

fibroblast – smartseq2 Trachea # detected genes 2000-% MT 30 20 10 3e+06 1e+05 3e+05 1e+06 Library size (log scale) 1.5density 0.0-1e+05 3e+05 1e+06 3e+06 Library size (log scale) density 0.10-0.05-0.00-20 10 30 % Mitochondrial features