## STAR: Summary Statistics 32 samples Total reads 0 M 5 M 10 M 15 M 20 M 25 M 30 M Aligned 50% 70% 0% 10% 30% 40% 60% 20% Uniq aligned 40% 0% 10% 20% 30% 50% 60% Avg. mapped len 0bp 160bp 180bp 20bp 40bp 60bp 80bp 100bp 120bp 140bp Annotated splices 5 M 0 M 10 M 15 M 20 M 25 M 30 M Mismatch rate 0.6% Del rate = 0.02% 0.03% 0.04% 0.05% 0.08% 0.09% 0.01% 0.06% 0.07% 0% Del len 2bp 1bp 1.5bp 2.5bp 0.5bp 0bp Ins rate 0.05% 0.1% 0.15% 0.2% 0.25% Ins len 2bp 0.5bp 1bp 1.5bp 0bp