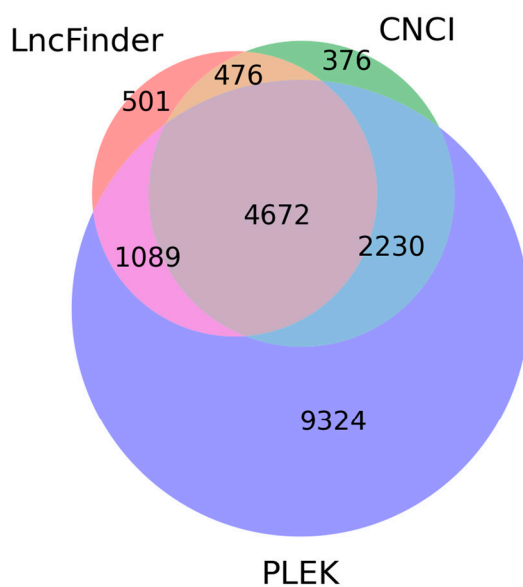
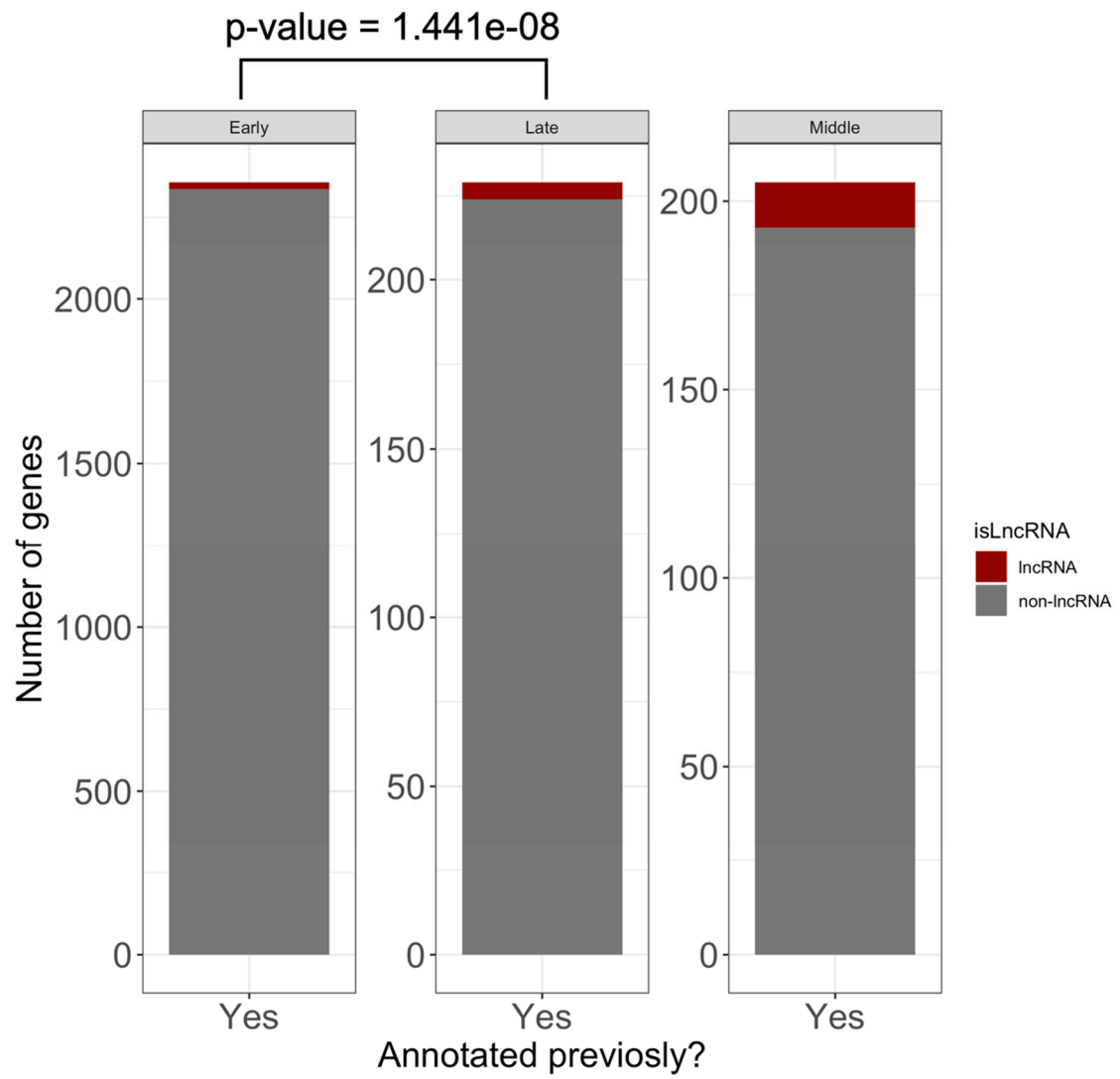


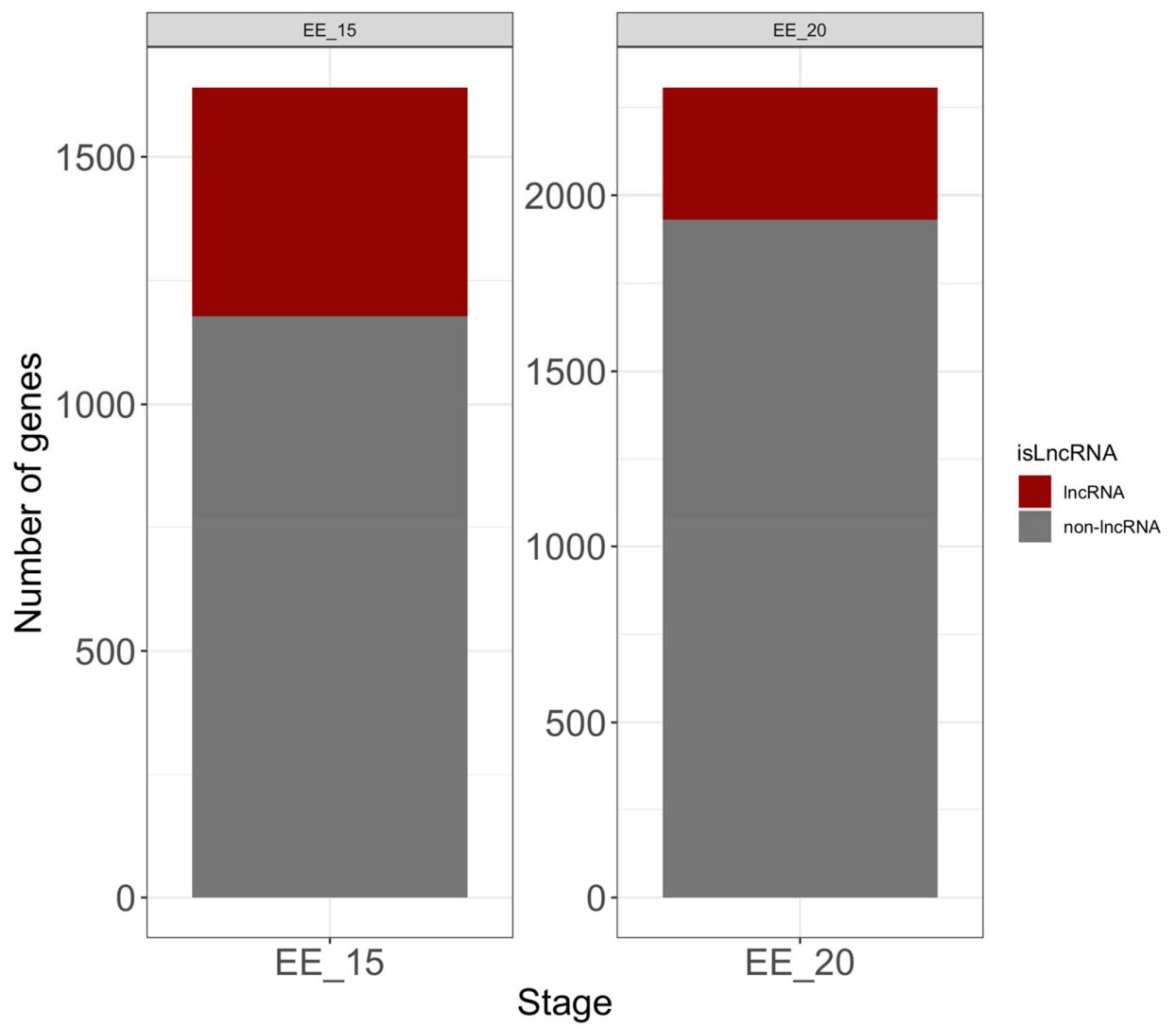
Supplementary Figure S1. Number of previously unannotated genes for each A, B and R chromosomes.



Supplementary Figure S2. Number of lncRNAs identified by three tools.



Supplementary Figure S3. Bar plot showing the fractions of lncRNAs among reference transcripts expressed during early, middle and late stages of triticle seed development.



Supplementary Figure S4. Bar plot showing the fractions of lncRNAs among the transcripts expressed specifically in EE samples (Germ + Endosperm).