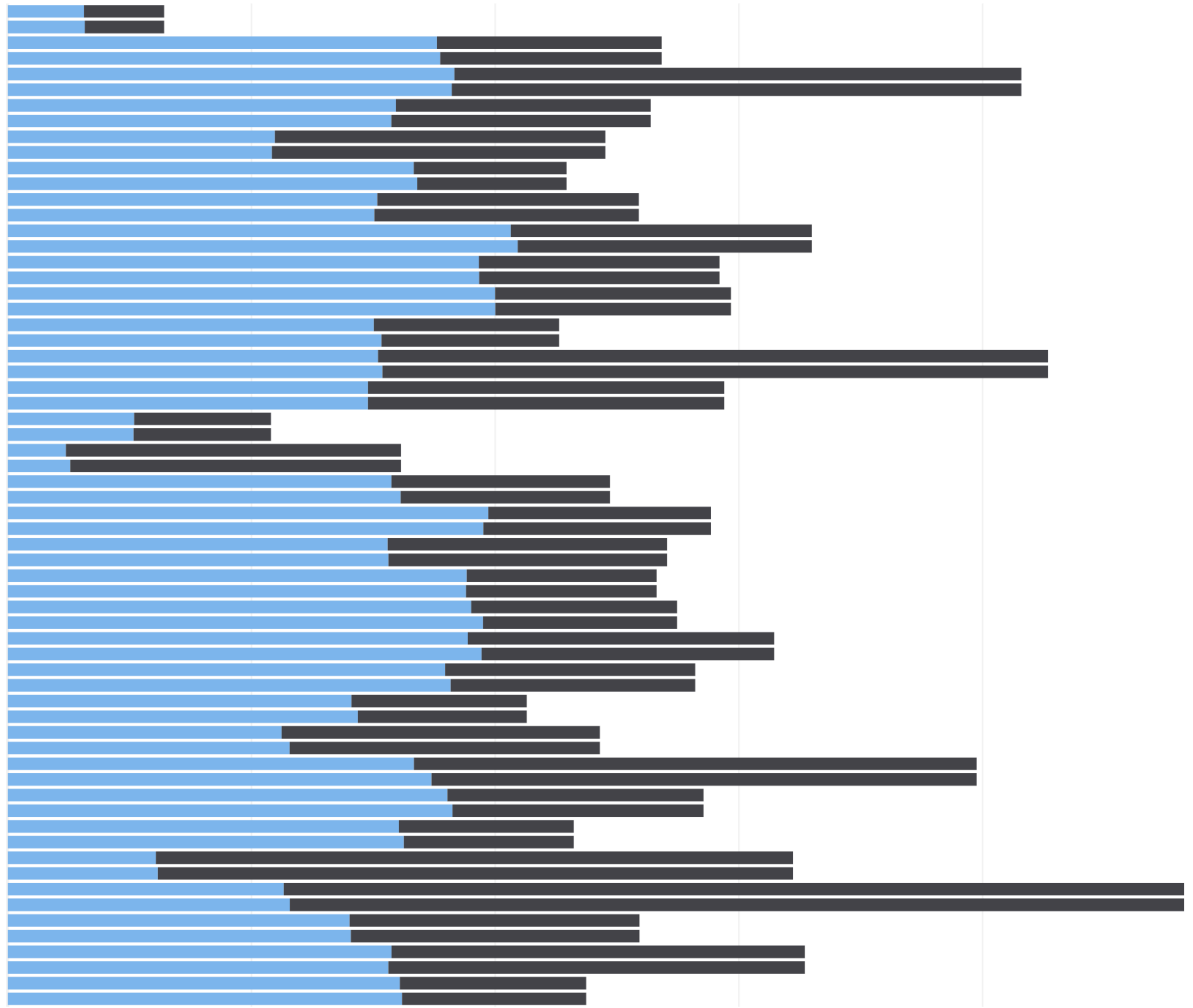


FastQC: Sequence Counts

64 samples

Unique Reads
Duplicate Reads

CF01-CM01-Zygote_2
CF01-CM02-Larvae_2
CF02-CM02-Zygote_2
CF03-CM03-Zygote_2
CF03-CM04-Larvae_2
CF03-CM05-Larvae_2
CF04-CM04-Zygote_2
CF05-CM02-Larvae_2
CF05-CM05-Zygote_2
CF06-CM01-Zygote_2
CF06-CM02-Larvae_2
CF07-CM02-Zygote_2
CF08-CM03-Zygote_2
CF08-CM04-Larvae_2
CF08-CM05-Larvae_2
EF01-EM01-Zygote_2
EF02-EM02-Zygote_2
EF03-EM03-Zygote_2
EF03-EM04-Larvae_2
EF03-EM05-Larvae_2
EF04-EM04-Zygote_2
EF04-EM05-Larvae_2
EF05-EM01-Larvae_2
EF05-EM05-Zygote_2
EF05-EM06-Larvae_2
EF06-EM01-Larvae_2
EF06-EM02-Larvae_2
EF06-EM06-Larvae_2
EF07-EM01-Zygote_2
EF07-EM03-Larvae_2
EF08-EM03-Larvae_2
EF08-EM04-Larvae_2



0 50M 100M 150M 200M

Number of reads