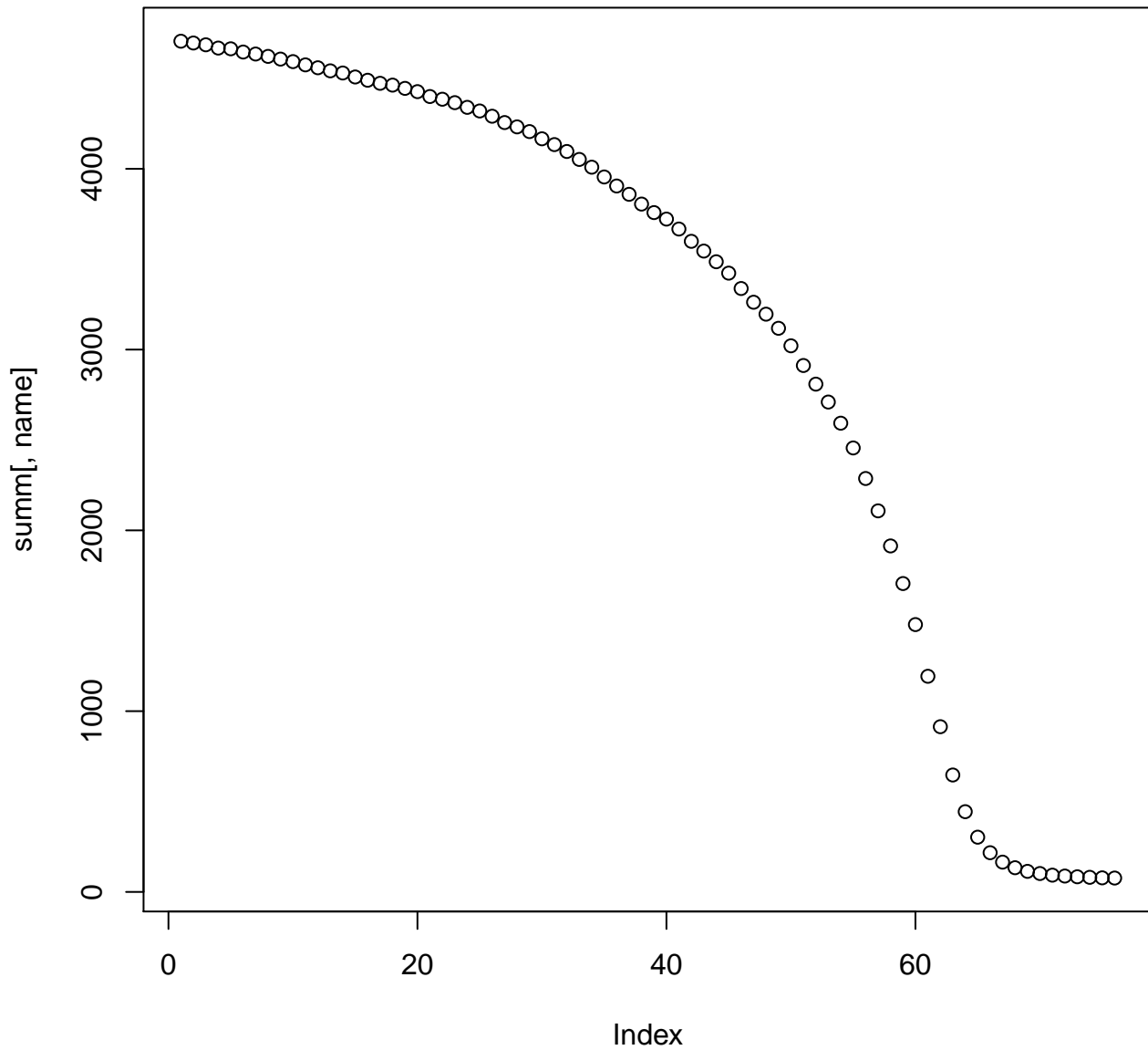
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summary

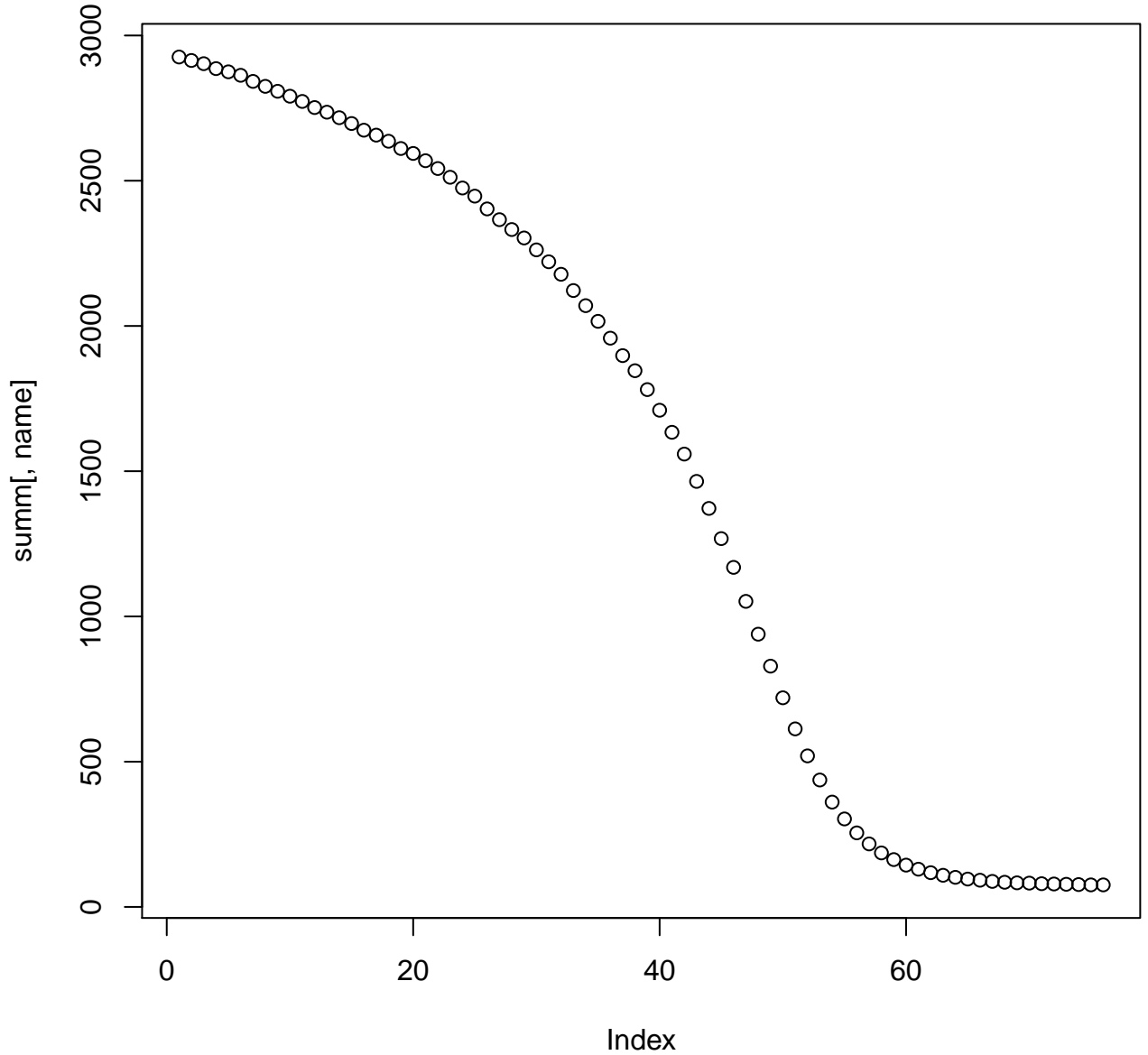


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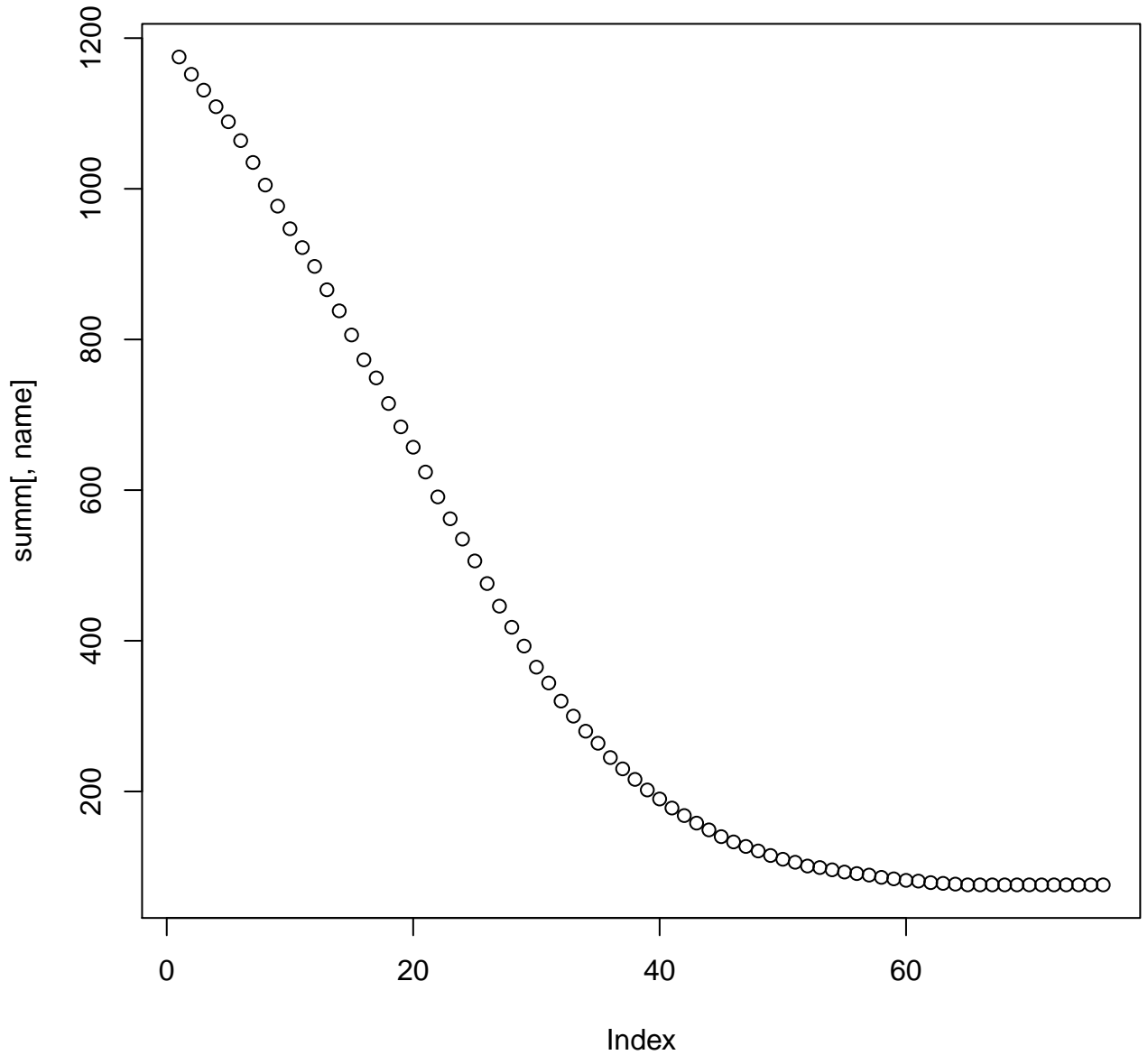
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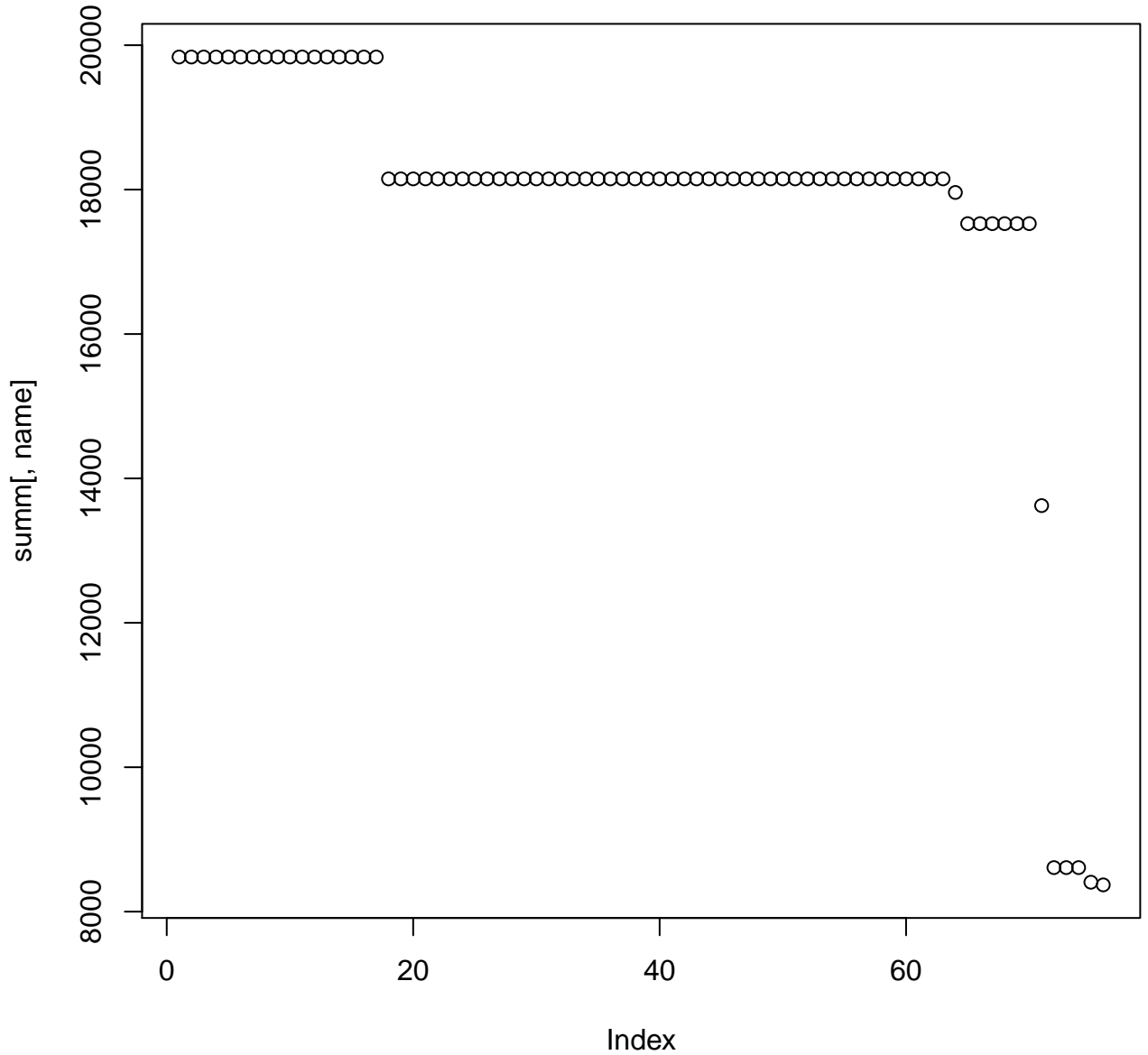
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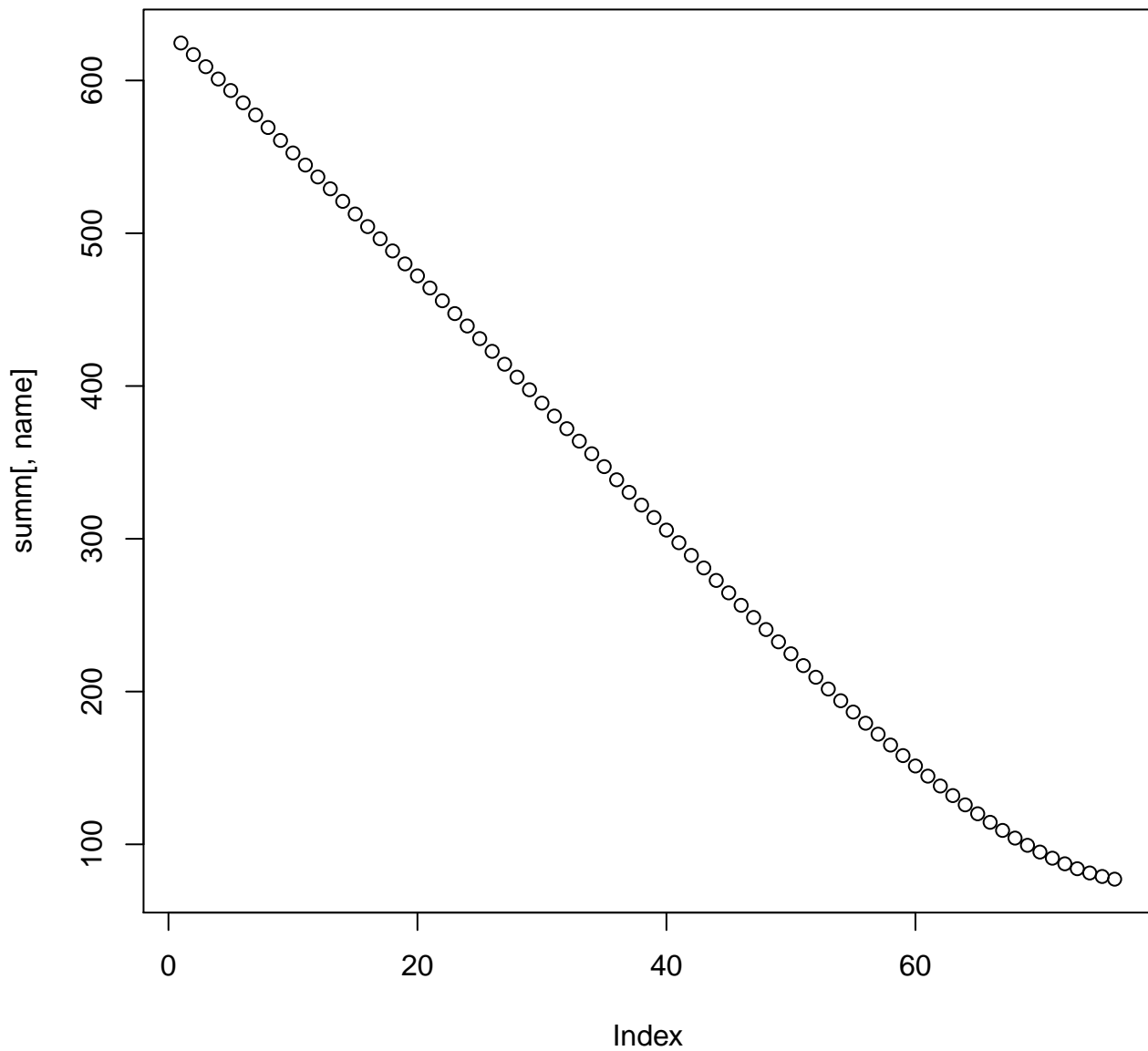
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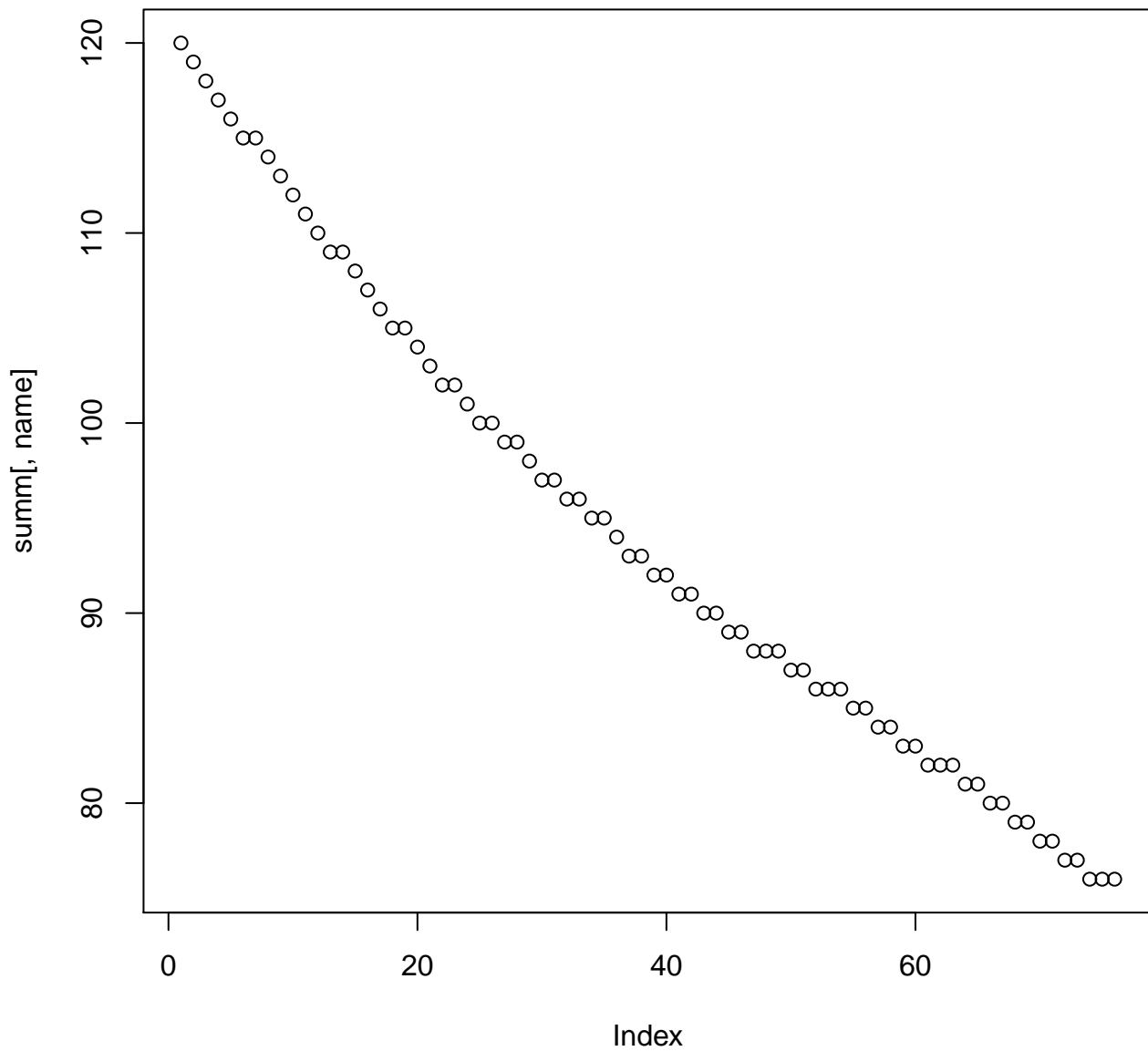
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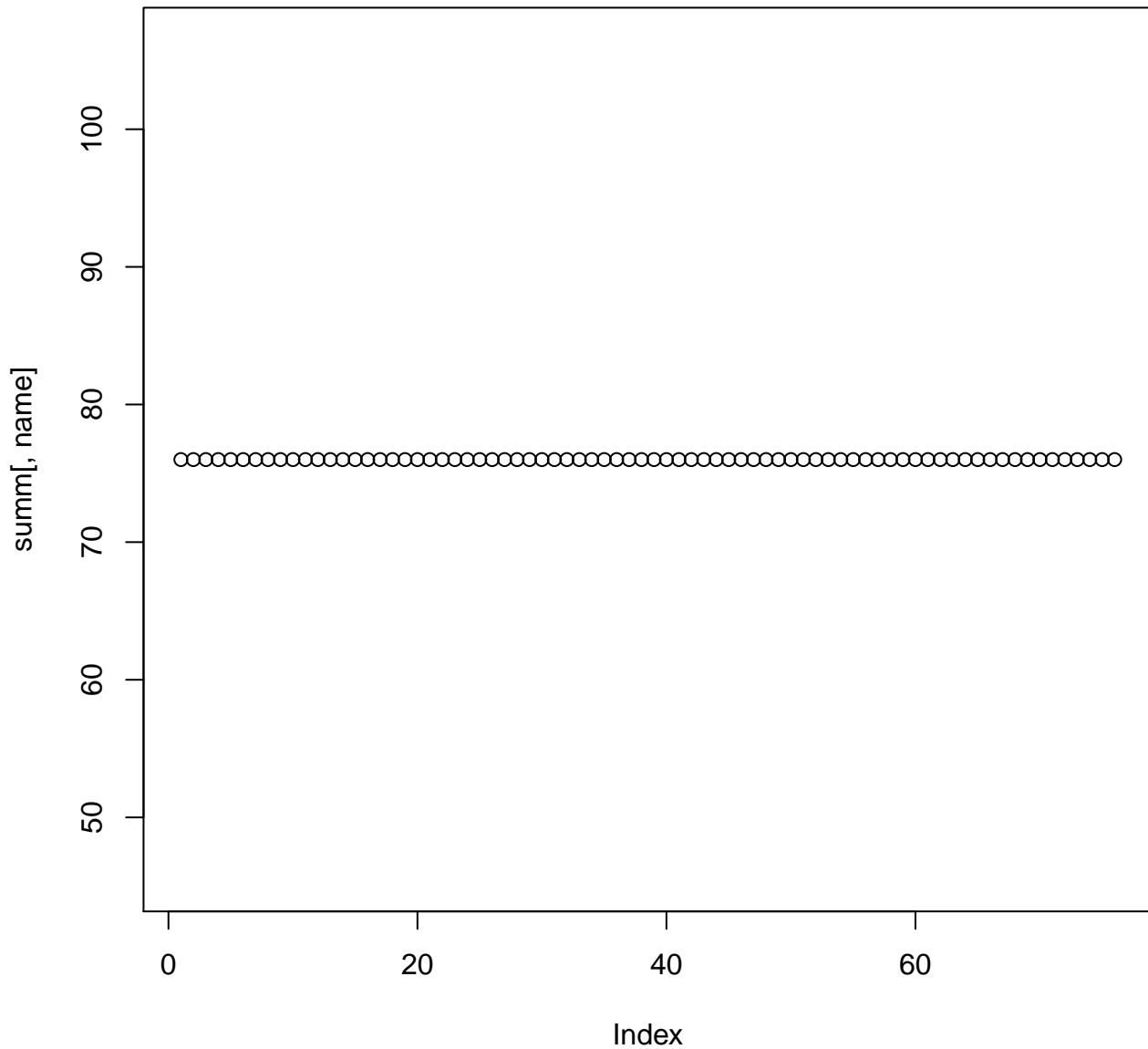
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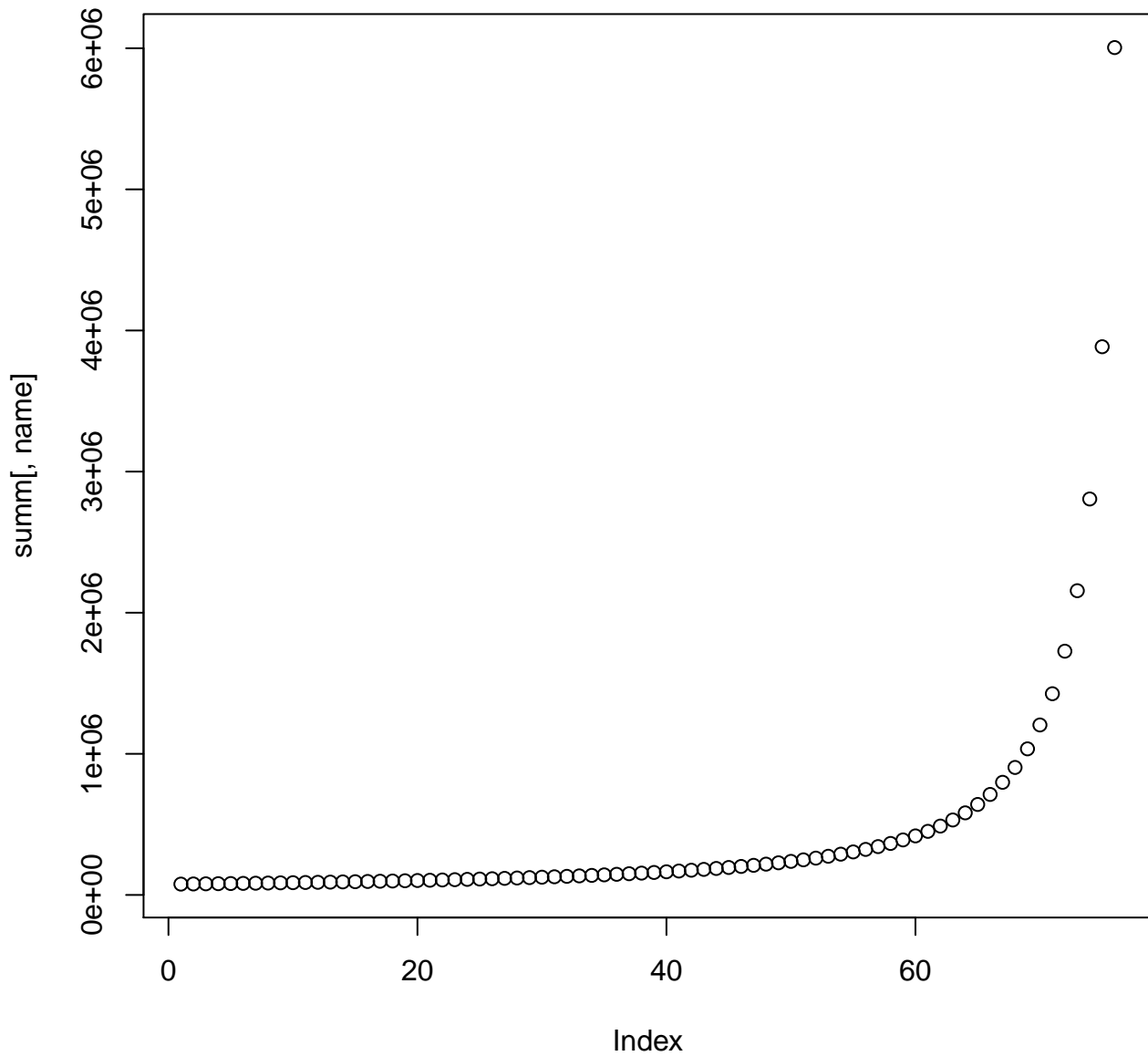
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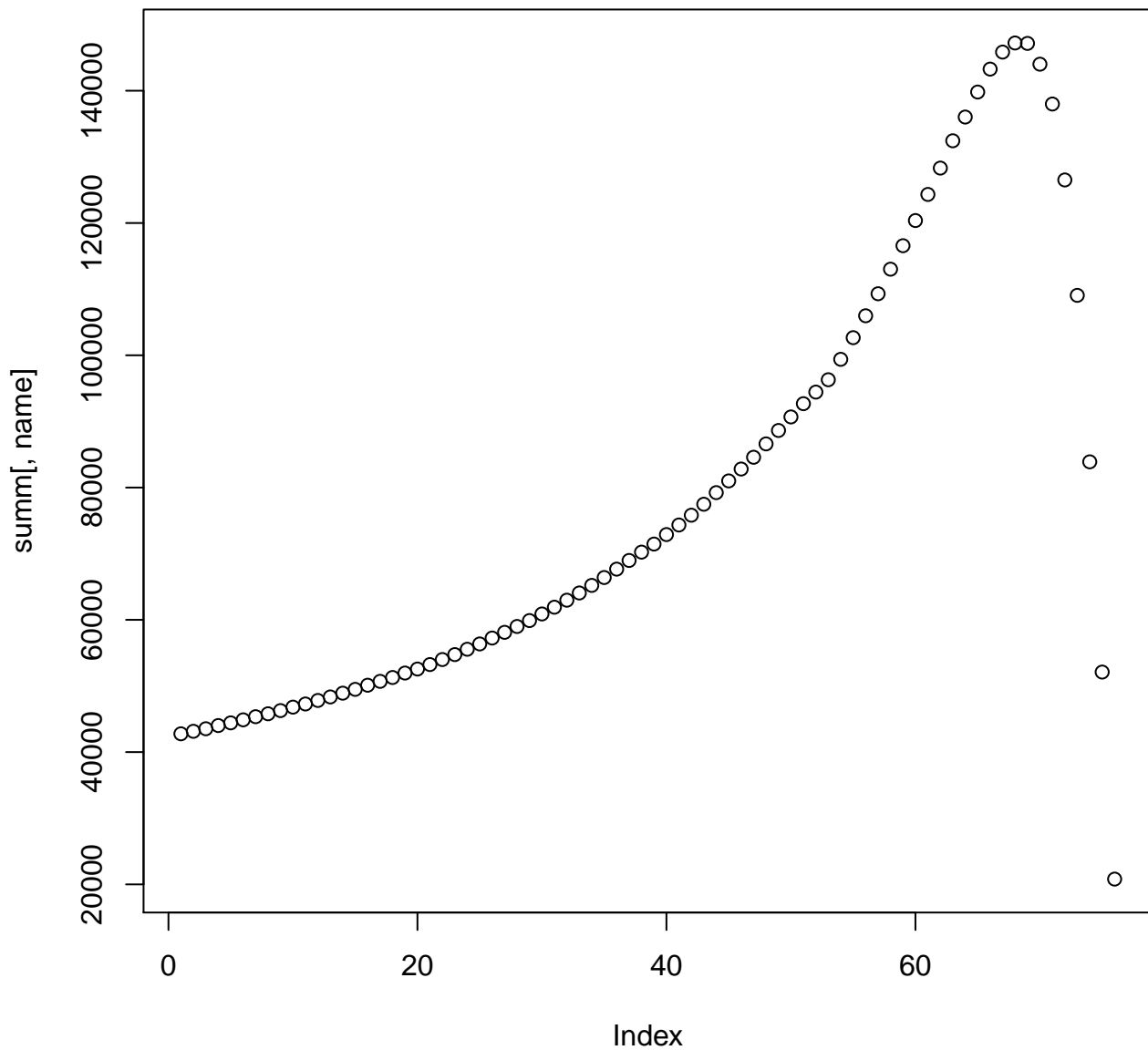
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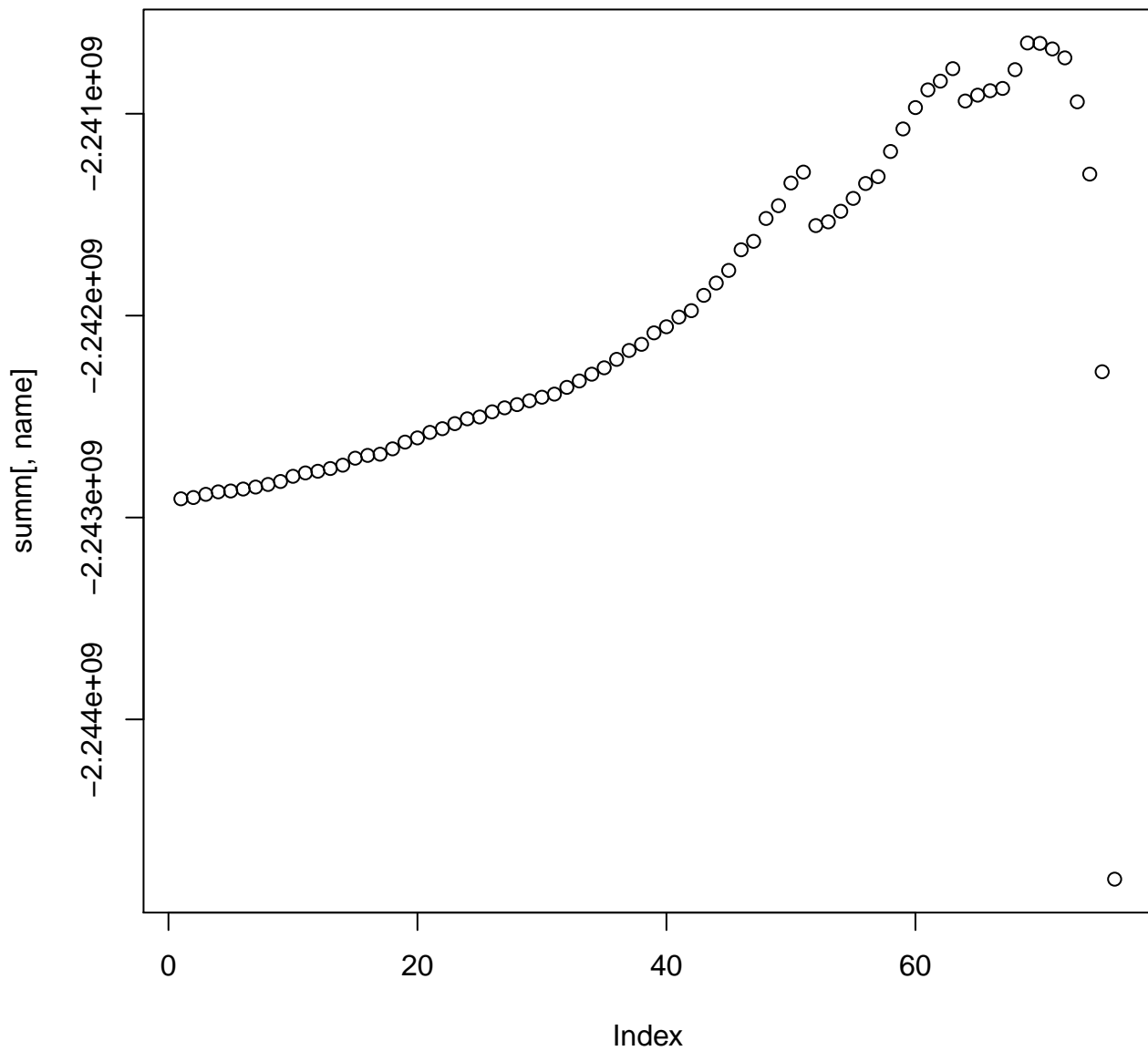
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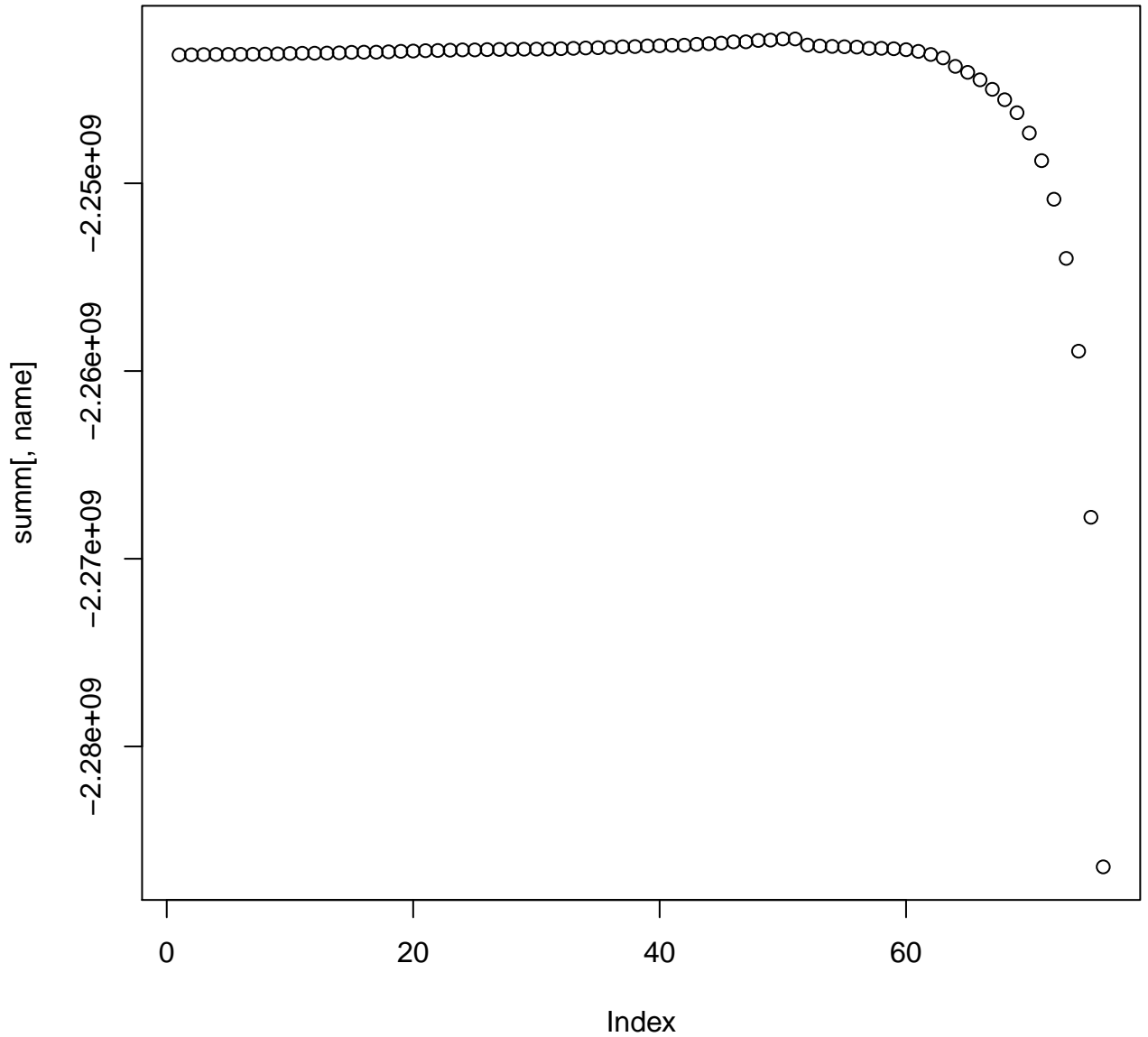
assembly.num.contigs.longer.than.100.bp



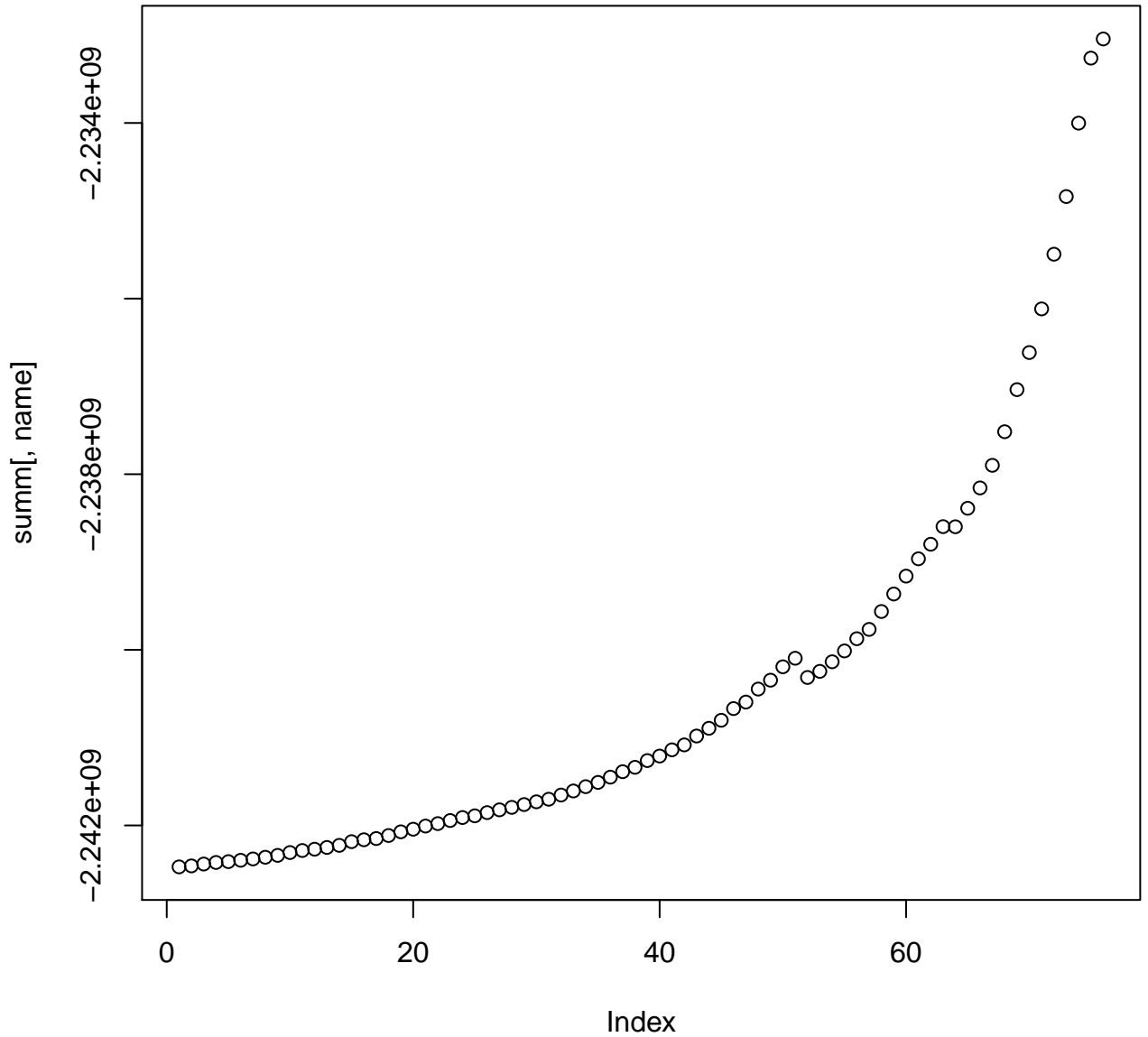
rsem.approx.approx



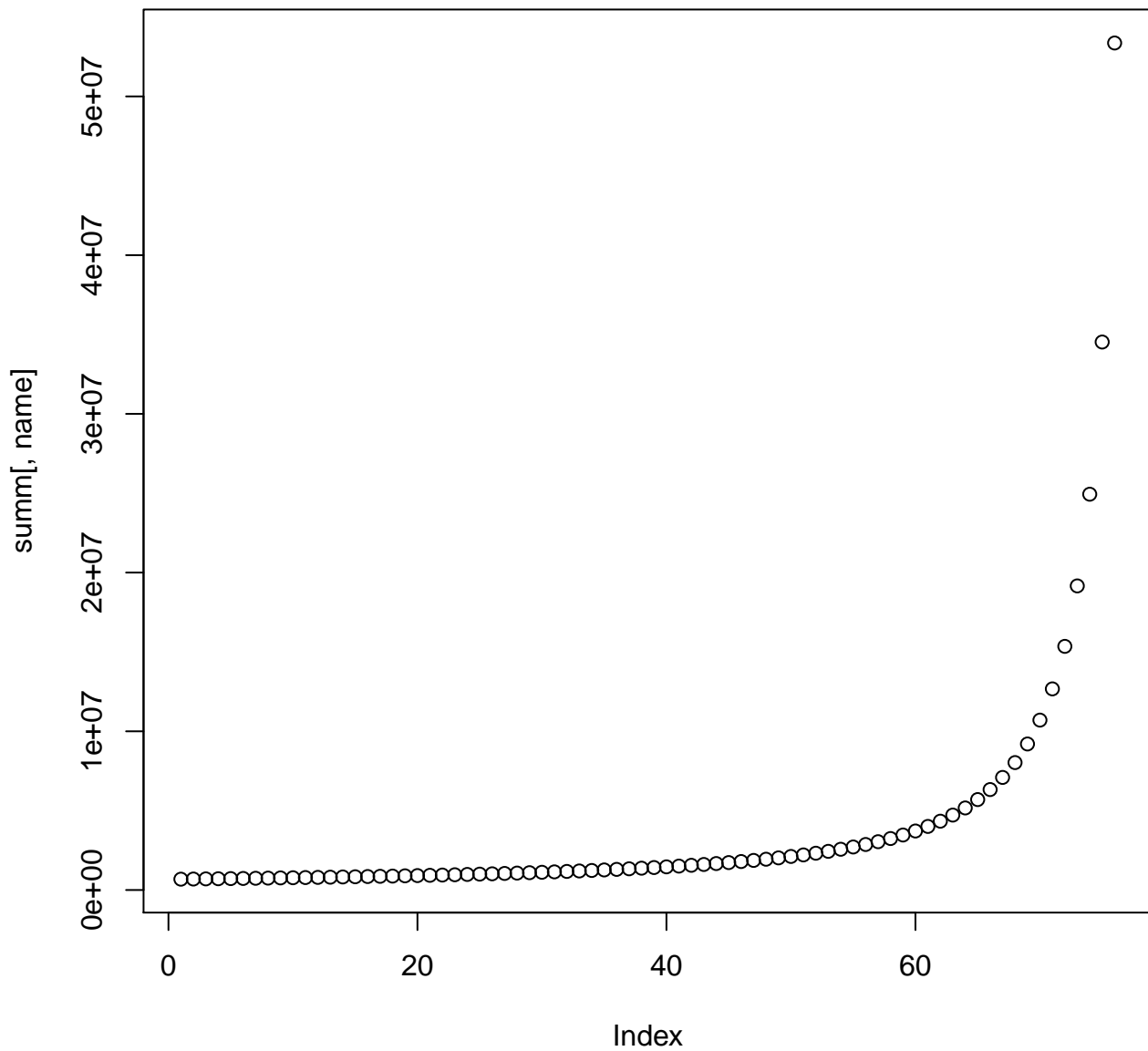
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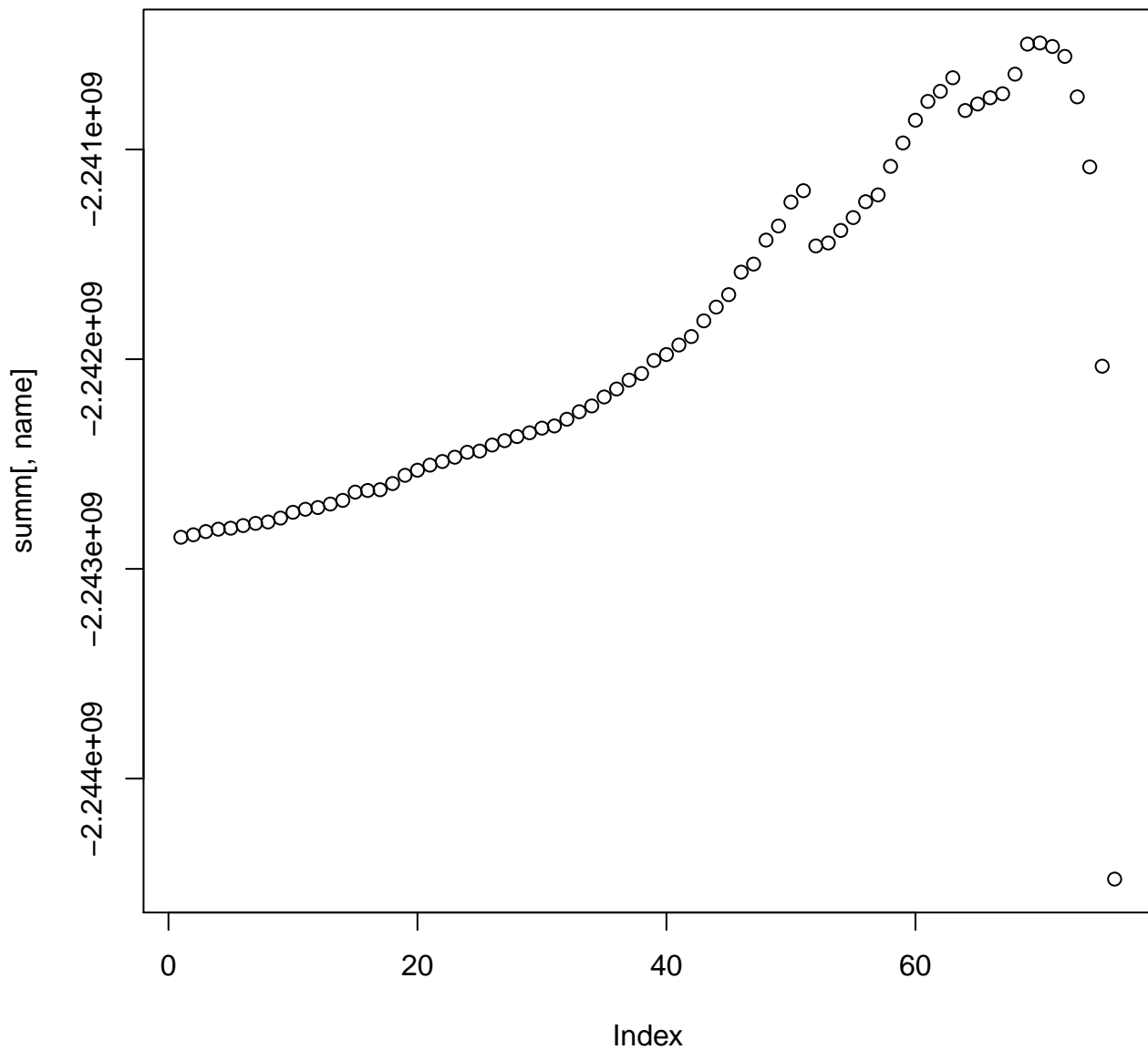
rsem.approx.loglikelihood



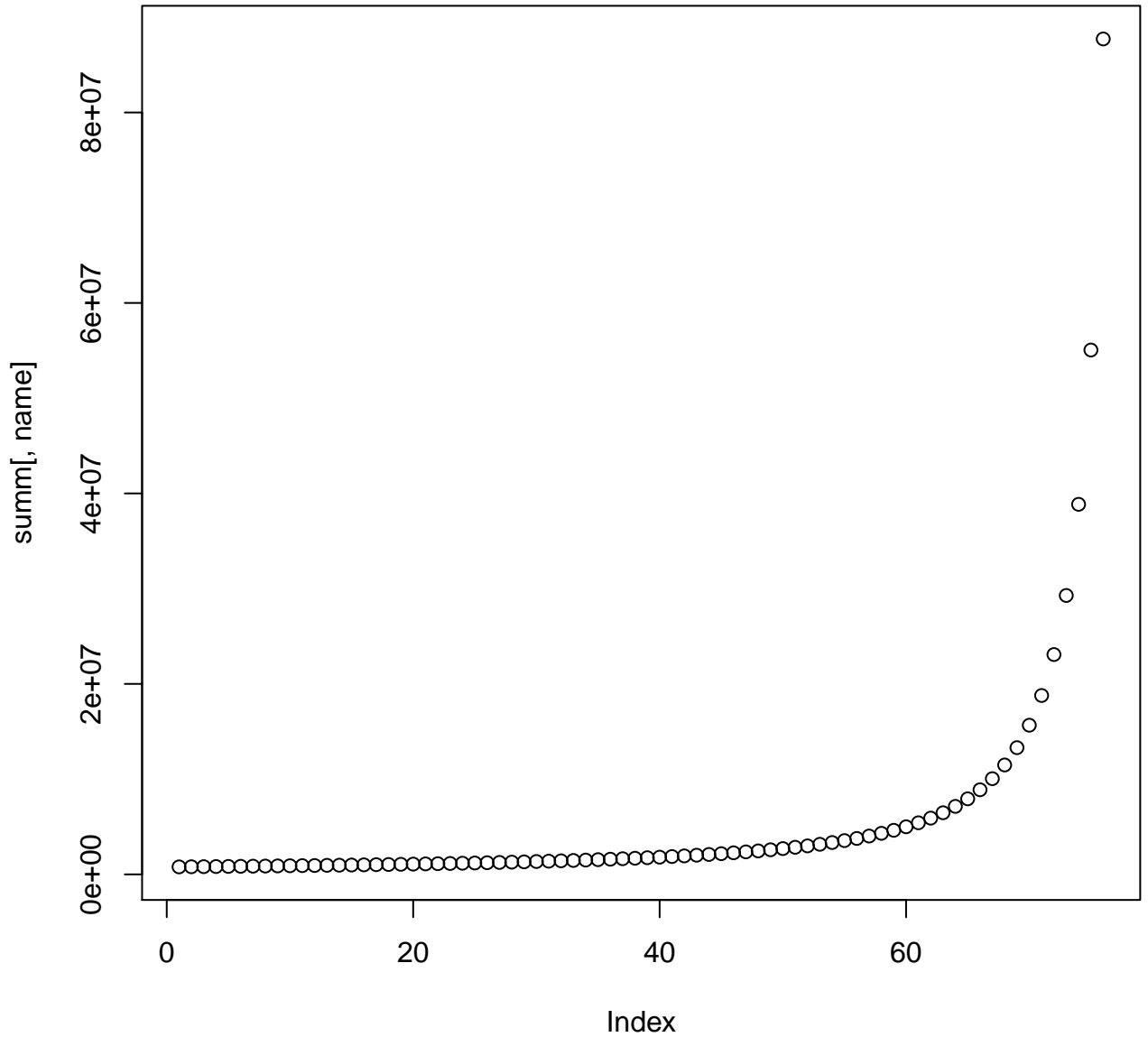
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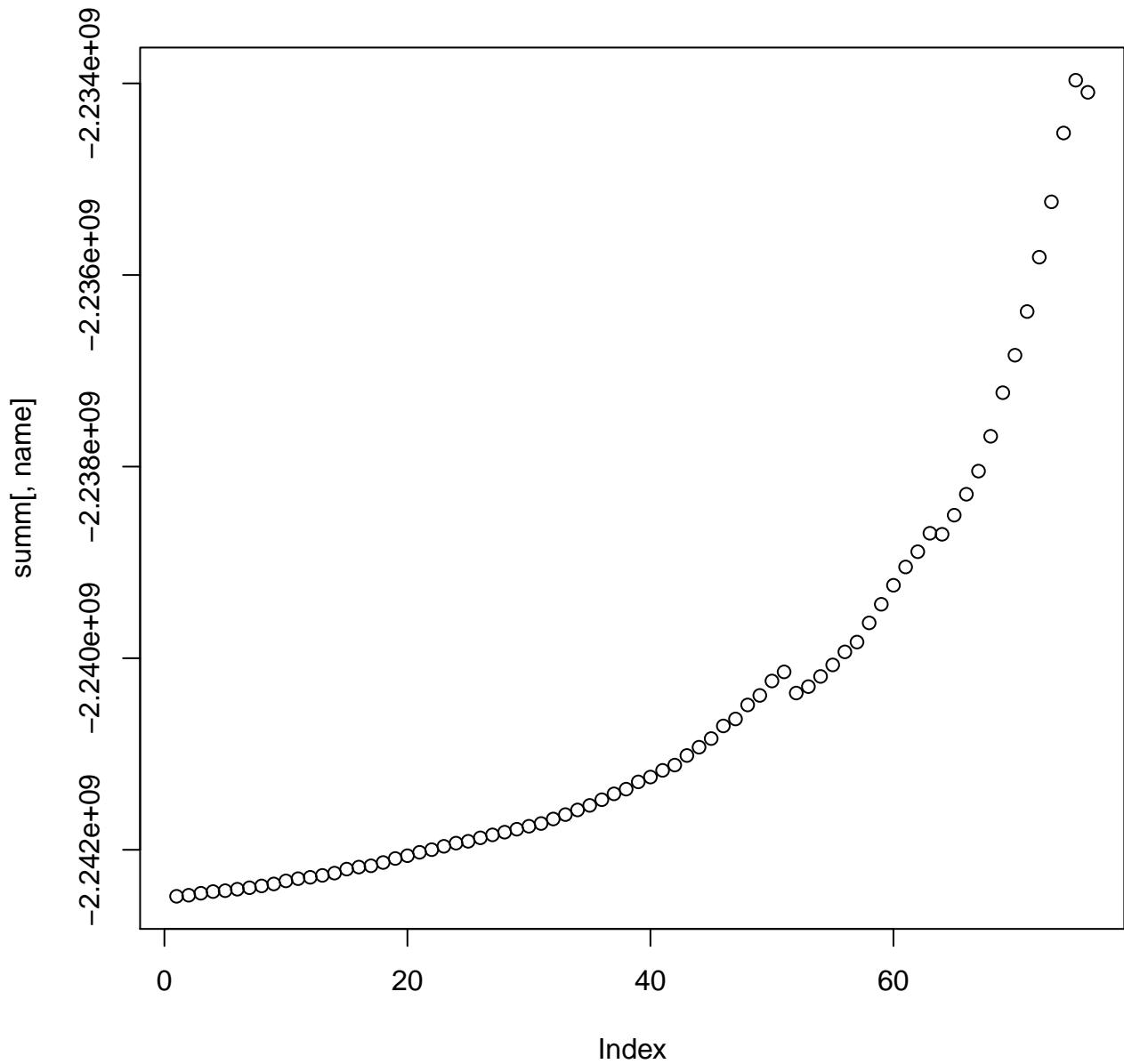
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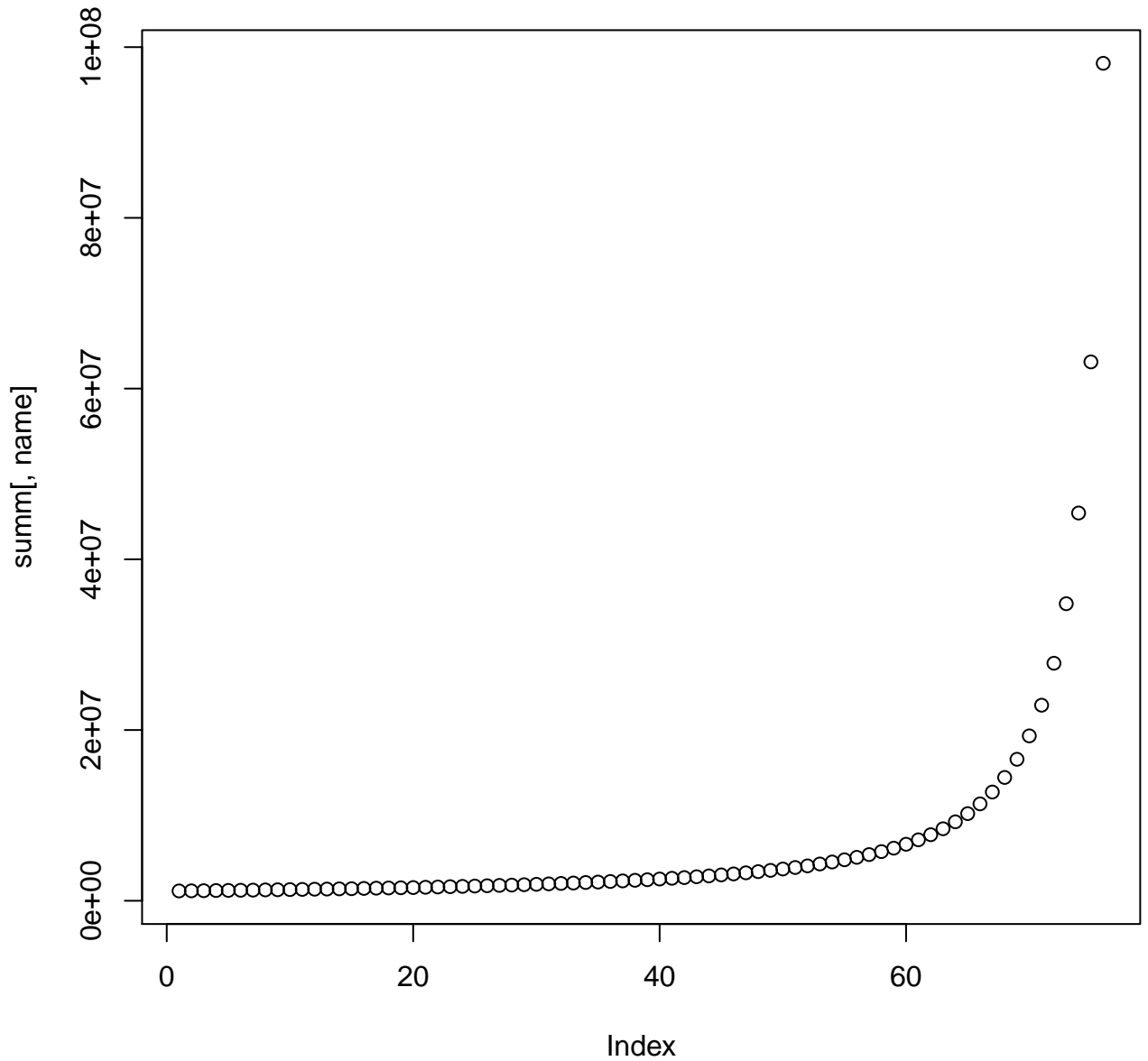
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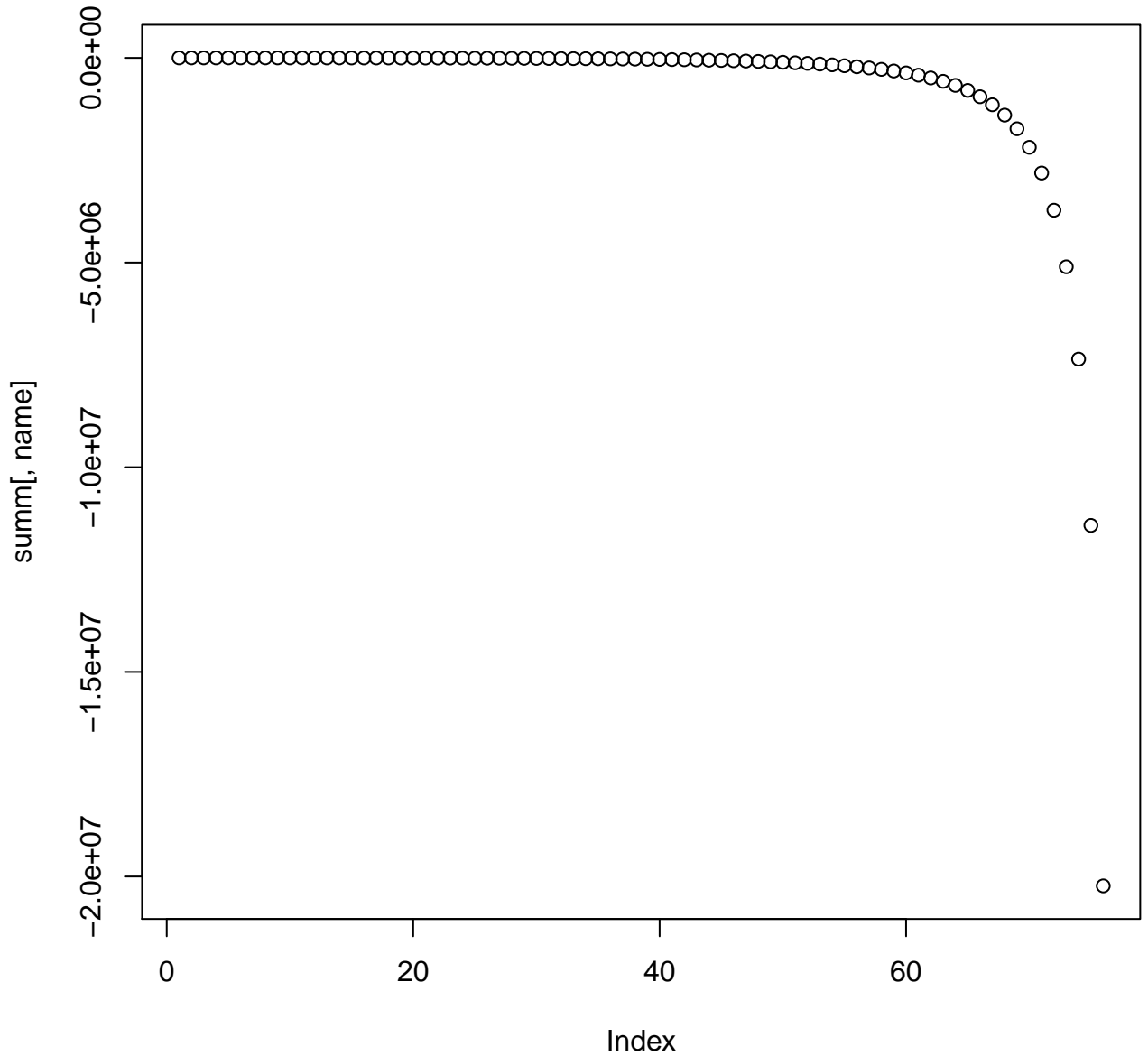
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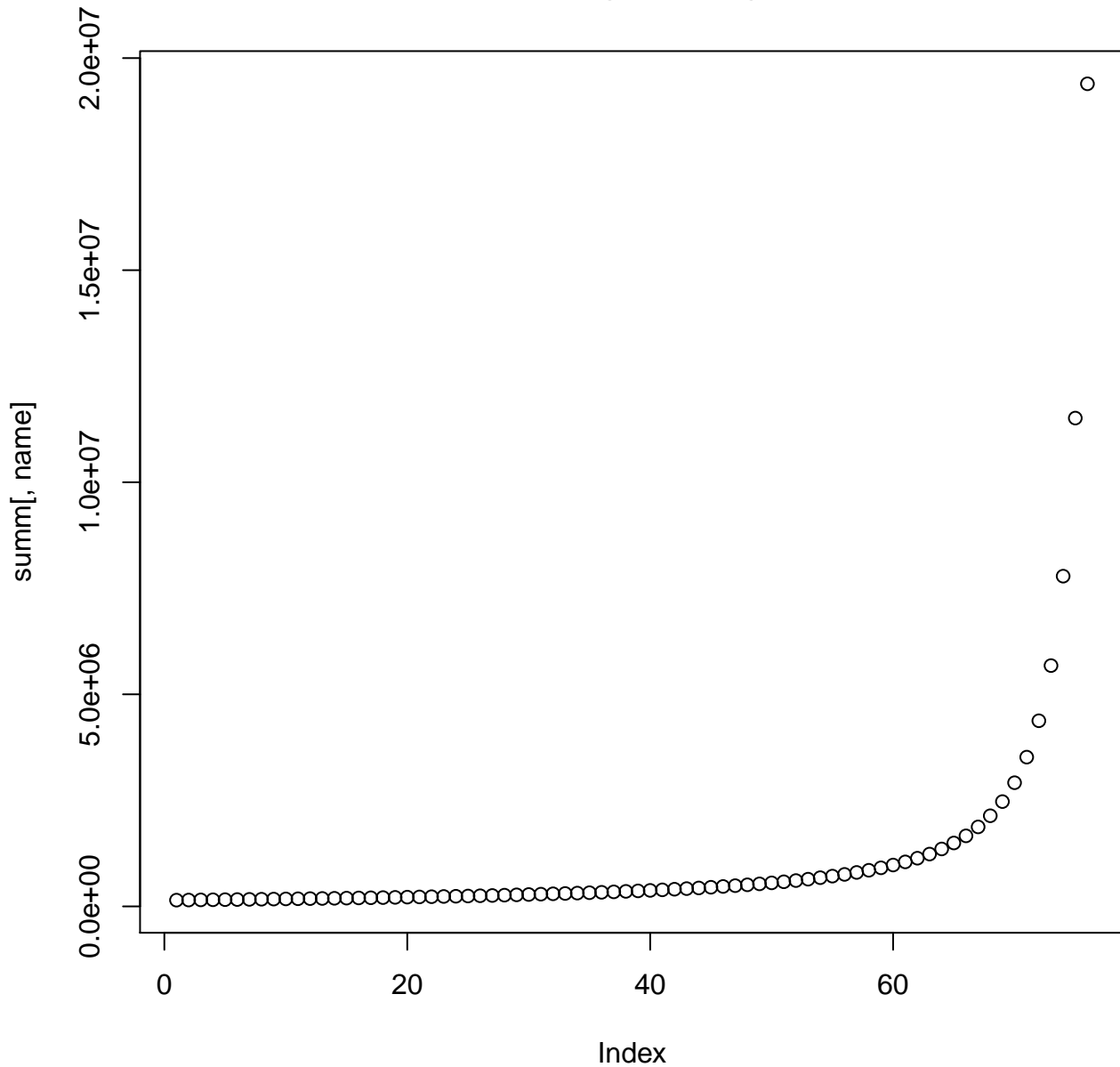
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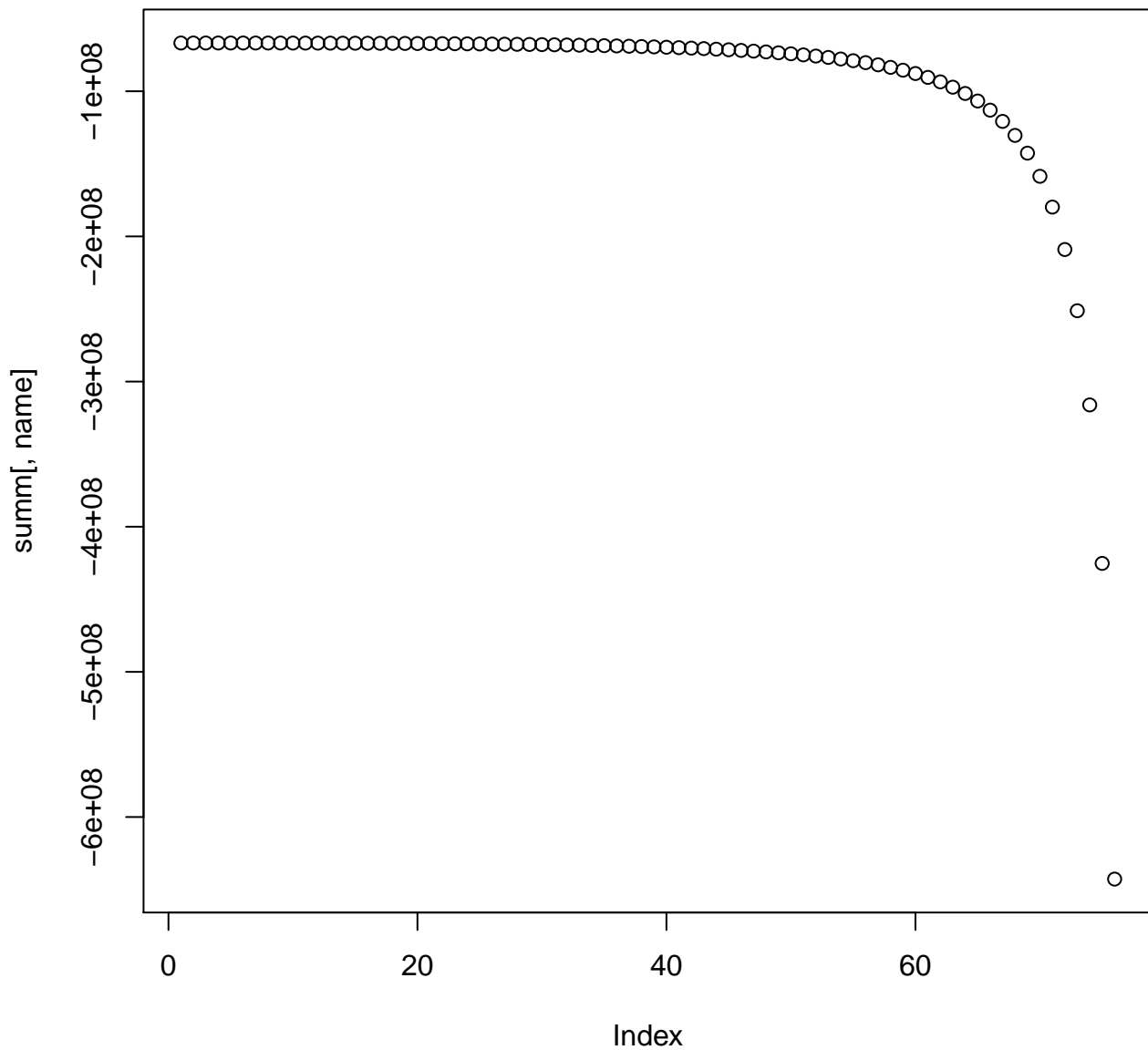
rsem.prior.log.prob.M



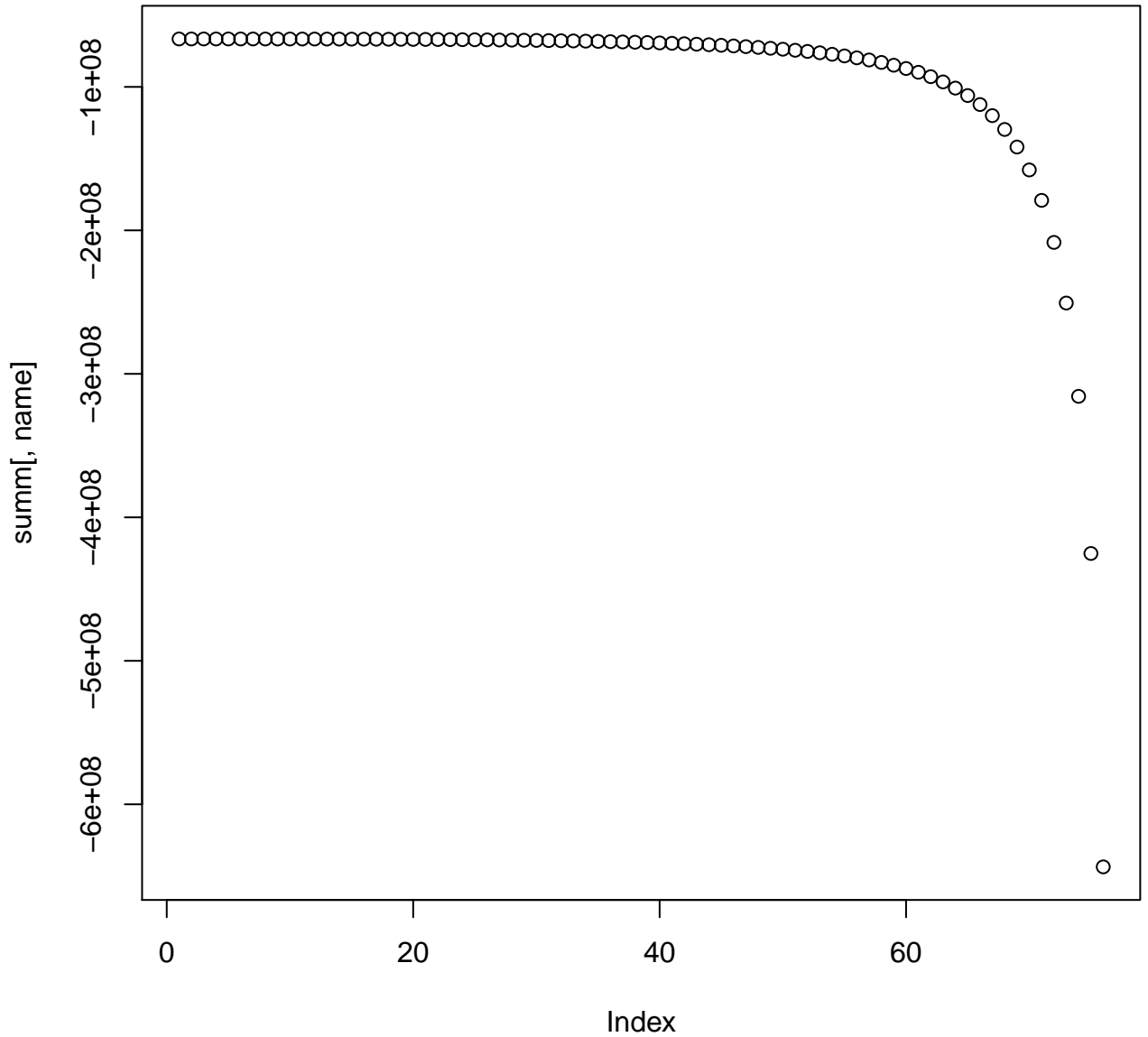
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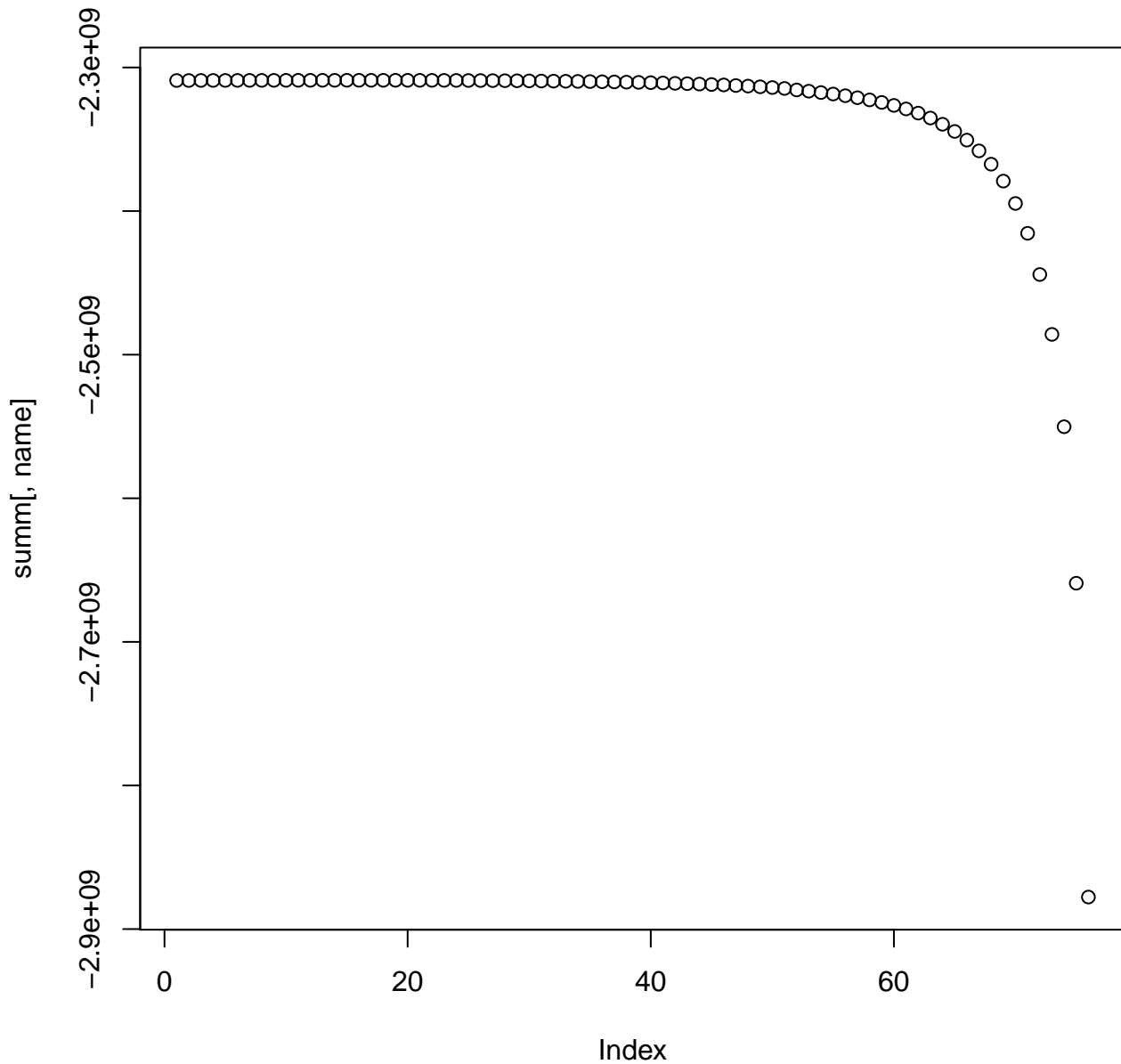
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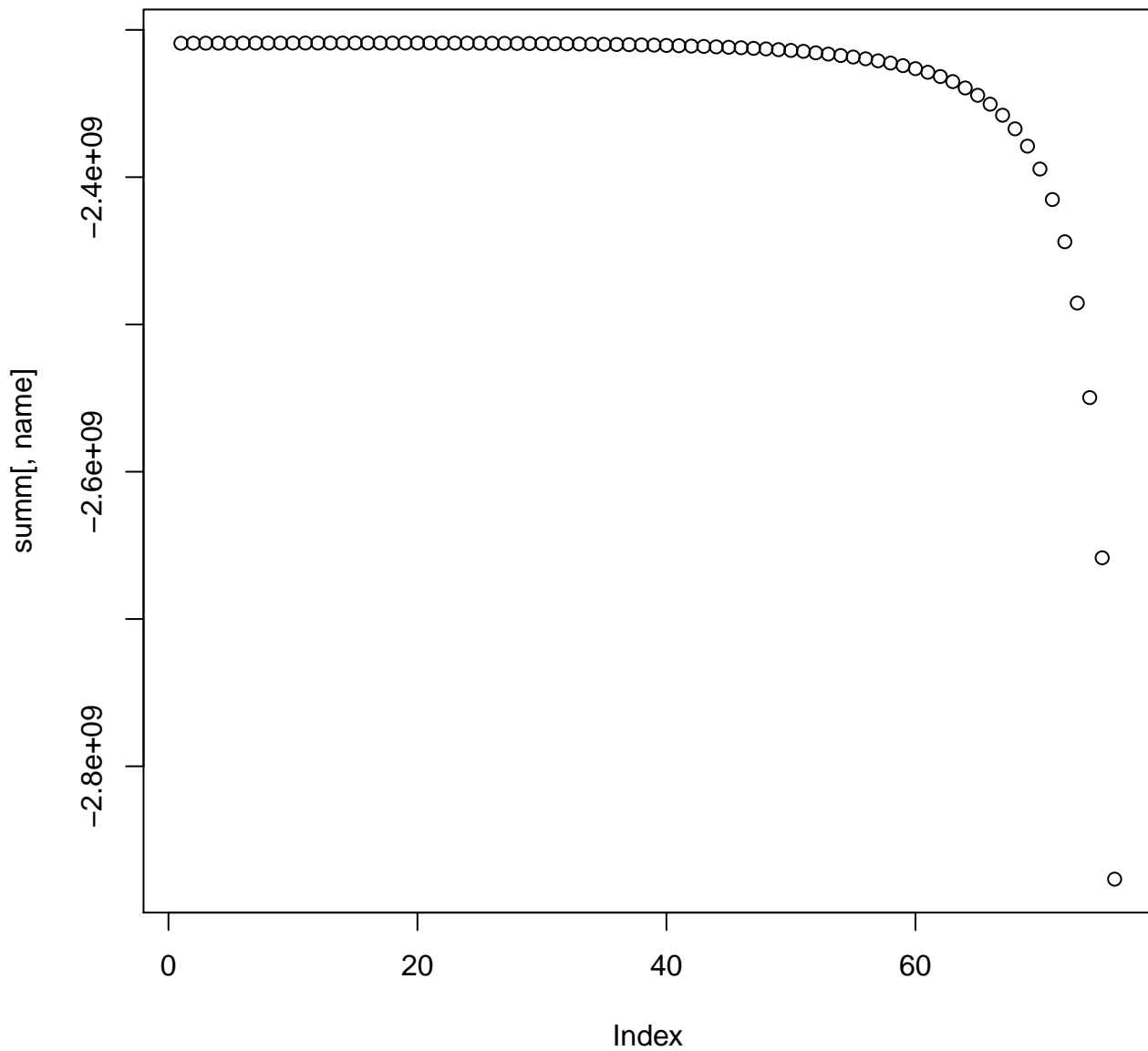
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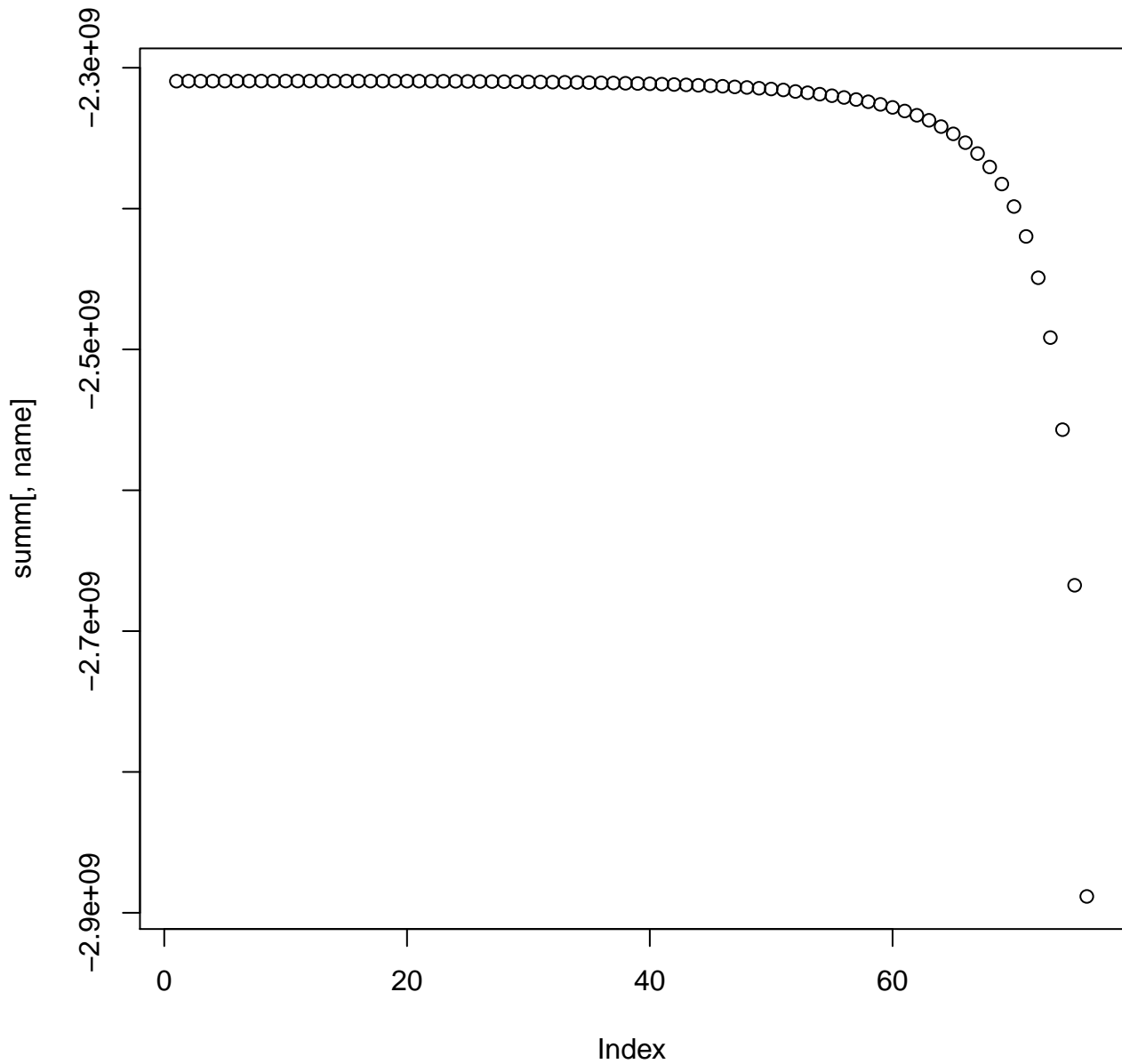
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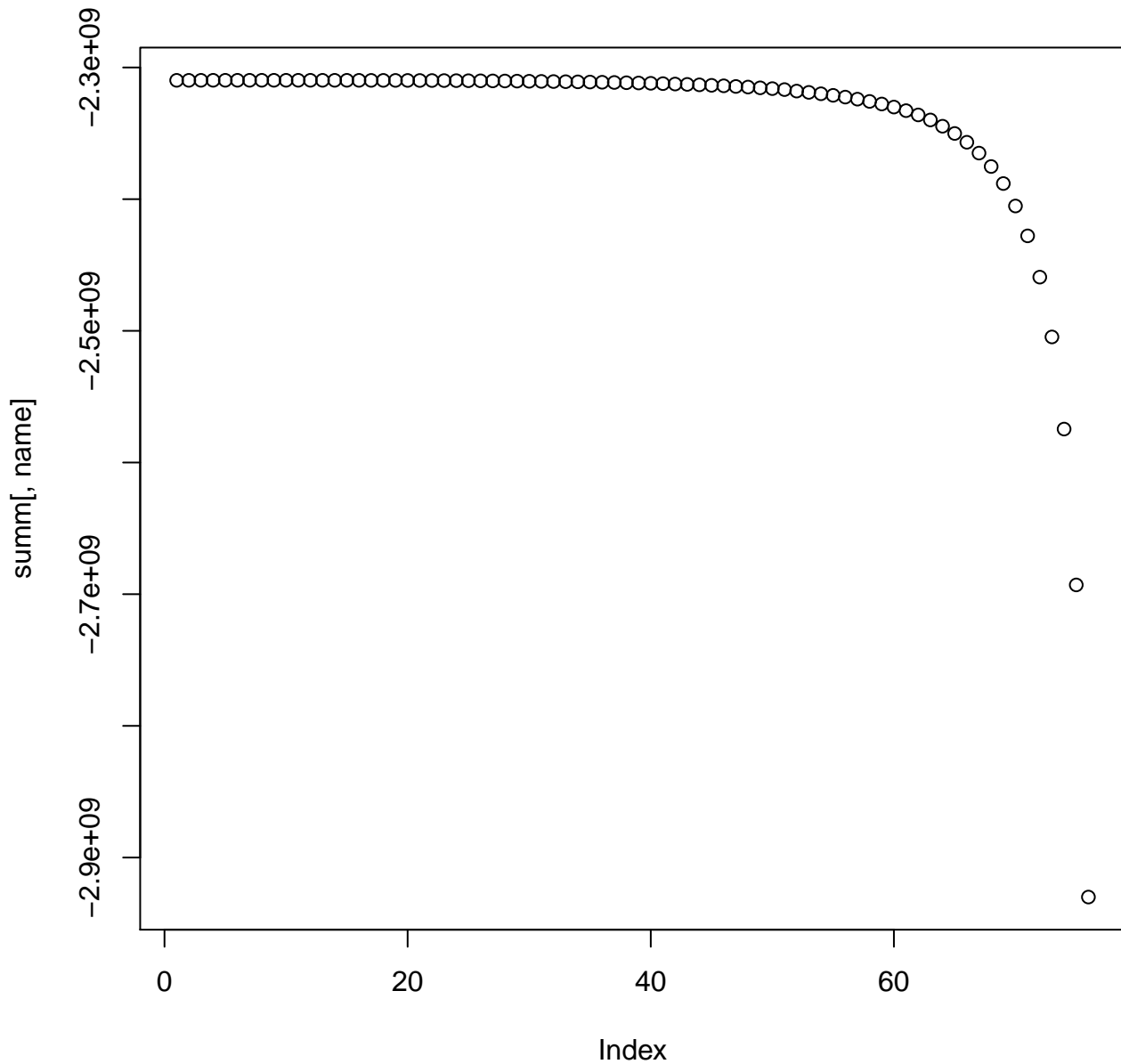
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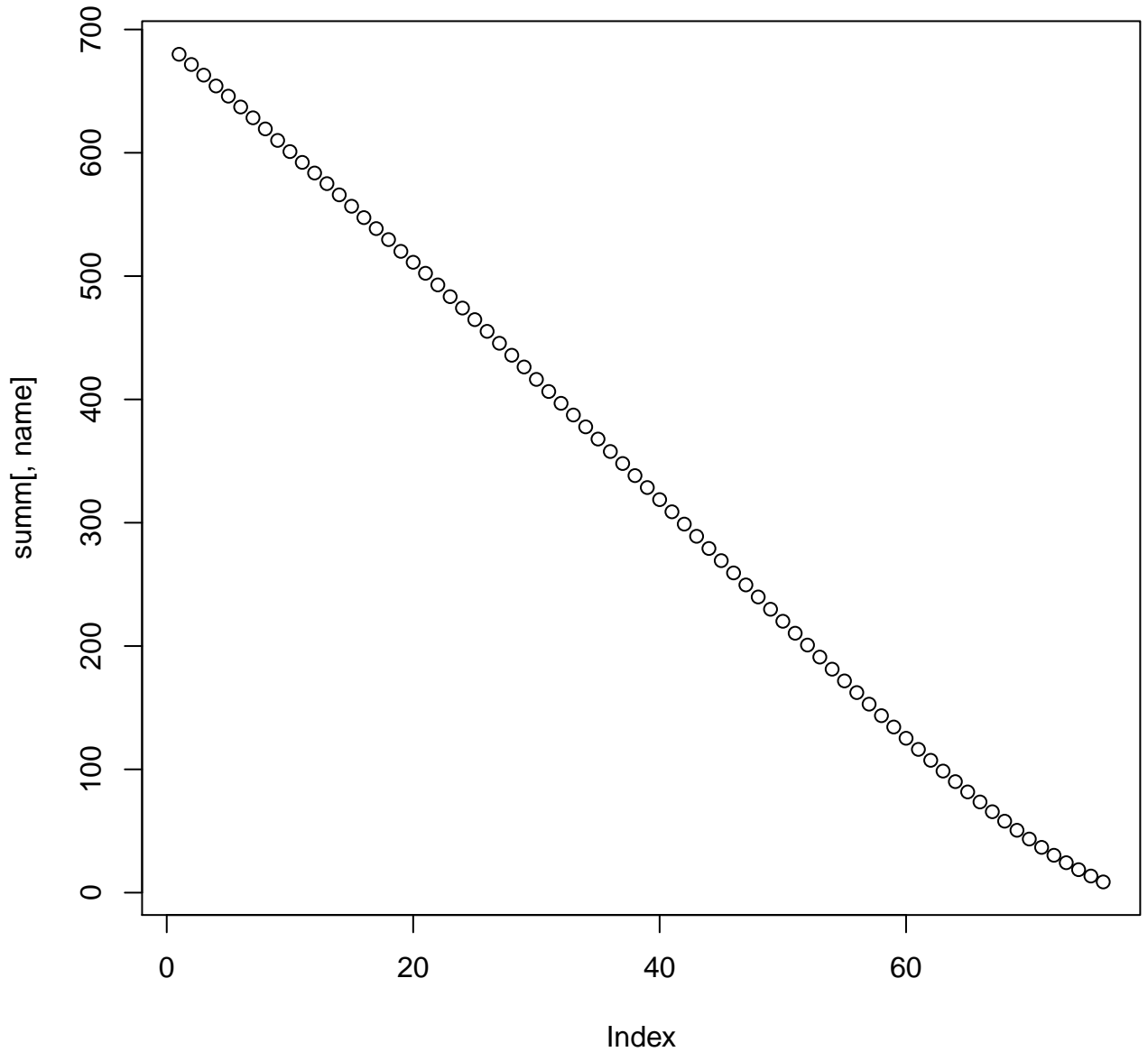
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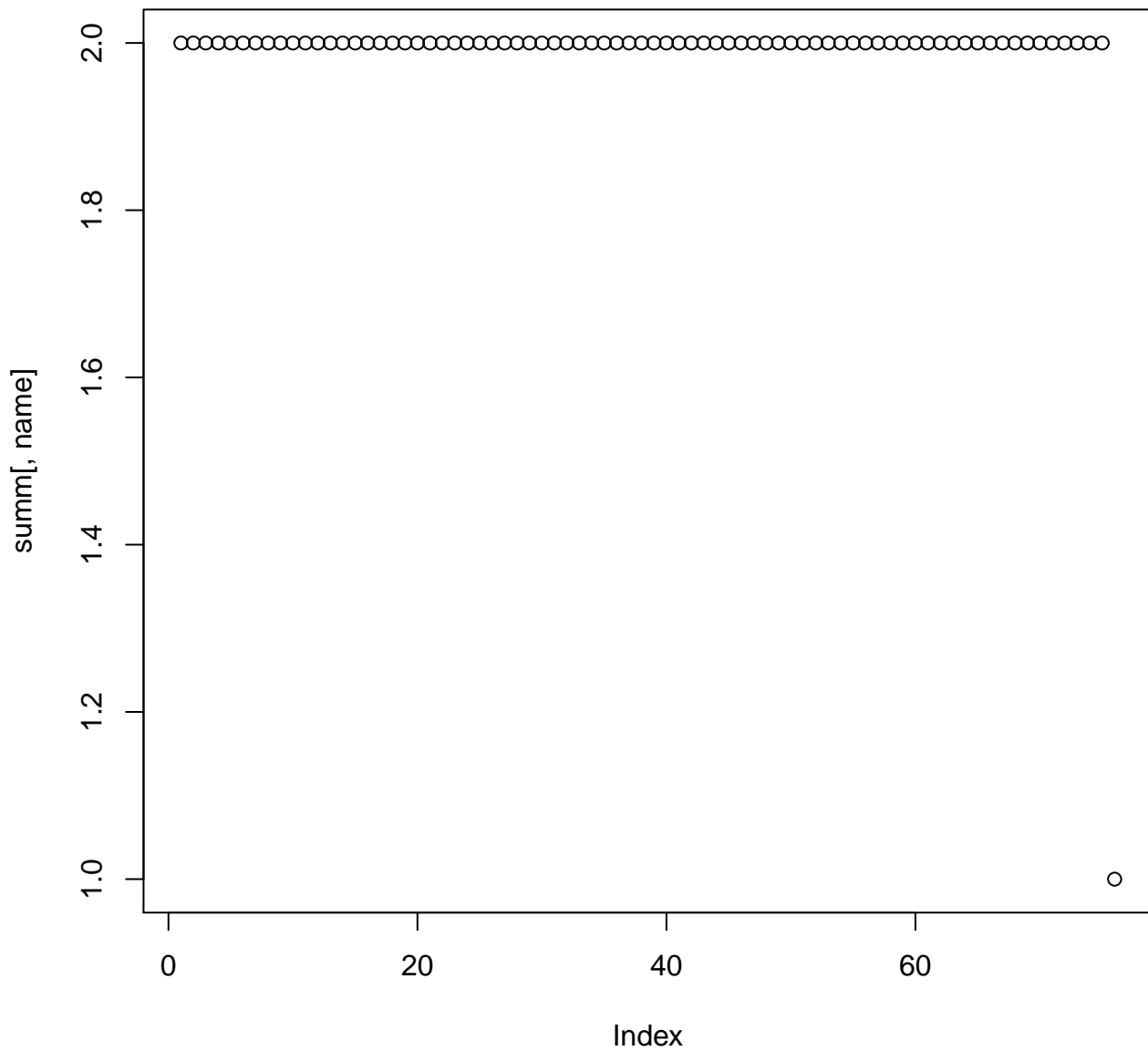
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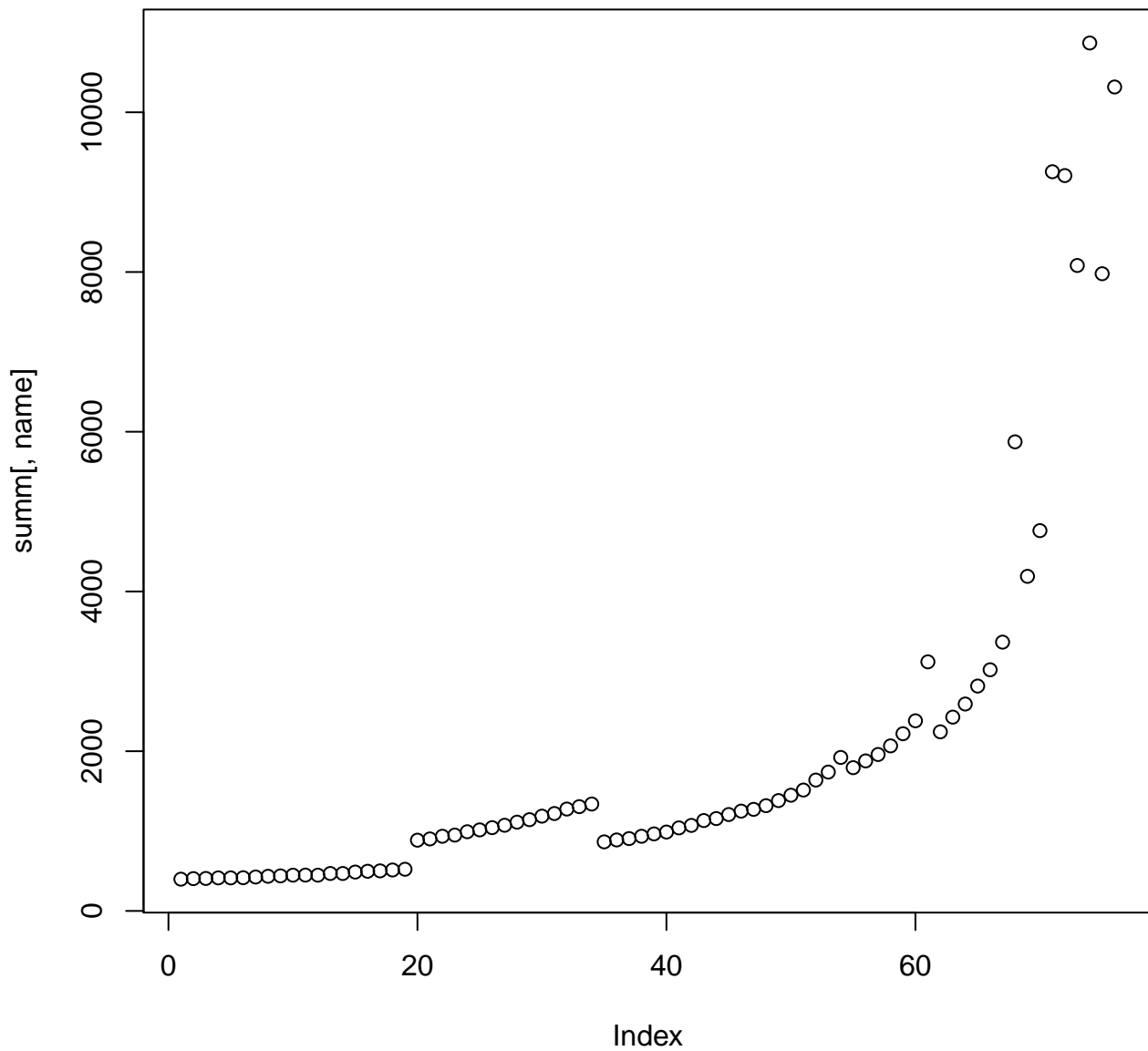
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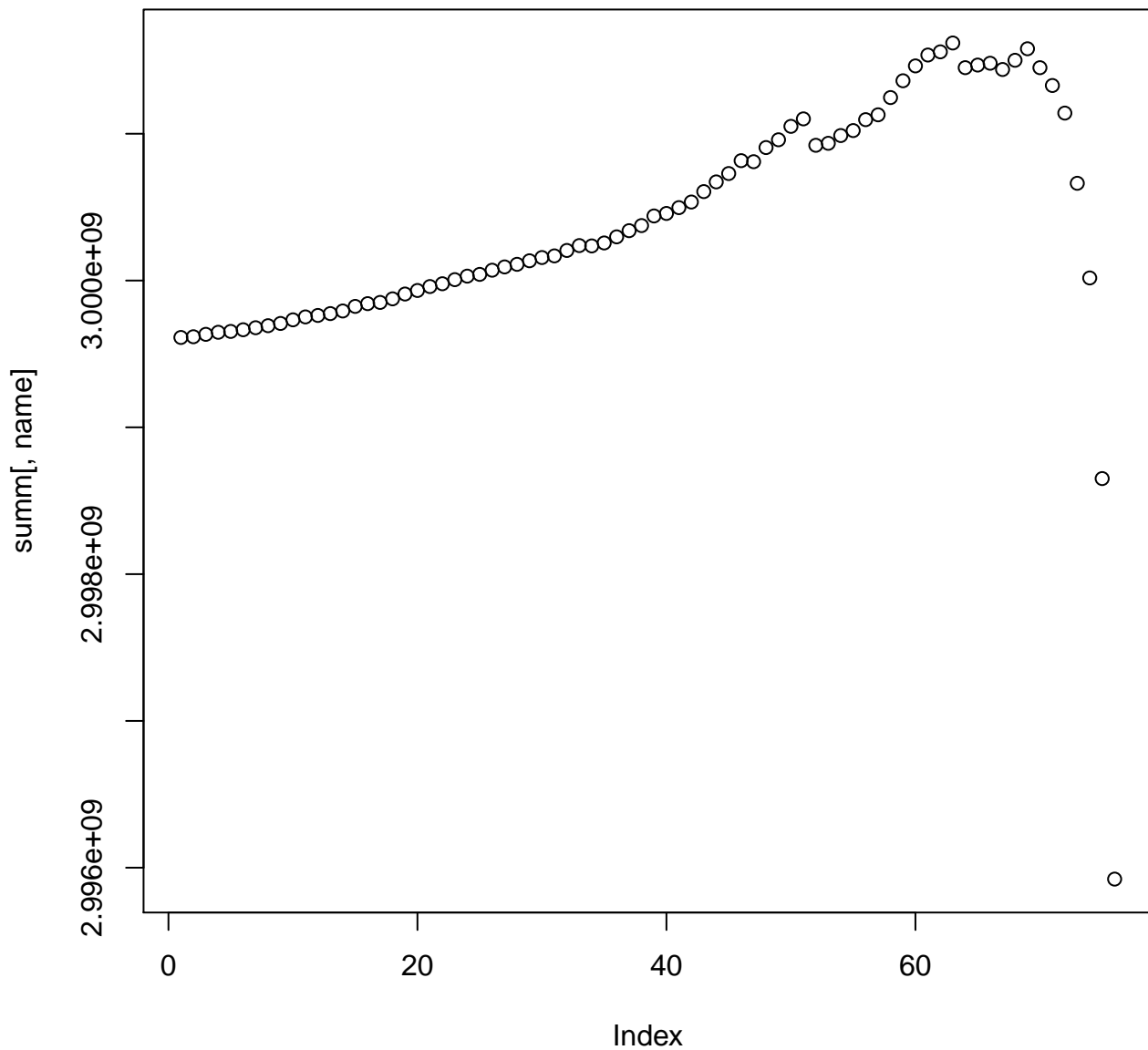
rsem.ss.median.num.reads.per.transcript



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



rsem.ss.num.matching.bases



rsem.ss.num.mismatching.bases

