STAR: Summary Statistics 38 samples Total reads 0 M 5 M 10 M 15 M 20 M 25 M 30 M Aligned 0% 10% 40% 50% 60% 20% 30% 70% Uniq aligned 30% 0% 10% 20% 40% 50% 60% Avg. mapped len 0bp 140bp 160bp 180bp 20bp 40bp 60bp 80bp 100bp 120bp Annotated splices 5 M 0 M 10 M 15 M 20 M 25 M 30 M Mismatch rate Del rate 0.04% 0.06% 0.08% 0.1% 0.02% 0% Del len 0.5bp 1.5bp 2bp 2.5bp 3bp 0bp 1bp Ins rate 0.05% 0.1% 0.3% 0.15% 0.2% 0.25% Ins len 1.5bp 1bp 2bp 0bp 0.5bp