

PEBP GENE FAMILY (PHOSPHATIDYL ETHANOLAMINE-BINDING PROTEINS)

TFS1 (Saccharomyces) RCN2 (Rice) RCN1 (Rice) SP (Lycopersicon esculentum) CENTRORADIALIS (ATC) BnTFL1-1 (Brassica napus) TERMINAL FLOWER 1 (TFL1) BROTHER OF FT (BFT) OsFT (rice) CiFT (Cichorium intybus)

Adapted from Kobayashi et al.,1999

Arabidopsis proteins

TWIN SISTER OF FT (TSF)

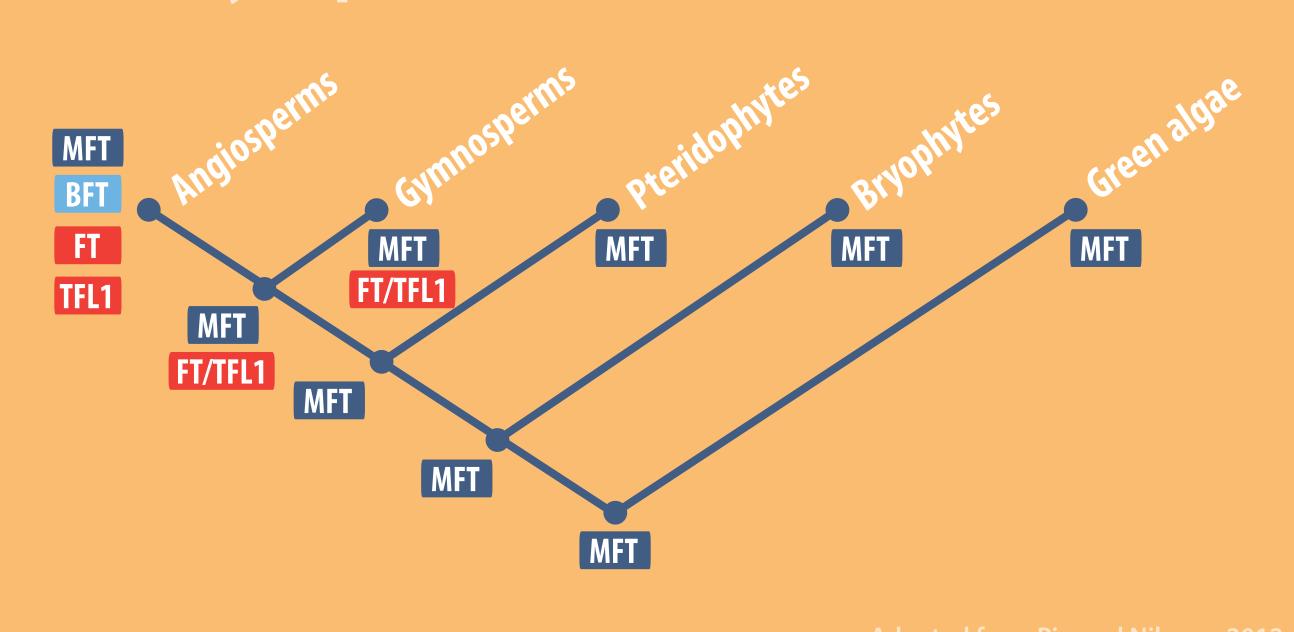
FLOWERING LOCUS T (FT)

MOTHER OF FT (MFT)

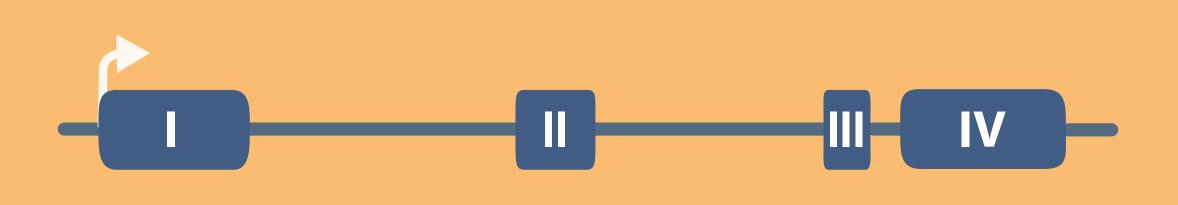
hPEBP (Human)

PEBP family in plant evolution

0,05

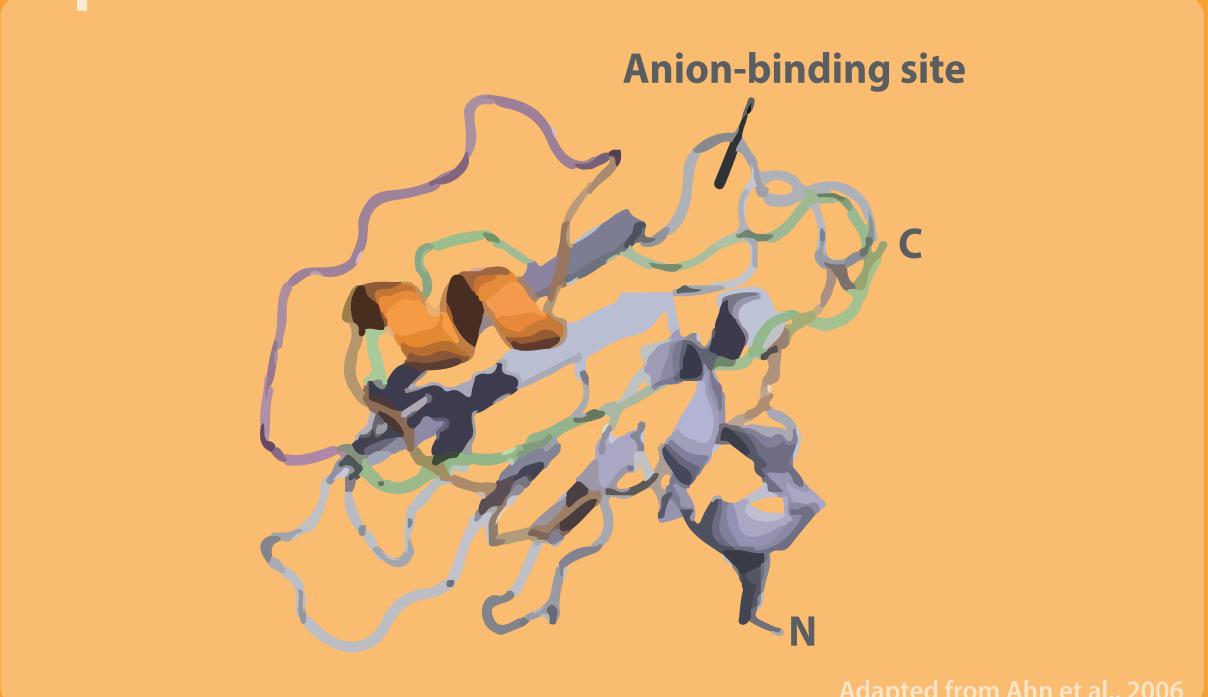


FT protein sequence



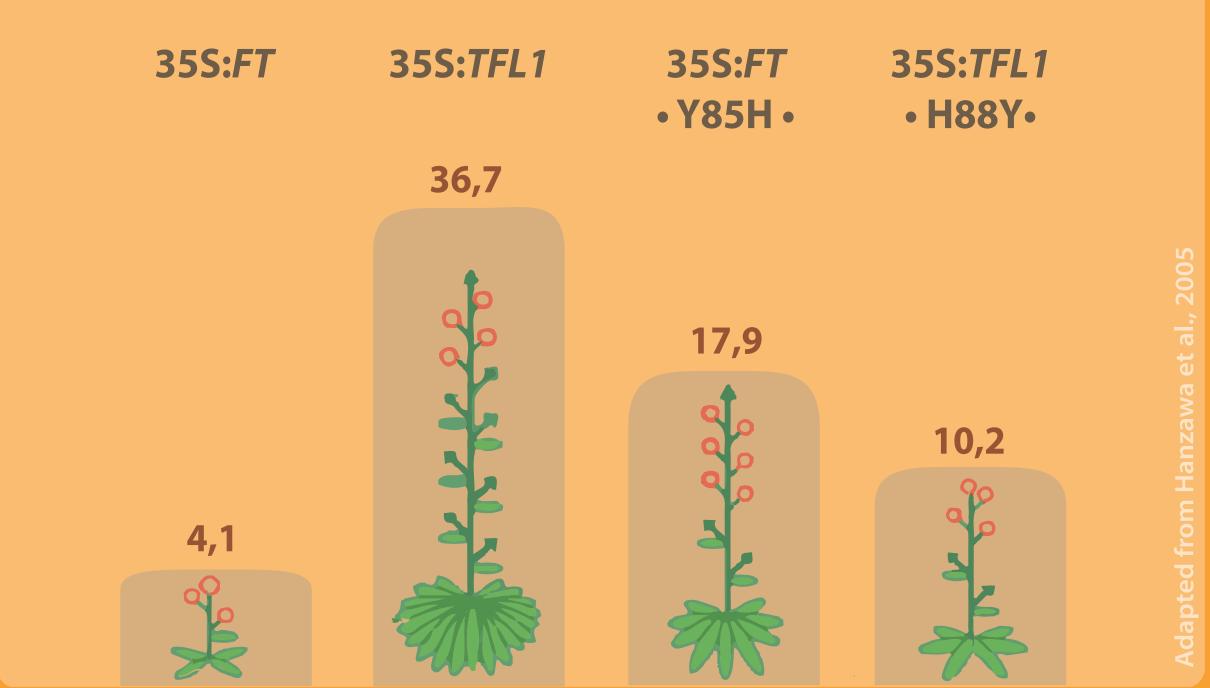
- Small protein of 175 amino acids (4 exons, 22 kD);
- No DNA binding domain.

FT protein structure

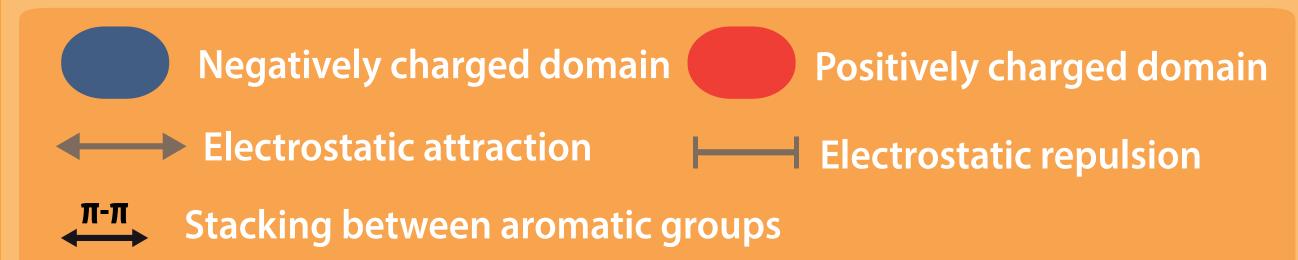


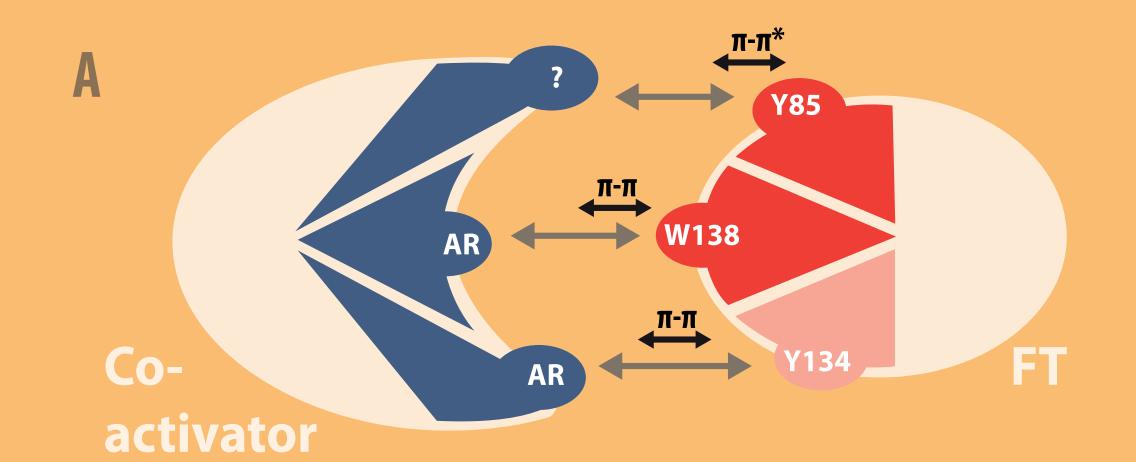
FT and TFL1 antagonistic functions depend on few amino acids

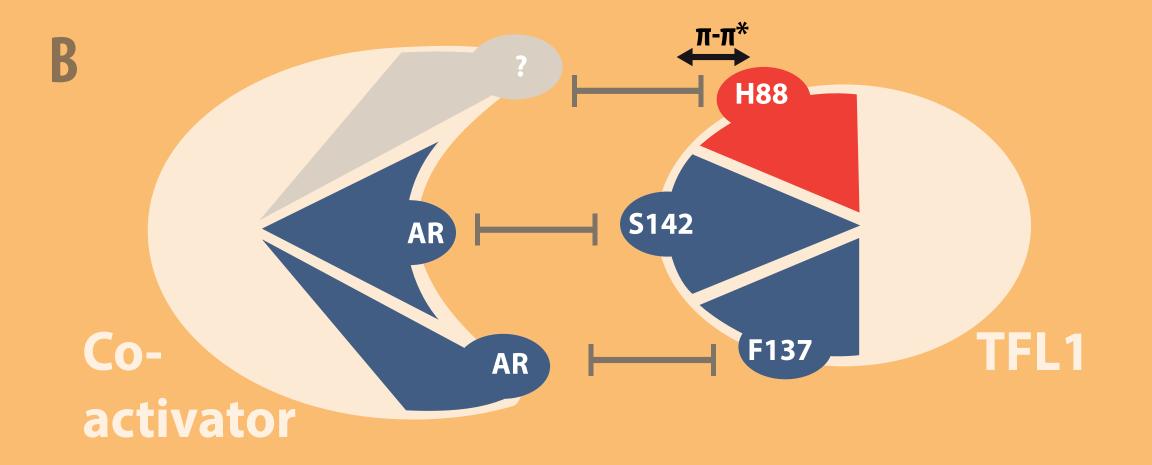
Number of leaves at flowering under 16 hours-long days

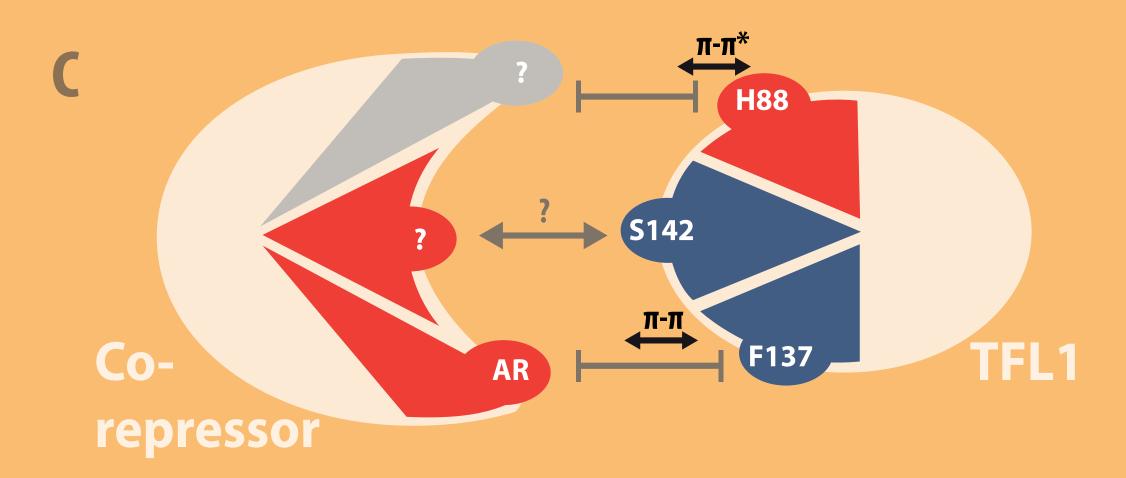


Mechanistic model for FT and TFL1 antagonistic function



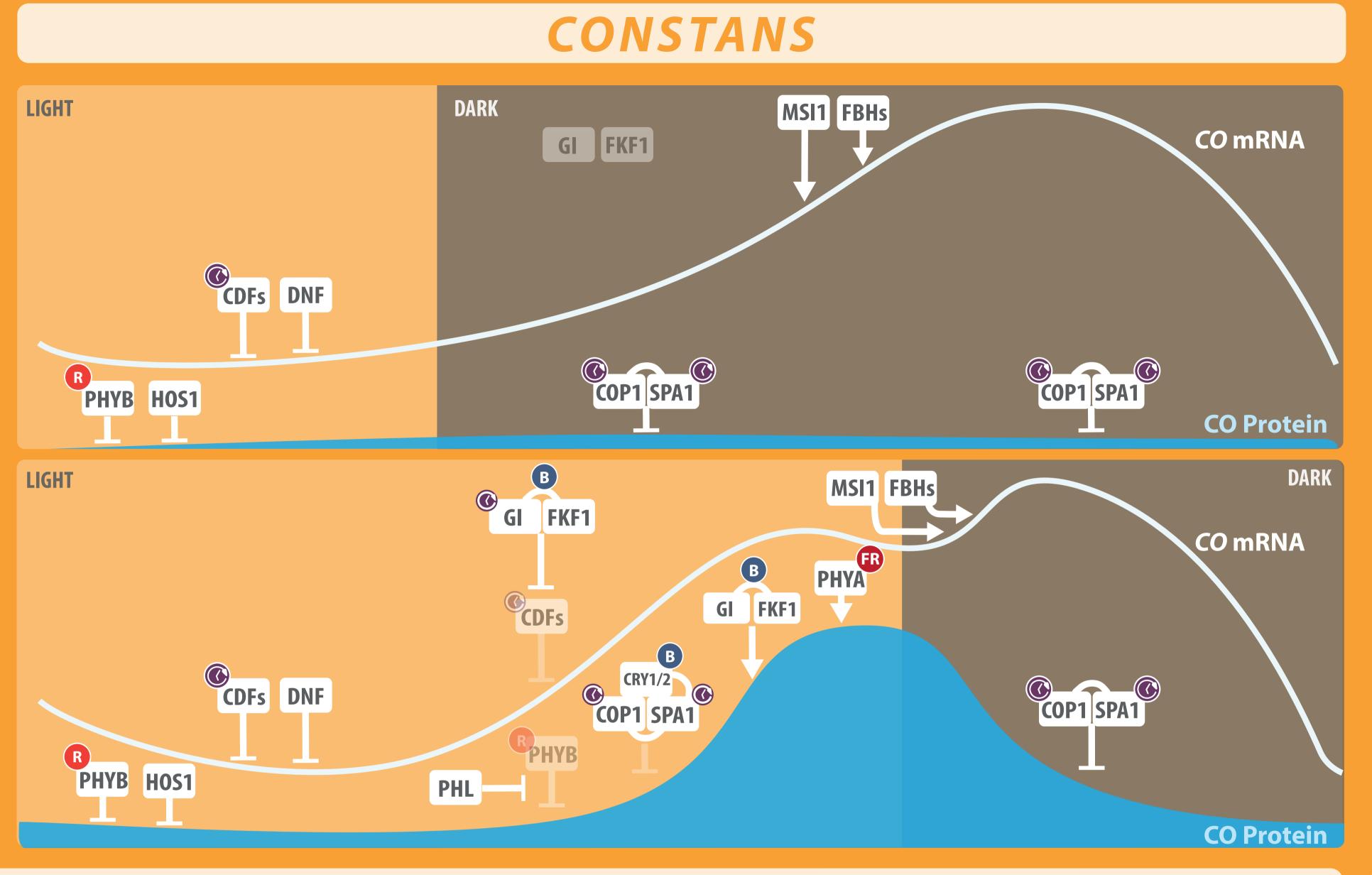






FT binds a coactivator (A), while TFL1 may either exlude a coactivator (B) or interact positively with a corepressor (C) (Ho and Weigel, 2014).

TIME-COURSE OF CONSTANS AND *FLOWERING LOCUS T* TRANSCRIPTIONAL AND POST-TRANSLATIONAL REGULATION



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