STAR: Summary Statistics 25 samples Total reads 0 M 30 M 40 M 10 M 20 M 50 M Aligned 0% 10% 20% 30% 40% 50% 60% 70% 80% 90% Uniq aligned 0% 10% 20% 30% 40% 50% 60% 70% 80% Avg. mapped len 0bp 250bp 50bp 100bp 150bp 200bp Annotated splices 0 M 10 M 20 M 40 M 30 M 50 M Mismatch rate Del rate 0.02% 0.03% 0.05% 0.01% 0.04% Del len 1bp 1.5bp 2bp 2.5bp 0bp 0.5bp Ins rate 0.02% 0.005% 0.01% 0.015% 0.025% 0.03% Ins len 2bp 0.5bp 1.5bp 2.5bp 0bp 1bp