

Fig. 1D* pBI121::chitEclipGFP:LAAQ				
	Fragments	Start	End	Category
	right Border	2478	2454	Miscellaneous
	CaMV 35S promoter	4974	5802	Promoter
	Chitinase signal	5844	5900	Localization Sequence
	ecliptic At-pHluorin	5907	6617	Gene
	low affinity Aequorin	6633	7196	Reporter Gene
	Terminator	7212	7413	Terminator
	left Border	8355	8331	Miscellaneous
	Restriction Sites cuts once / cuts twice	Location		Sequence
	BamHI	5822, 7155		ggatcc
	Cla I	4538, 6287		aagctt
	HindIII	4951		aagctt
	KpnI	7419		ggtacc
	NotI	14372		gcggccgc
	SacI (SstI)	7425		gagctc
	Sall	6618		gtcgac
	XbaI	5816		tctaga
	XhoI	6628		ctcgag

*refers to Figure 1 in Gao et al. 2004 (PubMed 15020753; DOI: 10.1104/pp.103.032508)

