## STAR: Alignment Scores 30 samples 24L007779\_S1\_L001 24L007779\_S1\_L002 24L007780\_S2\_L001 24L007780\_S2\_L002 24L007781\_S3\_L001 24L007781\_S3\_L002 24L007782\_S4\_L001 24L007782\_S4\_L002 24L007783\_S5\_L001 24L007783\_S5\_L002 24L007784\_S6\_L001 24L007784\_S6\_L002 2CLCmesc\_0-0-3\_S15 2CLCmesc\_0-0-4\_S16 2CLCmesc\_003\_RR\_S3 2CLCmesc\_004\_RR\_S4 2CLCmesc\_011\_RR\_S18 2CLCmesc\_016\_RR\_S17 2CLCmesc\_1\_RR\_S9 2CLCmesc\_1\_S1 2CLCmesc\_2\_RR\_S10 2CLCmesc\_2\_S2 2CLCmesc\_3\_RR\_S11 2CLCmesc\_3\_S5 2CLCmesc\_4\_RR\_S12 2CLCmesc\_4\_S6 2CLCmesc\_5\_RR\_S13 2CLCmesc\_5\_S7 2CLCmesc\_6\_RR\_S14 2CLCmesc\_6\_S8 5M 15M 10M 20M 0 # Reads

Uniquely mapped

Unmapped: other

Mapped to multiple loci

Unmapped: too short

Mapped to too many loci