

**Cover image:** Pictured is a topological representation of SeedNet, an online, searchable model of the network of transcriptional interactions among genes implicated in the dormancy and germination of seeds from the thale cress plant, *Arabidopsis thaliana*. George W. Bassel et al. created SeedNet to probe the genetic interactions that underpin seed germination, an ecologically and agronomically important plant trait. The authors' model suggests that seed dormancy likely evolved by coopting genetic mechanisms that control abiotic stress and help plant cells transition through developmental phases. See the article by Bassel et al. on pages 9709–9714. Image courtesy of George W. Bassel.

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