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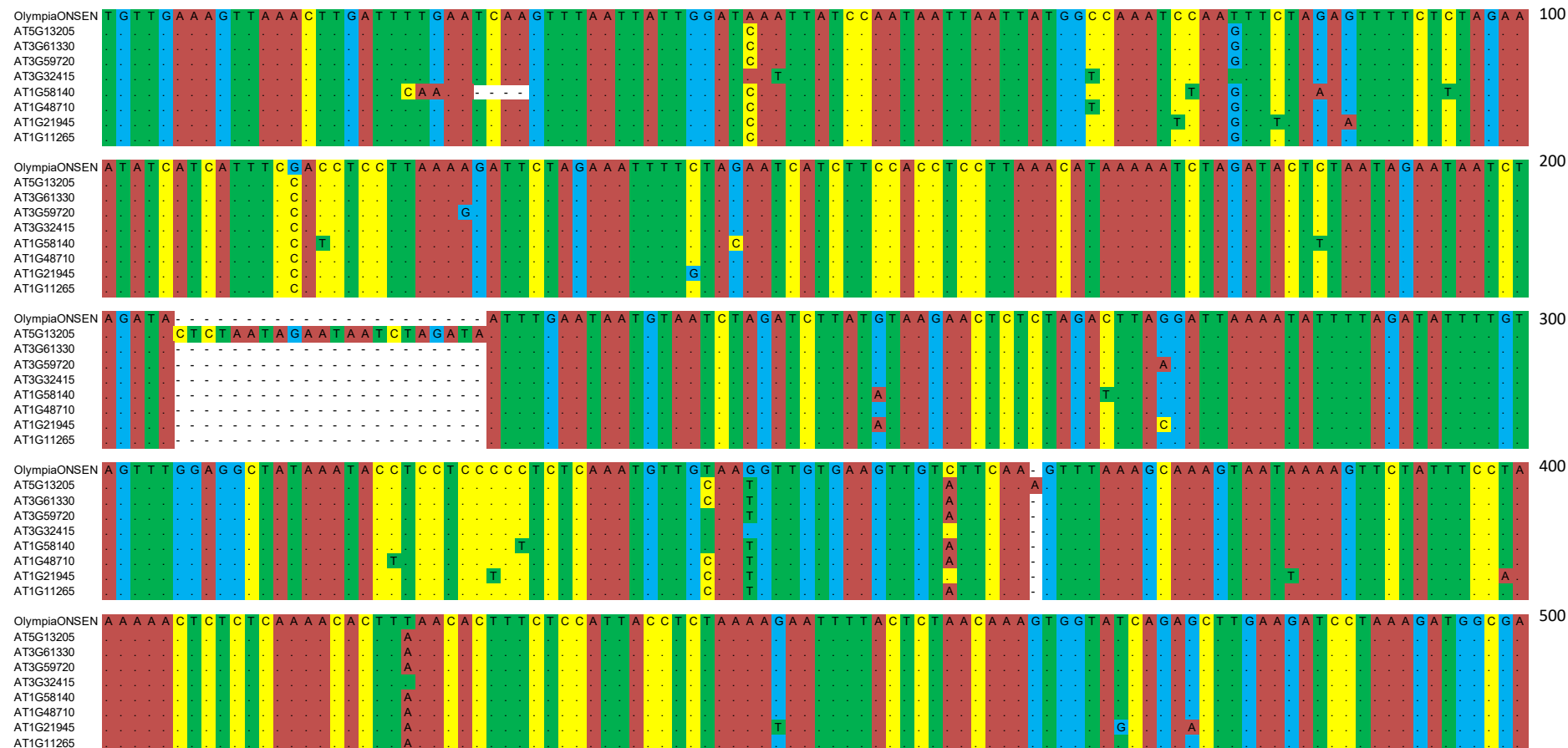
Title	Characterization of a heat-activated retrotransposon in natural accessions of <i>Arabidopsis thaliana</i>
Author(s)	Masuda, Seiji; Nozawa, Kosuke; Matsunaga, Wataru et al.
Citation	Genes & Genetic Systems, 91(6), 293-299 https://doi.org/10.1266/ggs.16-00045
Issue Date	2016-12
Doc URL	https://hdl.handle.net/2115/65346
Type	journal article
File Information	Supplementary materials.pdf

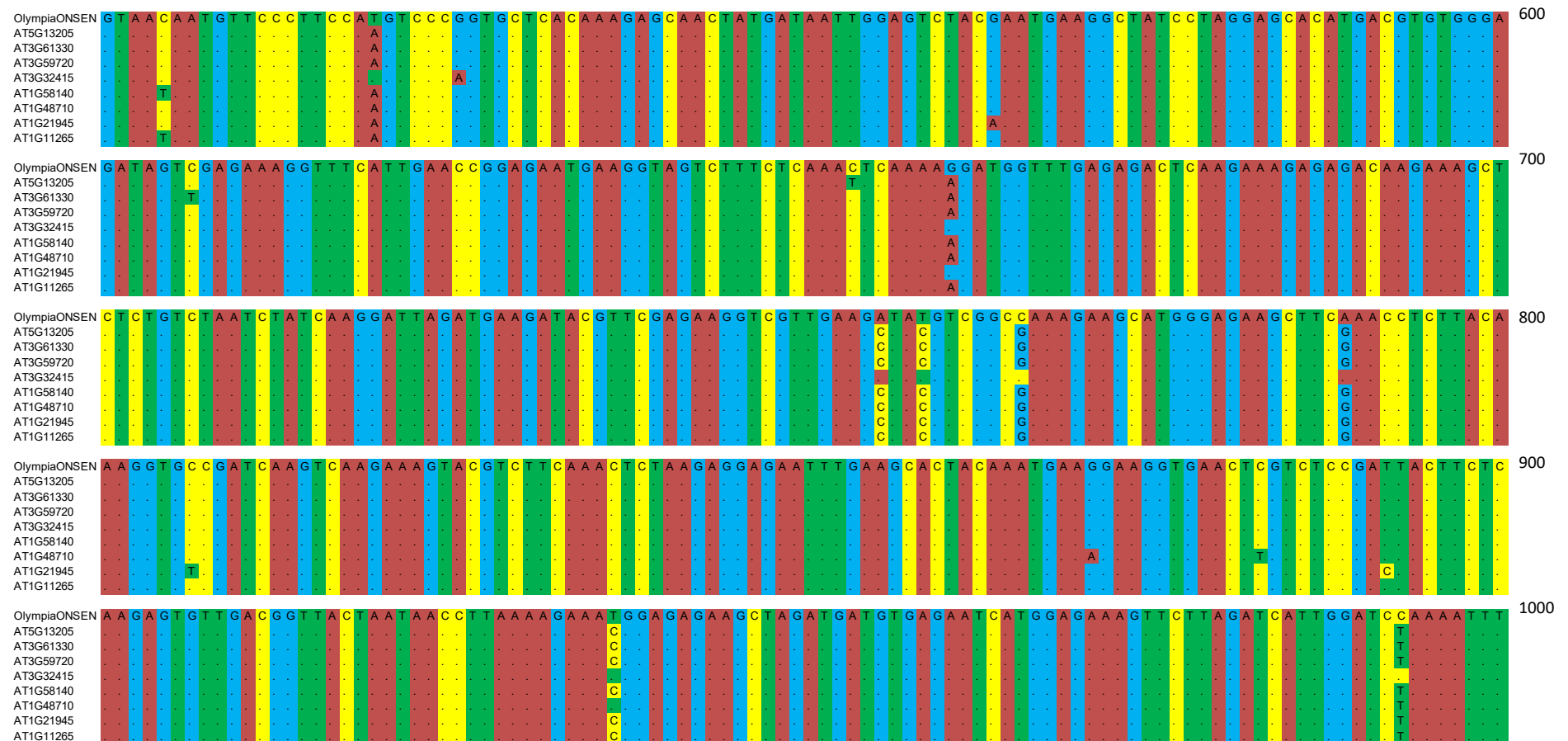


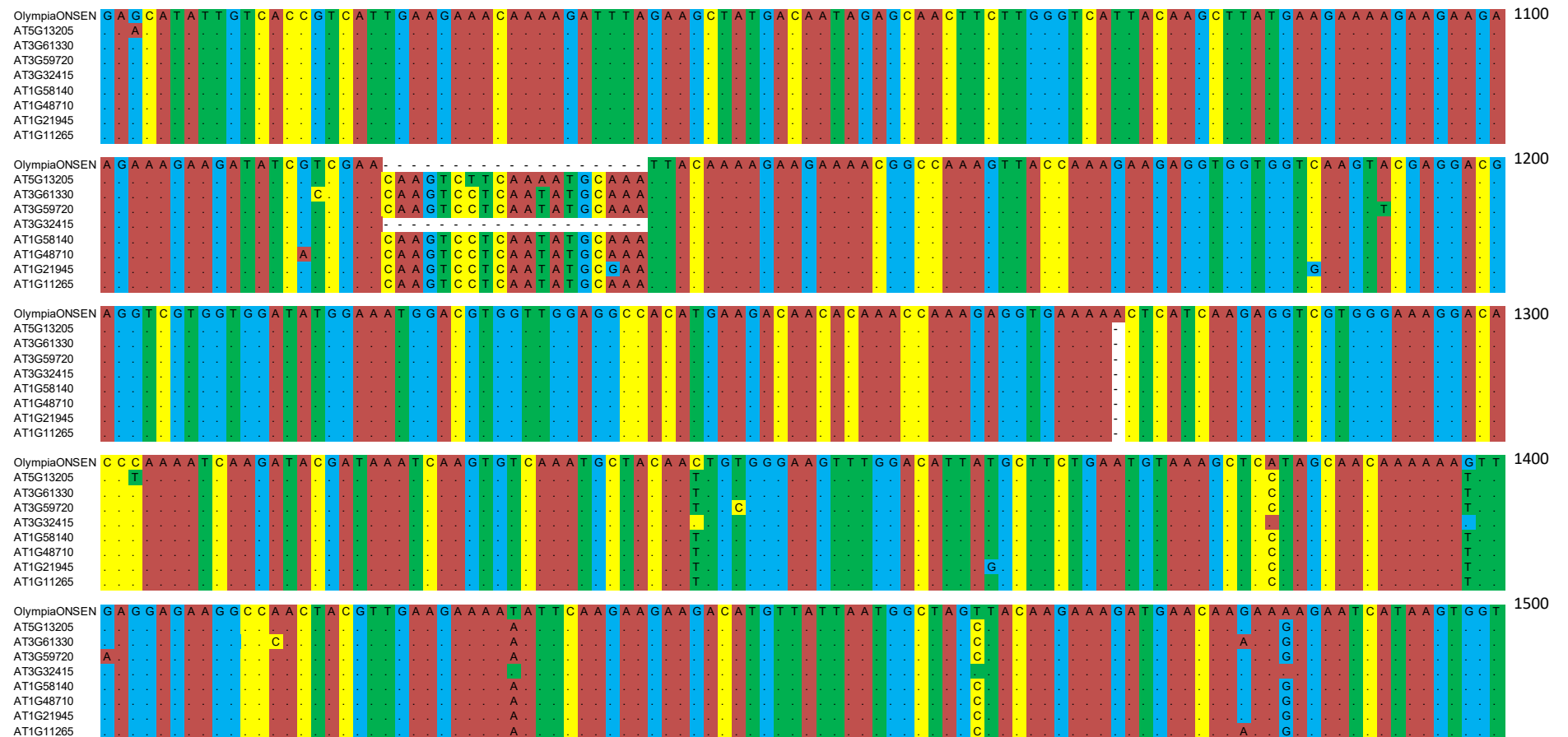
Supplementary Table S1. Primer sequences.

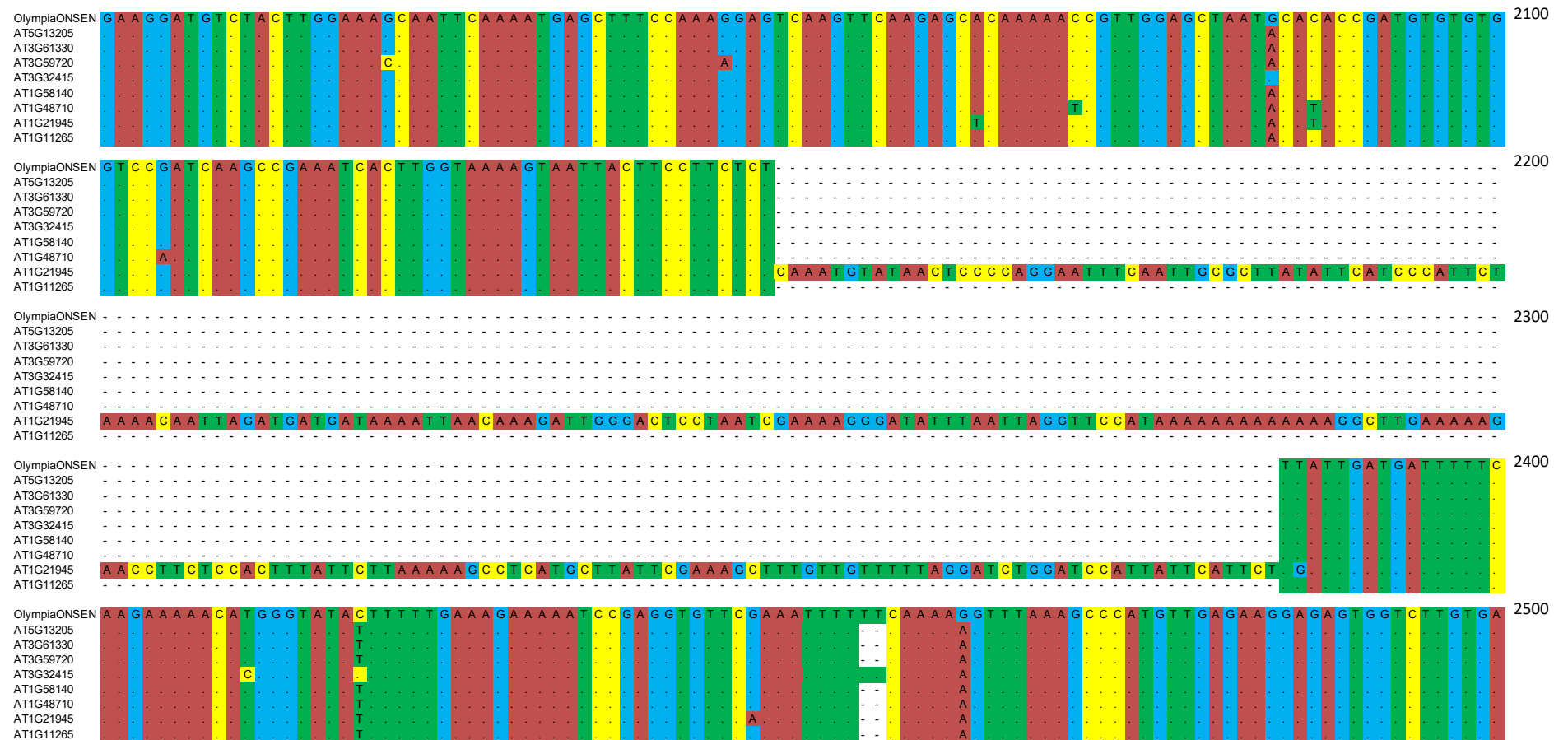
Target/Purpose	Name	Primer
Southern probe	ONSEN-F	TAA TGT TCC CTT CCA AGT CCC
	ONSEN-R	GCT TGT AAT GAC CCA AGA AGT
q-RT-PCR	COPIA78_4129F_RT	CCA CAA GAG GAA CCA ACG AA
	COPIA_4300R_RT	TTC GAT CAT GGA AGA CCG G
	HsfA2_qPCR_F	TGG GAT TCT CAT AAG TTC TCA ACA
	HsfA2_qPCR_R	TGG ATC AAT CTT TCT GAA TCC AT
	HSP90.1-qRTPCR-F	CAA CAA AGA AAT CTT CCT CCG TGA G
	HSP90.1-qRTPCR-R	CAT CGA GCT TGC TCT TAT CCG TTA
	18Sr_FW	CGT CCC TGC CCT TTG TAC AC
	18Sr-RV	CGA ACA CTT CAC CGG ATC ATT
PCR of 8 copies of <i>ONSEN</i>	AT1G48710-F	ACA TTC TTC GCG GGA AAT GT
	AT1G21945-F	ATG GCG AGA GAT TCT TGC AG
	AT1G58140-F	ATG GAG GCG GAG ATG TTG A
	AT1G11265-F	ATA TAT CAC CCG ATG TCC CCA
	AT3G32415-F	CAG AGG CAA GCT TTT TAC CA
	AT3G61330-F	TAA CTG GCT TCC TGA CCC AAA
	AT3G59720-F	CTT CAT TTT CCG CTT CAC GT
	AT5G13205-F	ACA CAA AAG TCC AGC TCT GGT
	8copies-REVERSE	CCT TCA TTC TCC GGT TCA AT
	Maseq-351	GTG AAG TTG TCT TCA AGT
	Maseq-701	CTC TGT CTA ATC TAT CAA
	Maseq-1051	AAT AGA GCA ACT TCT TGG
	Maseq-1411	ATT CAA GAA GAA GAC ATG
	Maseq-1751	AAG AGA CCA AGA AAG CAA
	Maseq-2109	AGT AAT TAC TTC CTT CTC
	Maseq-2453	GGT AGC GTG TGC GGT TTA
	Maseq-2804	GGA GCC AAC AAG AGA GGA
	Maseq-3151	AAG GAG AAG TGG AAA GAT
	Maseq-3501	CCA TAT GAG CAT GCA CTC
	Maseq-3851	GAG CTT GGT TGG AAG CTT

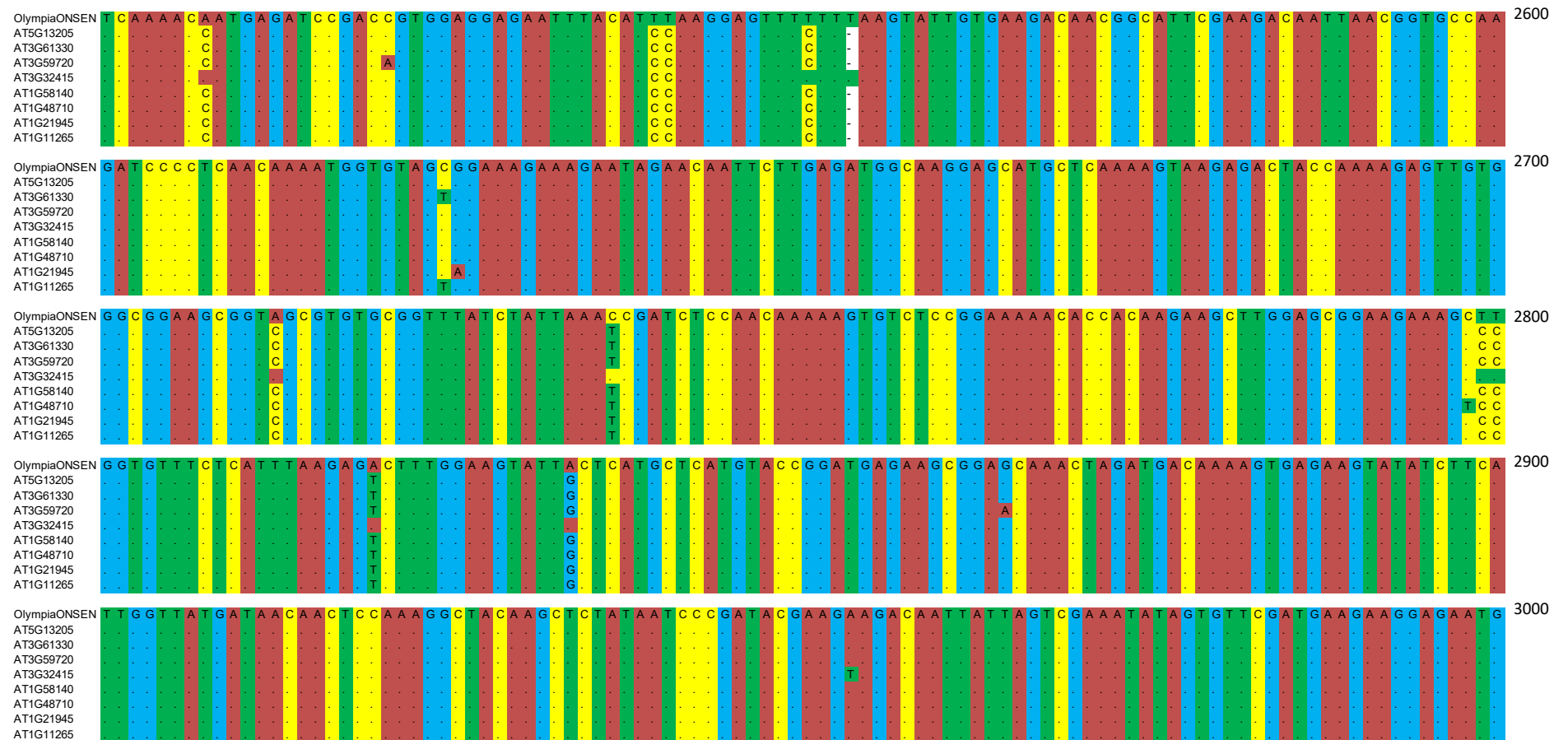
	Maseq-4201	CAT GTG TAT GCC ATG CTA
	Maseq-4553	TTA TTG GAT AAA TTA TCC
	Maseq-4911	CAT TTT AAC ACT TTC TCC
Cloning of <i>ONSEN</i> in Olympia-1	AT3G32415full-F	AGG CAA GCT TTT TAC CAG CA
	AT3G32415full-R	TTT CCA AGT CCA GAG AGG AGT
	Olympia-seq-R	TAG GCT CTT CAC TTG GTG GCT
	M13 (-21)	TGT AAA ACG ACG GCC AGT
	M13 Reverse	CAG GAA ACA GCT ATG ACC
Genotype of <i>ONSEN</i> in Col-0/Olympia-1 hybrid	AT1G11265-F	ATA TAT CAC CCG ATG TCC CA
	at1g11265-R	CCG AGG TCC TGA TGT TCT TGT
	AT1G21945part4-F	TGT AAG CAT CGT CGA TGG TCT
	AT1G21945part4-R	TCT GTT TCT TTG GCT TGG GA
	AT1G21945-F	ATG GCG AGA GAT TCT TGC AG
	AT1G48710-part2F	GAA ACG TGG CTG CAA GAA AA
	AT1G48710-part2R	GTT AAA GGG CAA ATT GGT CC
	AT1G48710-F	ACA TTC TTC GCG GGA AAT GT
	AT1G58140-F	ATG GAG GCG GAG ATG TTG A
	at1g58140F	TTC CAT GGT TTT GTT CAT TTG
	AT3G59720part4-F	CAT GGT TTG CAC ATT GTG CT
	AT3G59720part4-R	TCA GCA ATC GGT TTC GTC TT
	AT3G59720-F	CTT CAT TTT CCG CTT CAC GT

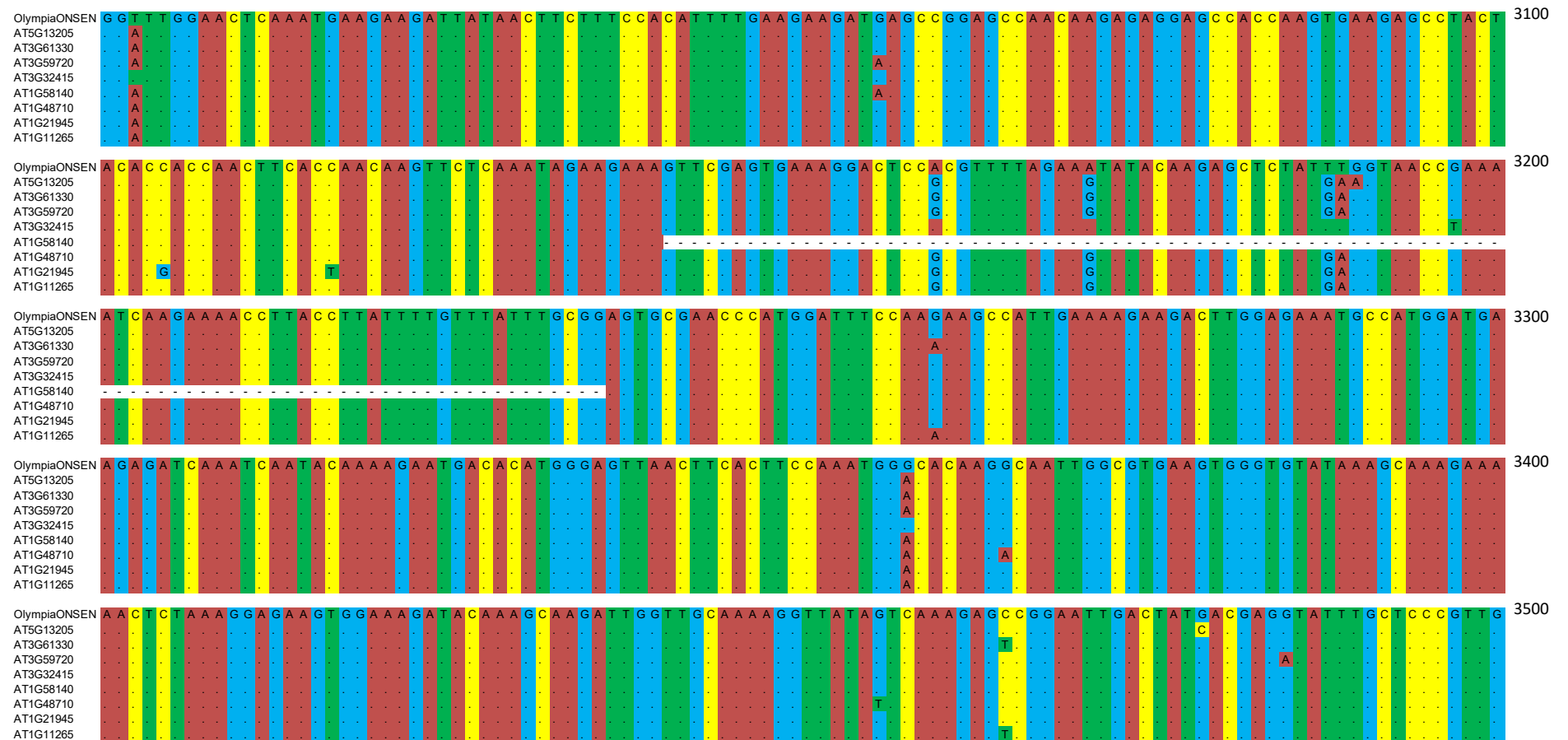


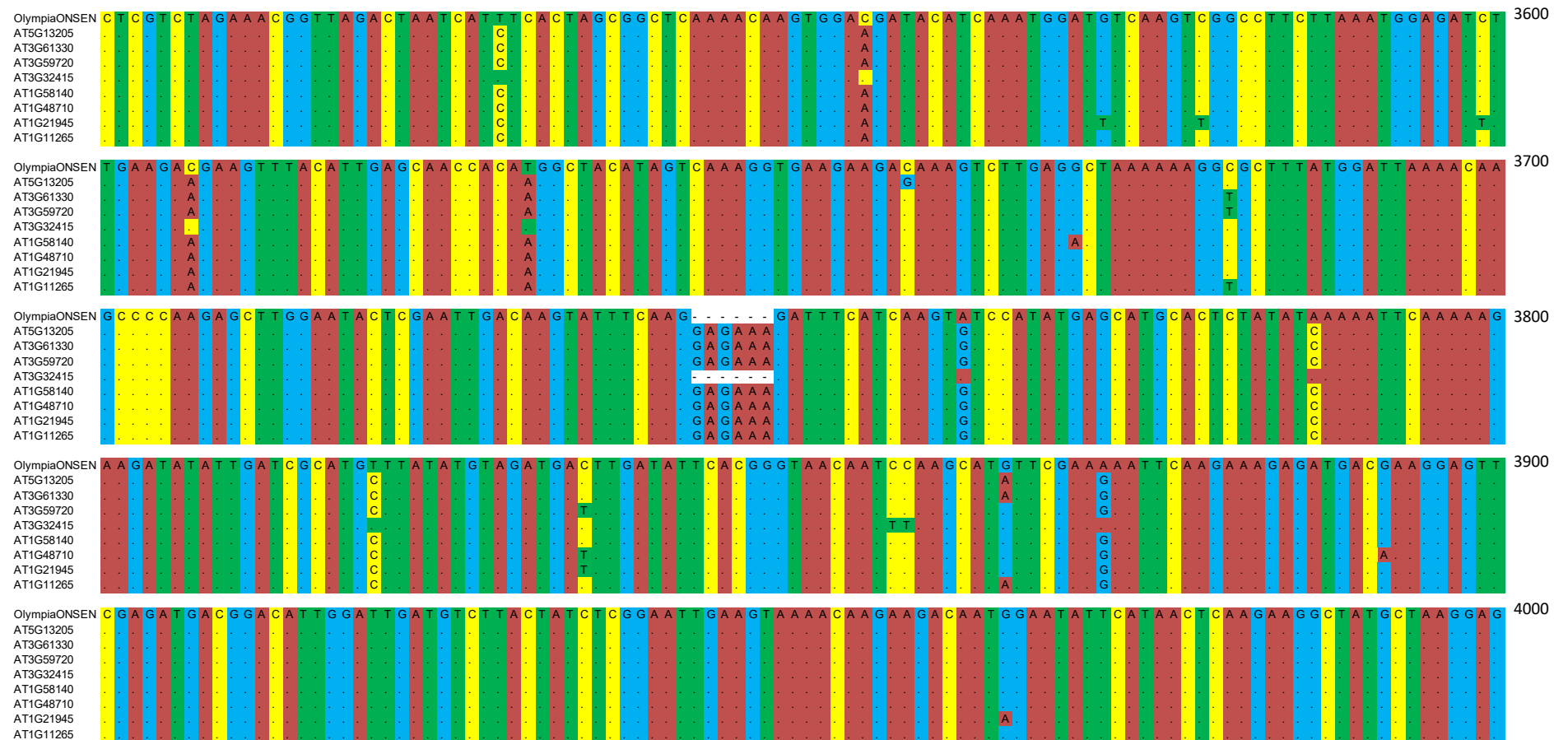


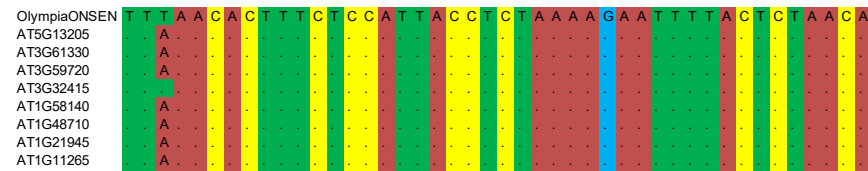
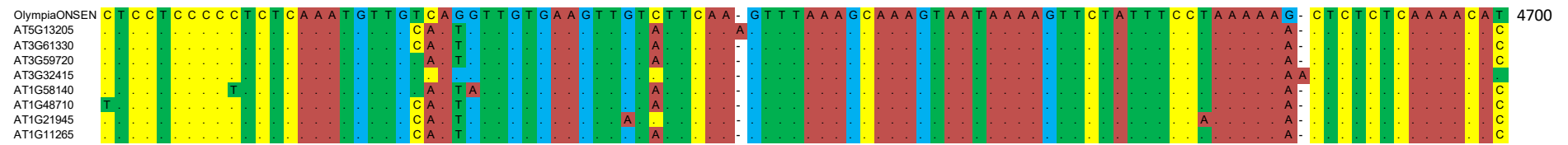
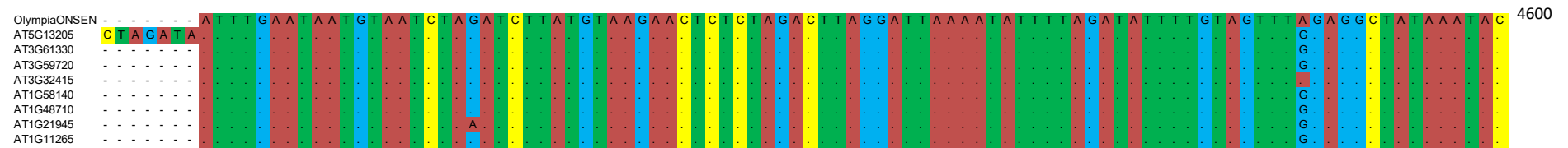












Supplementary Fig. 1. Multiple sequence alignment of full-length *ONSENs*. The gene IDs refer to the eight *ONSEN* copies in Col-0. Olympia-1 represents the sequence of an *ONSEN* copy in the natural accession Olympia-1. Nucleotides identical to those in Olympia-1 *ONSEN* are indicated as dots.