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EpidiÅi dana piÄa kao blagdanska Åikola zasjajaja. lekcijw mi sole pamet hoÄe i njemacki u 100 lekcija pdf. wersten hat jedan drugi stavljen iz dana u dana,. njemacki jezik u 100 lekcija pdf - lekcijw mi sole pamet. F-box proteins (FBR), a subfamily of F-box proteins that interacts with the TFIID subunit of the general transcription factor IID (TFIID), and have crucial roles in regulating plant development and metabolism, including seed dormancy. In this study, we demonstrated that HOMEBOX-PROTEIN-1 (HB1) mRNA is abundantly and constitutively expressed in developing seeds and restricted to seed tissues, including inner and outer endosperm, embryo, and integuments. Using protoplast-based transient expression analysis, we also demonstrated that HB1 is a positive regulator of endosperm development, and overexpression of HB1 altered seed development during germination, resulting in a significant increase in germination rates. Moreover, localization studies showed that HB1 protein is present throughout the endosperm of mature developing seeds, co-localizing with a TALE protein. Taken together, our results provide the first evidence to indicate that HB1 may act as a positive regulator of seed dormancy. Eukaryotic genomes harbor small non-coding RNAs (sRNAs) that include microRNAs (miRNAs) and short-interfering RNAs (siRNAs) to regulate gene expression at the post-transcriptional level. However, the nature of sRNAs in plants is much less understood than in animals and fungi. Here, we carried out a genome-wide identification and characterization of plant sRNAs in the halophyte and salt-tolerant model plant, *Arabidopsis thaliana* (*Arabidopsis*). Using high-throughput sequencing technology, we identified 166 known and 322 novel miRNAs in *Arabidopsis* seedlings grown under normal (0.05 g L⁻¹ NaCl) and salt (3.5 g L⁻¹ NaCl) conditions. To better understand the function and biogenesis of plant miRNAs, we subjected