mature enterocyte – smartseq2 Small_Intestine 5000-# detected genes 3000-2000-1000-% MT 75 50 25 1e+05 1e+06 1e+04 Library size (log scale) density 0.6-0.3-0.0 1e+05 1e+06 1e+04 Library size (log scale) 0.020density 0.015-0.005-0.000-50 100 % Mitochondrial features