

COP1 is a negative regulator of seed germination in Strigolactone signaling.

Shigeo Toh^{1,2}, Duncan Holbrook-Smith¹, Yuichiro Tshchiya^{1,2,3}, Toshinori Kinoshita^{2,3} and Peter McCourt¹

¹*Cell and Systems Biology, University of Toronto*, ²*Division of Biological Science, Graduate School of Science, Nagoya University*, ³*Institute of Transformative Bio-Molecules, Nagoya University*

Strigolactones (SLs) are host factors that stimulate seed germination of parasitic plant species *Striga*. To understand the roles of SLs in seed germination, it is necessary to develop a tractable experimental system using model plants such as *Arabidopsis*. We reported that thermoinhibition, which involves exposing seeds to high temperatures, uncovers a clear role for SLs in promoting *Arabidopsis* seed germination. Both SL biosynthetic and signaling mutants showed increased sensitivity to seed thermoinhibition. The synthetic strigolactone GR24 does not rescue *max2-1* seed germination.

Hormone analysis revealed that SLs alleviate thermoinhibition by modulating levels of the two plant hormones, GA and ABA. Recently, we also reported that GR24 directly binds the HTL α/β hydrolase in *Arabidopsis* in vitro. Strigolactones promoted an interaction between HTL and the F-box protein MAX2 in yeast. We also found *htl* mutant in our GR24 insensitive screening using thermoinhibitor. These results suggest that HTL is involved in SL signaling during seed germination in *Arabidopsis*. Molecular analysis using a hypocotyl elongation assay showed SLs regulate the nuclear localization of the COP1 ubiquitin ligase, which in part determines the levels of light regulators such as HY5. Genetic analysis revealed that *cop1* single mutant, *cop1 max2* double mutant and *cop1 htl-3* double mutant showed thermo-tolerant seed germination phenotype. These results indicated that COP1 is a negative regulator of seed germination and is genetically at or downstream of MAX2 and HTL in SL signaling with respect to seed germination.