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What is the significance of SAS-MAKE.DAT and SAS-OPTIONS.EXE ? A: SAS-MAKE.DAT is a computer-readable and writeable text file that contains the SAS version information that can be used by the installer software. When the SAS install software detects the SAS version information in the make.dat file, it automatically modifies the installation options. For example, if the SAS version in the make.dat file is Version 9.3, it automatically enters the version 9.3 option in the options.txt file. The default name for this file is OPTIONS. Ref: sas-options.exe: This is a tool used by the SAS Installing Software to control SAS options before the installation. For detailed description about SAS options, refer to the SAS 9.2 and later Release Notes. For example, if the SAS installation is to run with the SAS/ACCESS Configuration Option, enter the ACCESS option to the SAS options tool. HELLA1 and HELLA2 genes control leaf size and flowering time in Arabidopsis. The floral transition is an important step in plant development that leads to transition from the vegetative to the reproductive phase. The timing of the transition is associated with diverse abiotic and biotic stresses. We have identified the HELLA (HEARTLESS, LECUCHE, LAFL) genes as being strongly down-regulated during the floral transition. Expression analysis of HELLA mutants demonstrated their tissue specific and stress-dependent regulation. A genome-wide screen and genetic analysis revealed that both HELLA genes are essential for the regulation of plant growth and development. Furthermore, mutations in both HELLA genes result in larger leaves and plants with shorter petioles, bigger sepals, an early flowering phenotype and larger siliques. Consistent with our data, several studies have shown that overexpression of HELLA genes in plants leads to altered vegetative and floral development. Moreover, HELLA genes may control petiole elongation by the regulation of cytokinin levels. Genetic and biochemical experiments showed that HELLA proteins are localized to the cytoplasm and participate in the control of cell growth and cell division in conjunction with the RAS/ROP-GTP f3e1b3768c

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