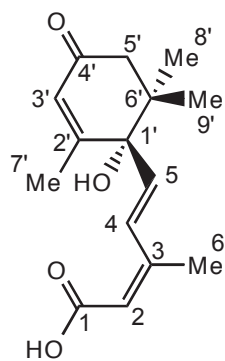
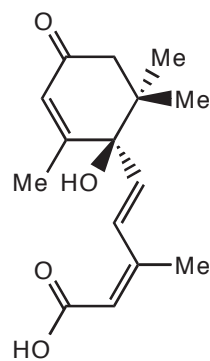


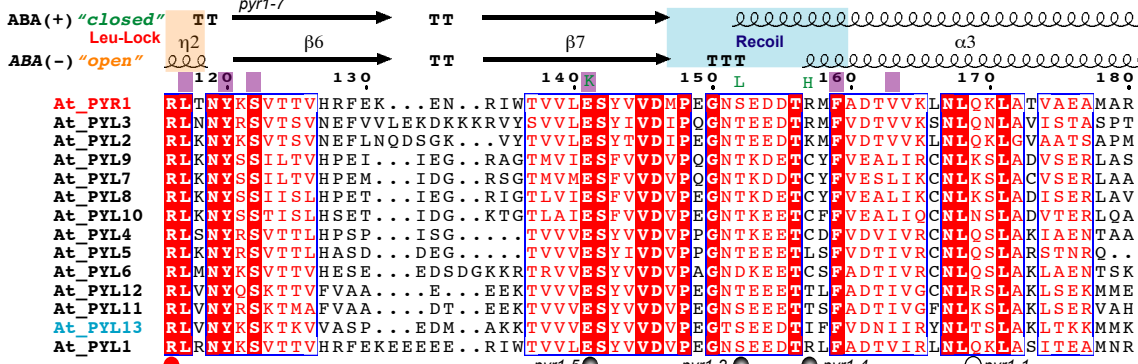
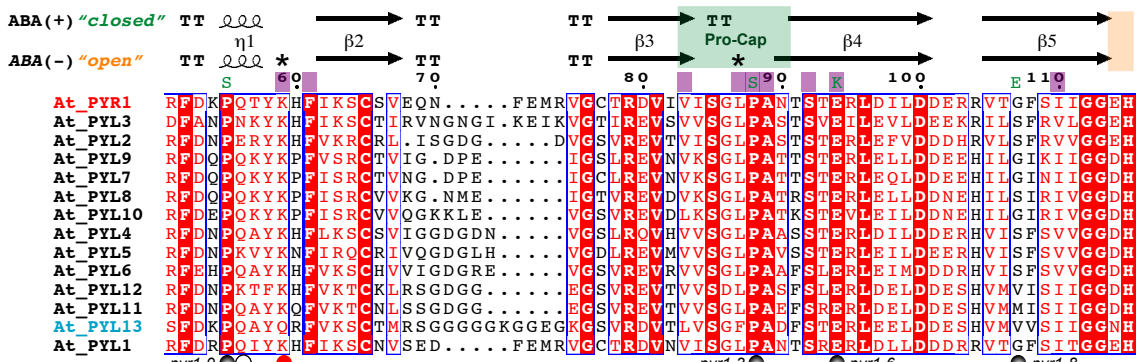
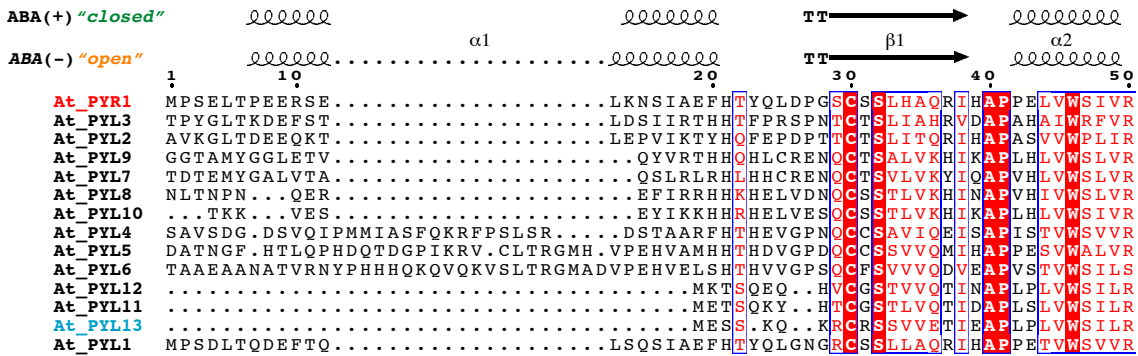
A

(+)-ABA

B

(-)-ABA

Nishimura et al. Figure S1

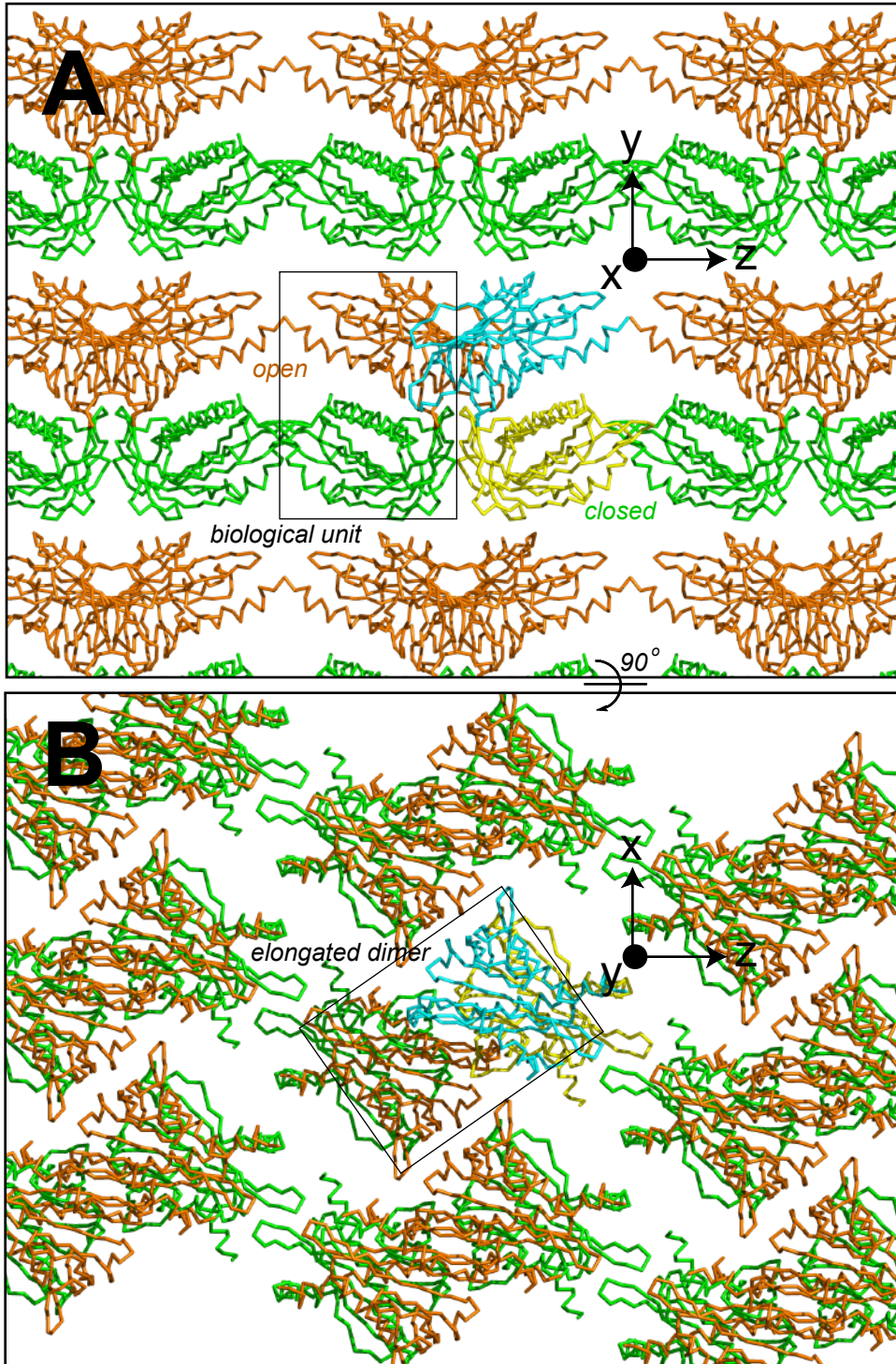


190 Gene id:

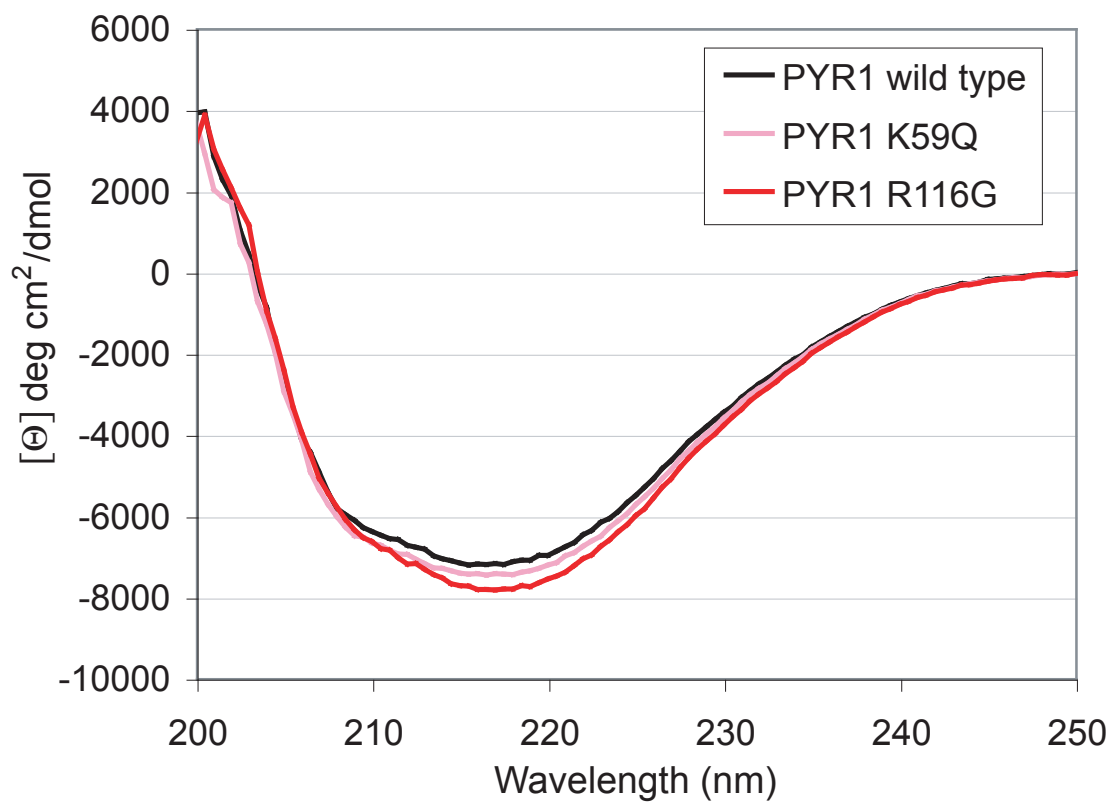
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At_PYL3	AT1G73000
At_PYL2	HDDE.....	AT2G26040
At_PYL9	QDITQ.....	AT1G01360
At_PYL7	QDITNSIATFCNASNGYREKNHTE TNL	AT4G01026
At_PYL8	QDTTESRV.....	AT5G53160
At_PYL10	ES.MEKKI.....	AT4G27920
At_PYL4	ESKKKMSL.....	AT2G38310
At_PYL5	AT5G05440
At_PYL6	FS.....	AT2G40330
At_PYL12	LT.....	AT5G45870
At_PYL11	LKL.....	AT5G45860
At_PYL13	AT4G18620
At_PYL1	NNNNNSSQVR.....	AT5G46790

Pro-Cap ● mutational sites
 Leu-Lock ● point mutation
 Recoil ○ stop (green letters)
 ABA binding ● designed mutation

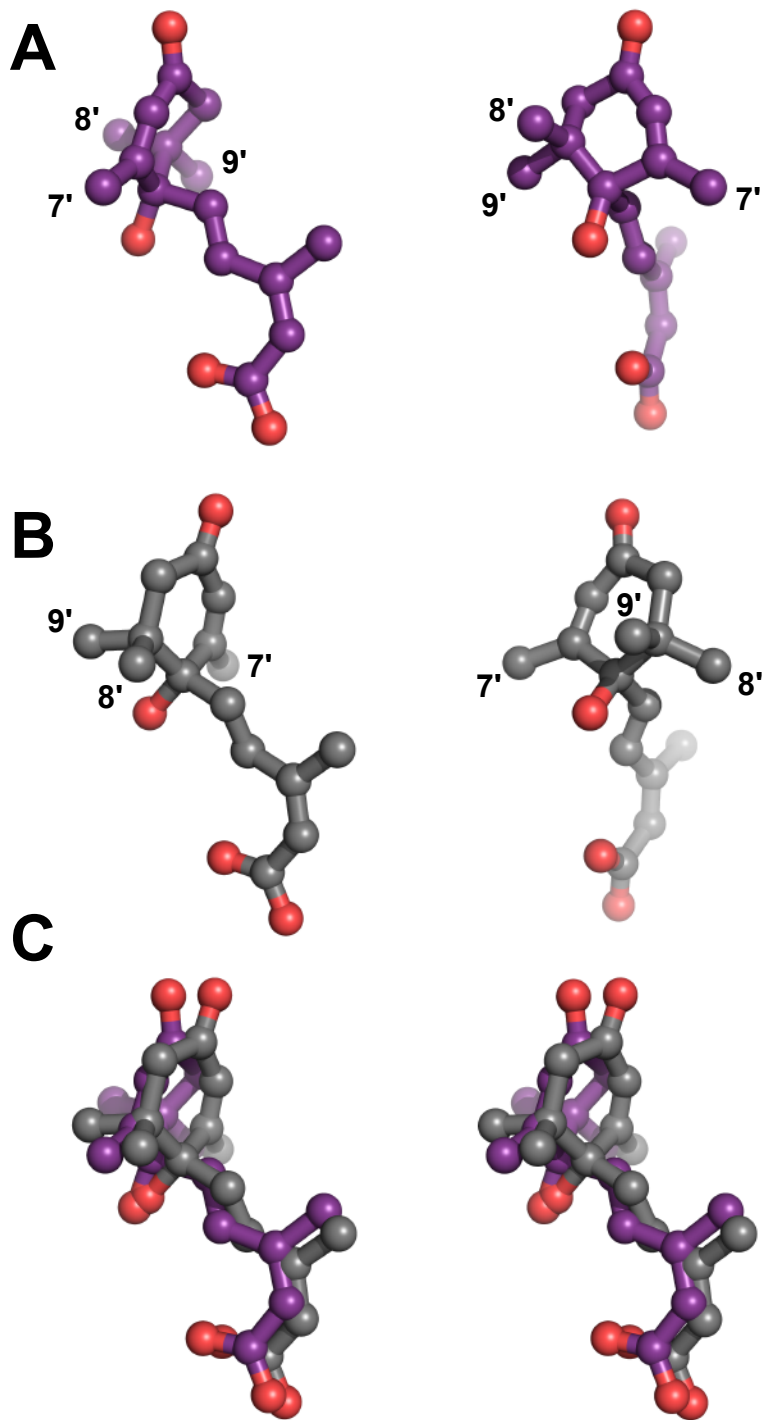
Nishimura et al. Figure S2



Nishimura et al. Figure S3



Nishimura et al. Figure S4



Nishimura et al. Figure S5