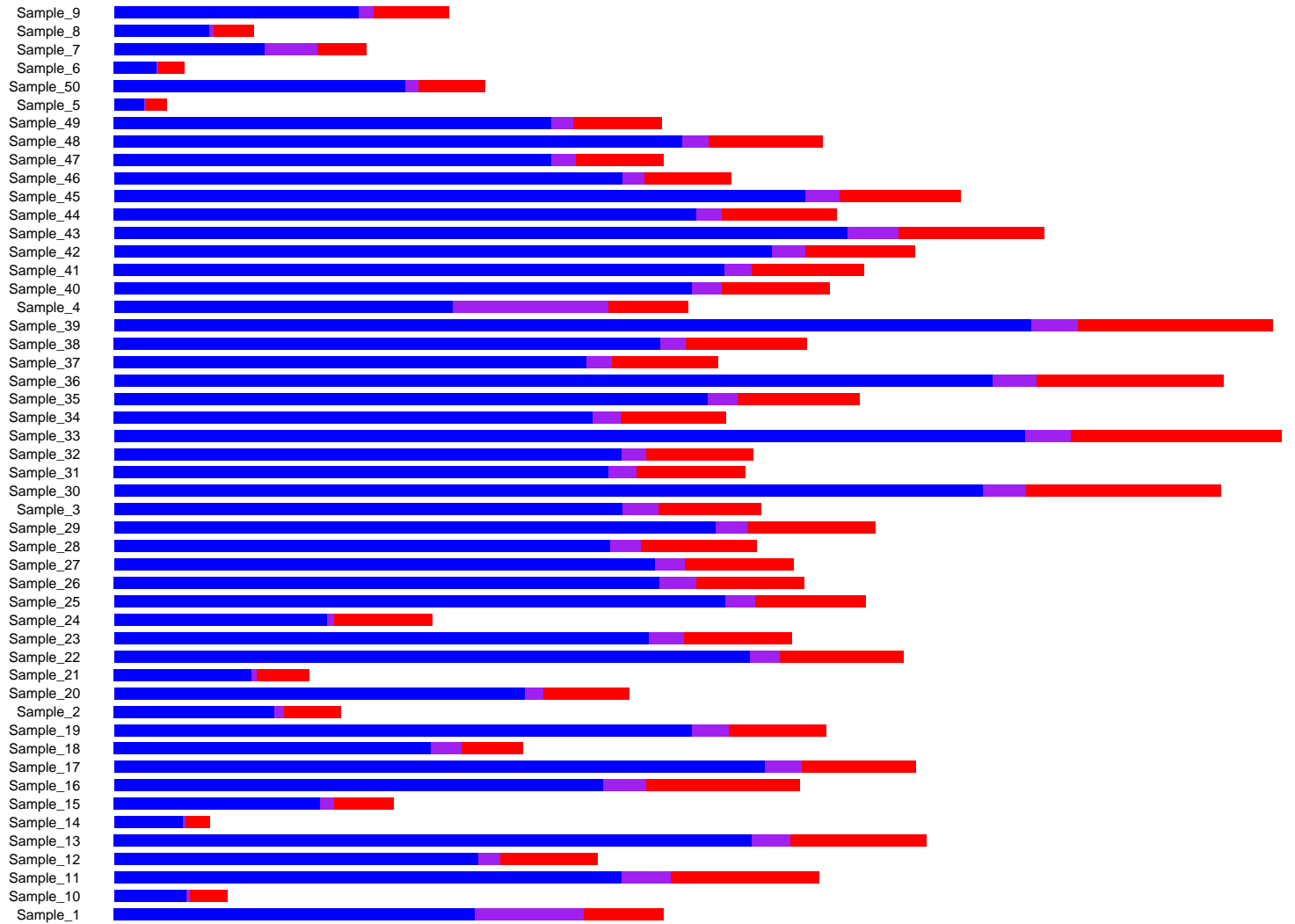


Sequencing Reads

■ Deduplicated ■ Duplicated ■ Unaligned



0 10 20 30 40 50

Number of reads (millions)