STAR: Summary Statistics 38 samples Total reads 0 M 5 M 10 M 15 M 20 M 25 M 30 M Aligned 40% 20% 50% 60% 0% 10% 30% 70% Uniq aligned 40% 0% 10% 20% 30% 50% 60% Avg. mapped len 0bp 120bp 160bp 180bp 20bp 40bp 60bp 80bp 100bp 140bp Annotated splices 5 M 0 M 10 M 15 M 20 M 25 M 30 M Mismatch rate 0.8% Del rate 0.06% 0.1% 0.18% 0.02% 0.04% 0.08% 0.12% 0.14% 0.16% 0% Del len 1bp 3bp 4bp 0bp 2bp 5bp Ins rate 0.05% 0.1% 0.15% 0.2% 0.4% 0.25% 0.3% 0.35% Ins len 1.5bp 1bp 3bp 3.5bp 0bp 0.5bp 2bp 2.5bp