

Fig. 1B* pBI121::chitsmGFP5:AQ (14467 bp)				
	Fragments	Start	End	Category
	right Border	2478	2454	Miscellaneous
	CaMV 35S promoter	4974	5802	Promoter
	Chitinase signal	5844	5900	Localization Sequence
	smGFP5	5907	6617	Gene
	Aequorin	6633	7196	Reporter Gene
	Terminator	7212	7413	Terminator
	left Border	8355	8331	Miscellaneous
	Restriction Sites	Location		Sequence
	cuts once / cuts twice			
	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)

