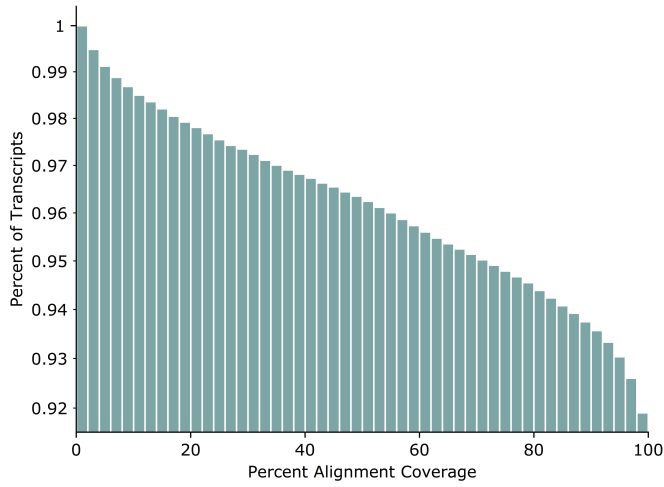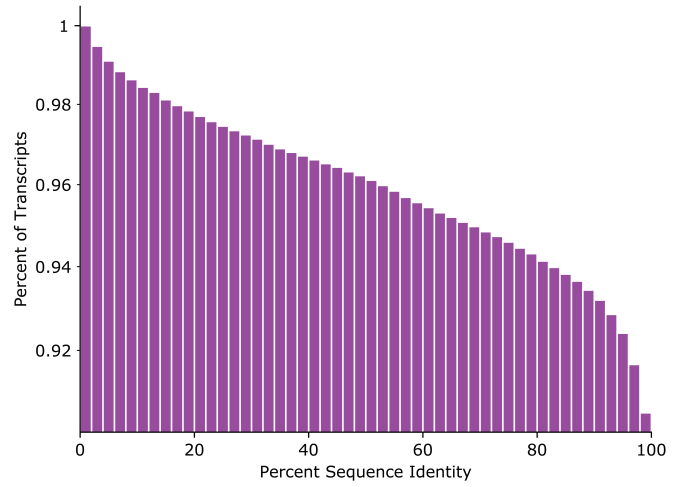**Figure S4. Alignment coverage and sequence identity cumulative distribution. (A)** Cumulative distribution showing how much of the IWGSC CS v1.0 transcripts map onto Triticum_aestivum_4.0. The y-axis shows the fraction of transcripts with percent coverage greater than or equal to coverage on the x-axis. **(B)** Cumulative distribution showing the sequence identity of IWGSC CS v1.0 transcripts mapped onto Triticum_aestivum_4.0. The y-axis shows the fraction of transcripts with sequence identity greater than or equal to sequence on the x-axis.