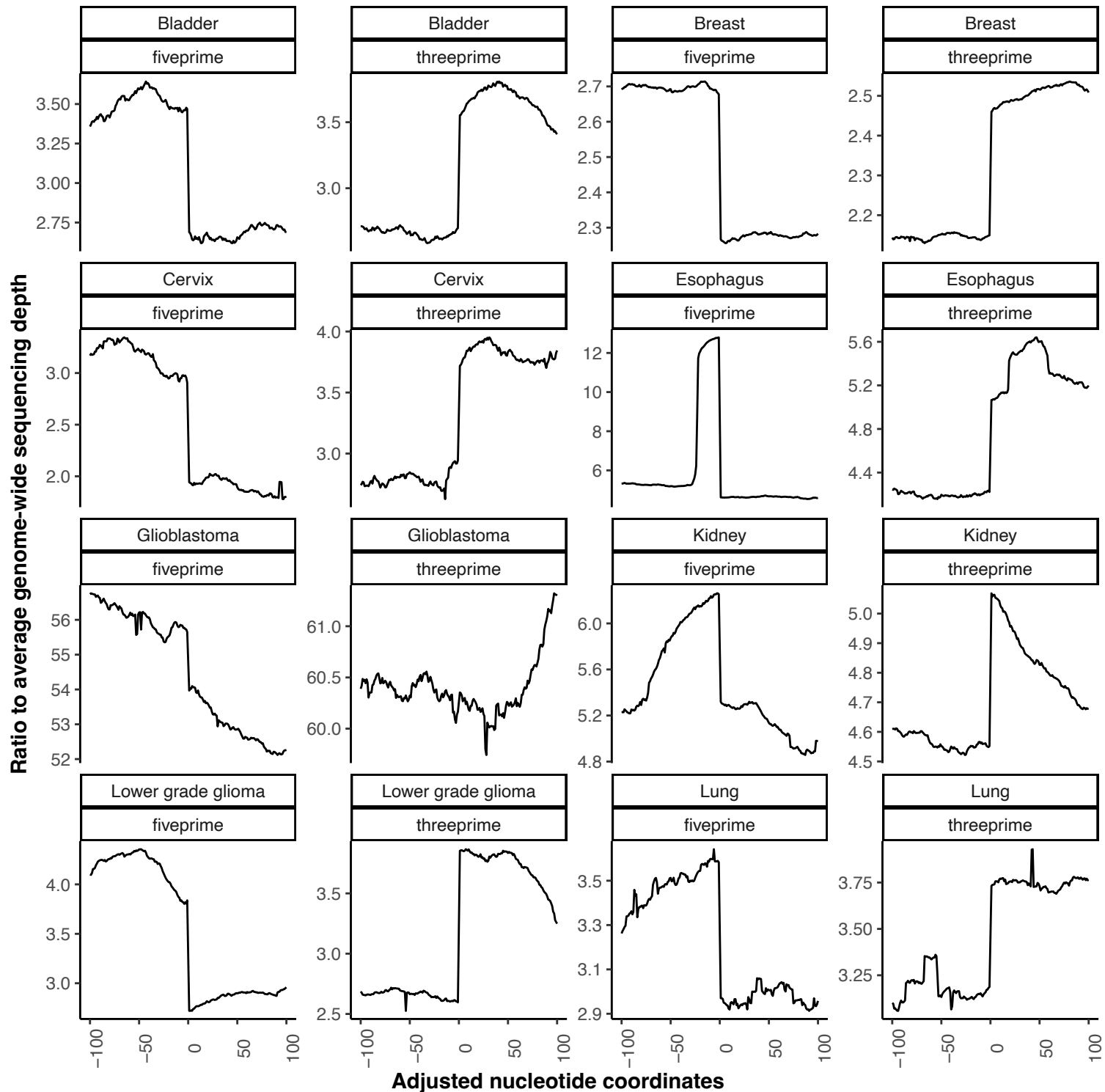




Additional file 6. Sequencing depth around genomic breakpoints in WGS data, expressed as a ratio to the genome average.