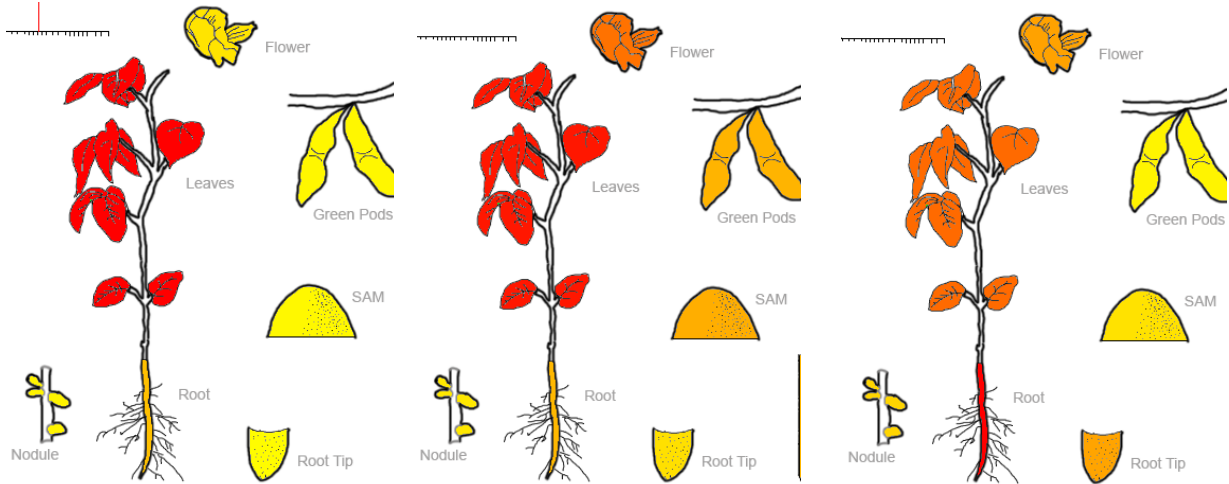


Glyma08g08810 (FLS2-like)

Glyma01g40130 (ACA4,11-like)

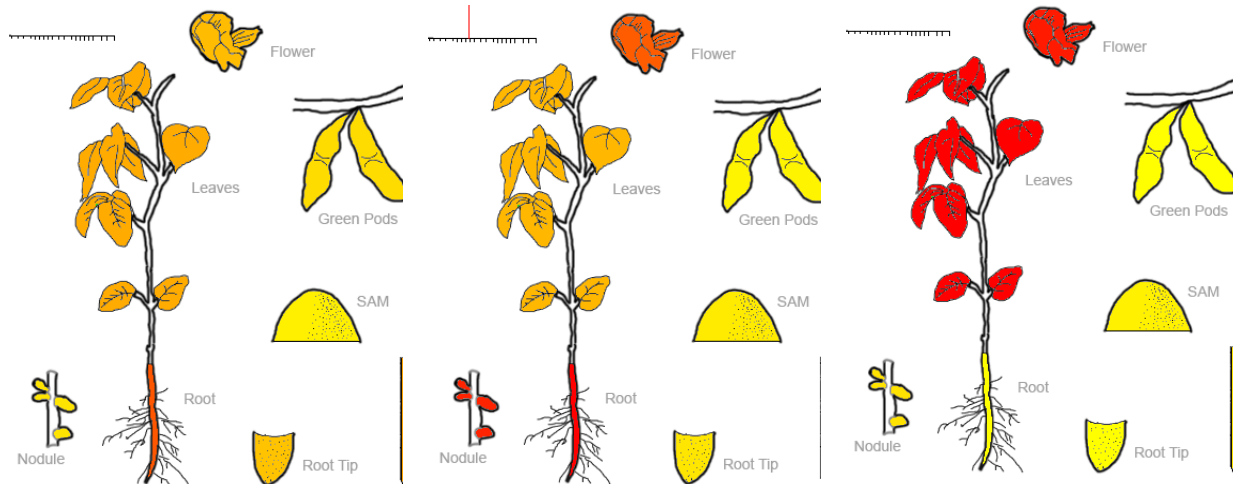
Glyma03g31420 (ACA4,11-like)



Glyma11g05190 (ACA4,11-like)

Glyma19g34250 (ACA4,11-like)

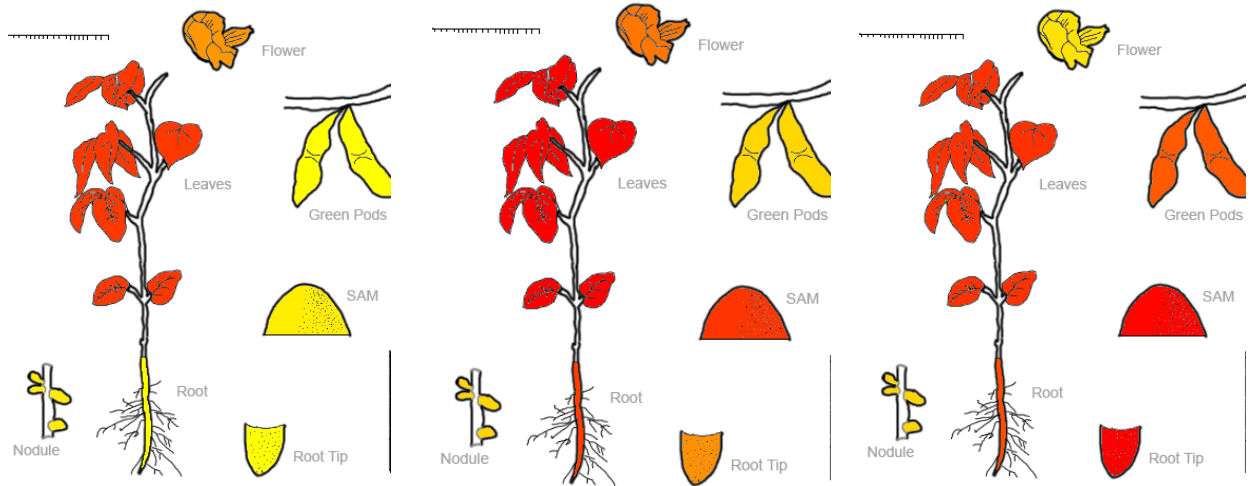
Glyma11g15700 (MPK4,6-like)



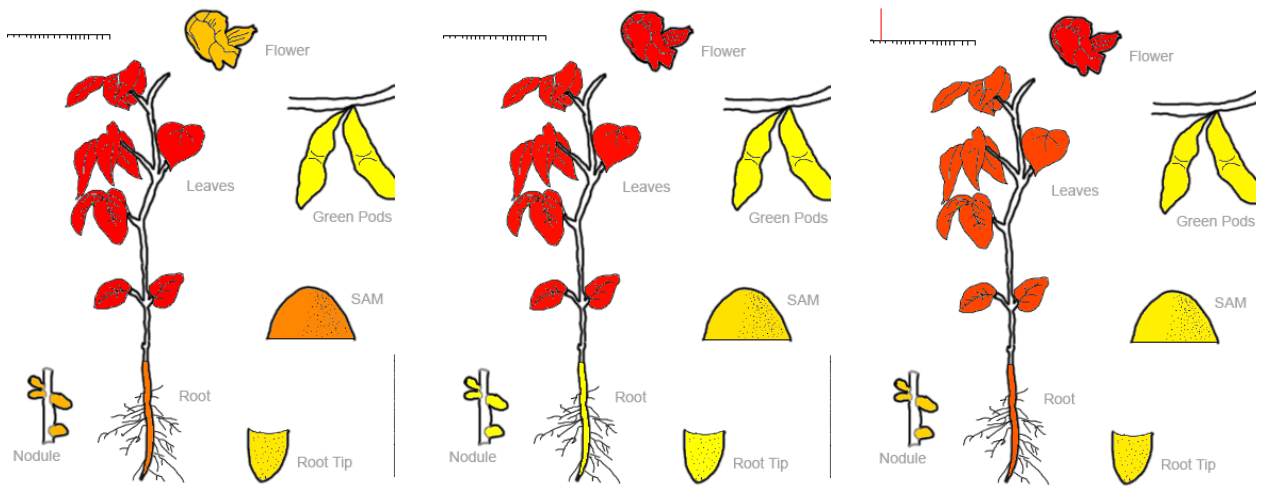
Glyma12g07770 (MPK4,6-like)

Glyma06g09520 (PEPR1,2-like)

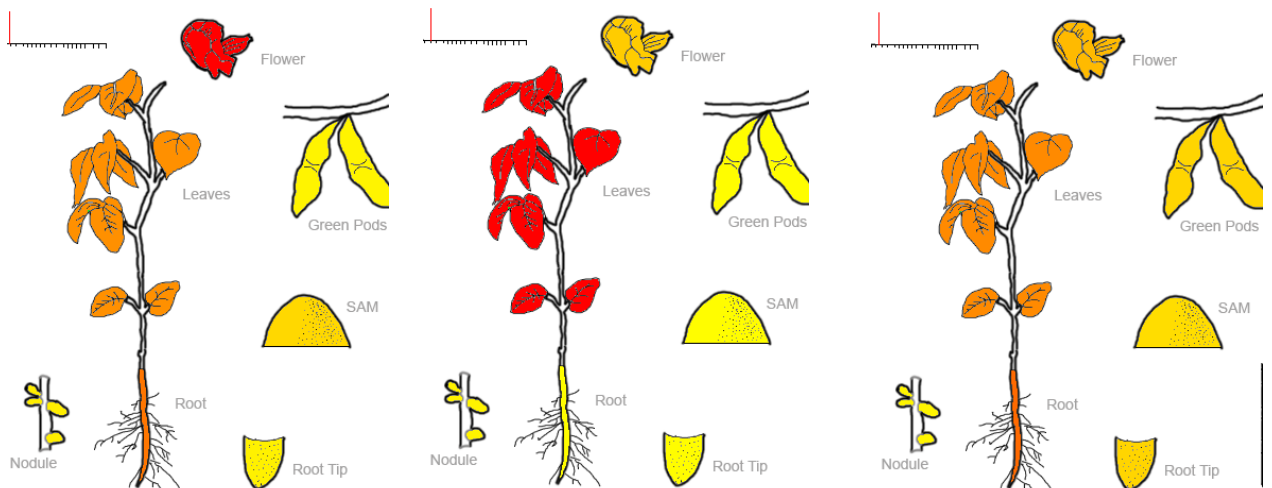
Glyma08g08810 (PEPR1,2-like)



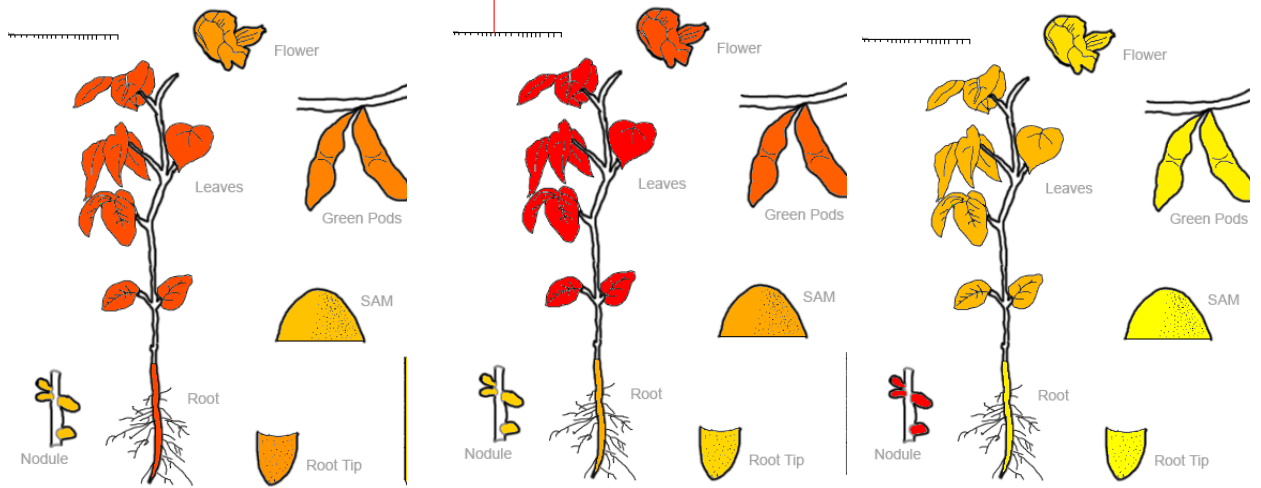
Glyma06g40690 (RPP1,4,5-like) Glyma06g40950 (RPP1,4,5-like) Glyma06g40980 (RPP1,4,5-like)



Glyma16g10340 (RPP1,4,5-like) Glyma03g33660 (MLO-like) Glyma02g39870 (WRKY33-like)



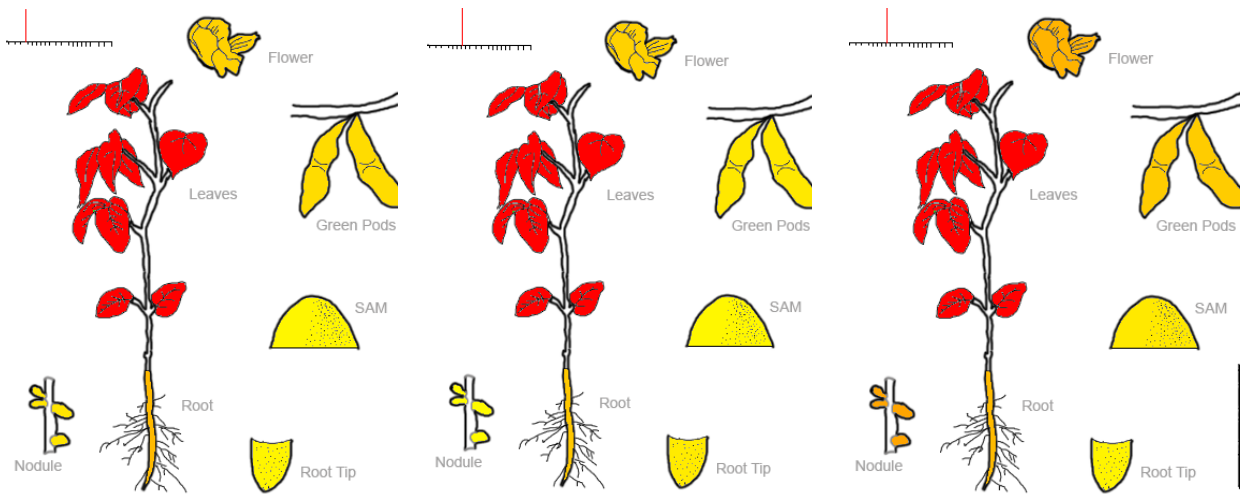
Glyma14g38010 (WRKY33-like) Glyma13g31280 (LOX1-like) Glyma02g35230 (SYP121,122-like)



Glyma10g10200 (SYP121,122-like)

Glyma08g36720 (MYC2-like)

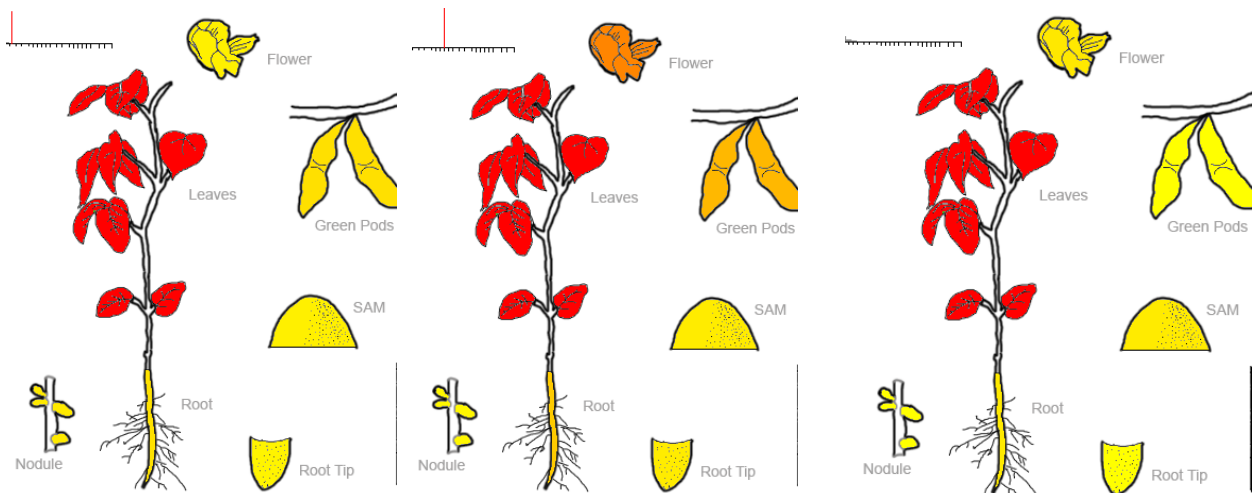
Glyma18g05890 (PR14-like)



Glyma09g08290 (JAZ1-like)

Glyma13g17180 (JAZ1-like)

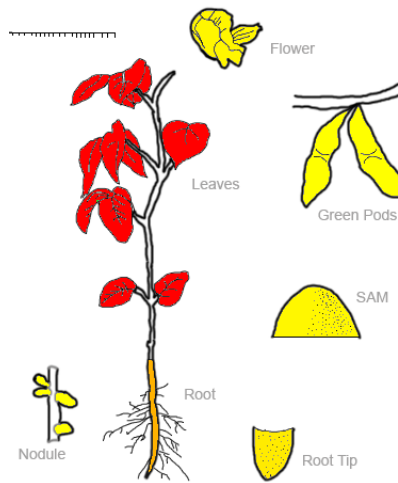
Glyma15g19840 (JAZ1-like)



Glyma17g05540 (JAZ6-like)

Glyma16g01220 (JAZ6-like)

Glyma13g29070 (JAZ8-like)



Glyma15g09980 (JAZ8-like)

Supplemental Fig. 1. Expression pattern of several PTI (PAMP-triggered immunity)-, ETI (effector-triggered immunity)-, JA (jasmonic acid)-, and defence-related genes in different tissues of soybean analyzed using the soybean eFP Browser (http://bar.utoronto.ca/efp_soybean/cgi-bin/efpWeb.cgi). Sixty-seven defense genes, whose expression was significantly differed between BLP (bacterial leaf pustule)-susceptible and BLP-resistant near-isogenic lines after *Xanthomonas axonopodis* pv. *glycines* (Kim et al 2011), were analyzed using the soybean eFP Browser. Among 67 genes, 28 genes, whose expression was less in green pods than in leaves, were depicted in this supplemental figure. Expression of 2 genes was higher in green pods than in leaves, and others were similar between green pods and leaves.