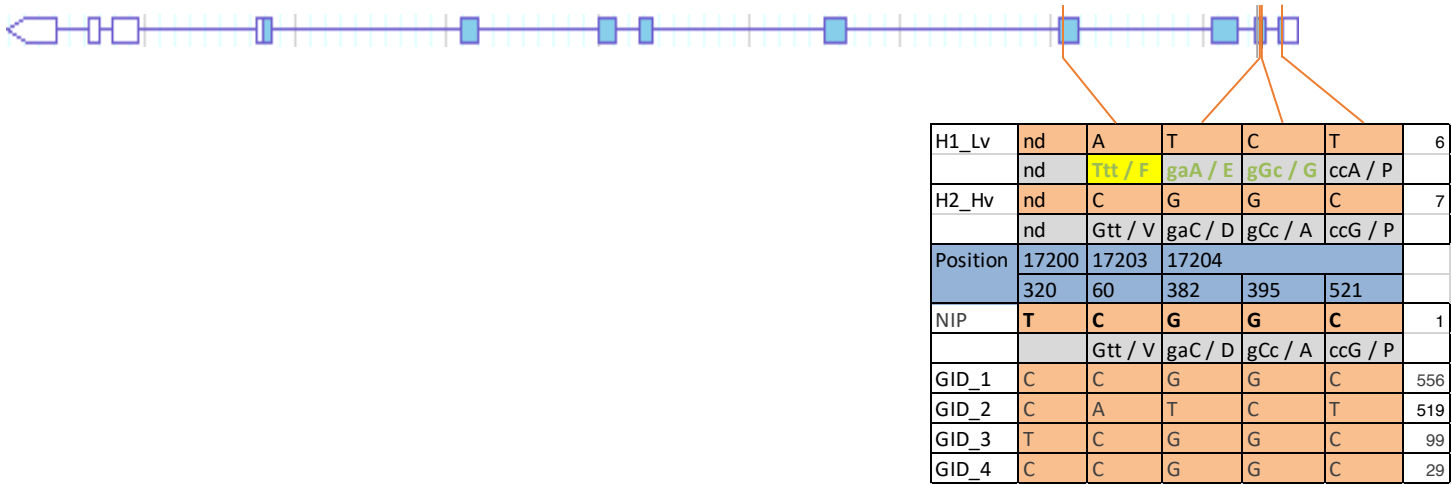
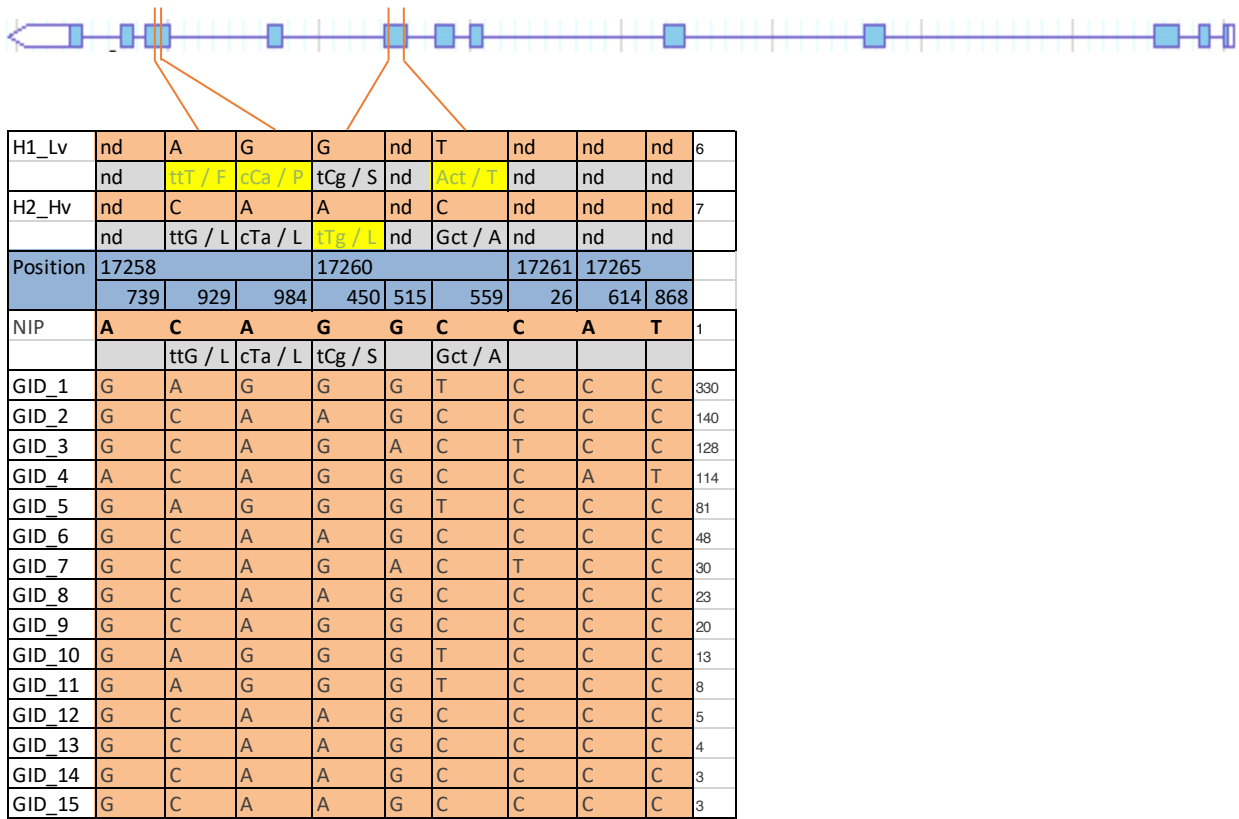


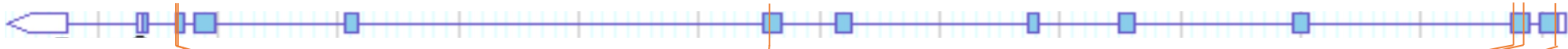
LOC_Os02g29040



LOC_Os02g29140

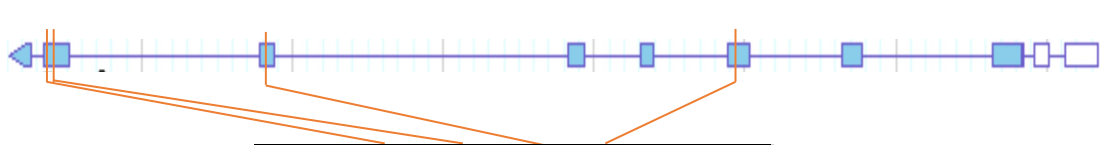


LOC_Os02g29160



H1_Lv	G	G	nd	nd	nd	C	nd	nd	nd	C	C	nd	nd	T	6
	Ctc / L	Ctc / L	nd	nd	nd	Gta / V	nd	nd	nd	gaG / E	Ggc / G	nd	nd	cAc / H	
H2_Hv	A	A	nd	nd	nd	G	nd	nd	nd	G	A	nd	nd	C	7
	Ttc / F	Ttc / F	nd	nd	nd	Cta / L	nd	nd	nd	gaC / D	Tgc / C	nd	nd	cGc / R	
Position	17278					17280				17283					
	694	706	822	98	123	621	660	987	11	772	843	18	39	97	
NIP	A	G	A	G	A	G	A	A	G	C	A	G	C	C	1
	Ttc / F	Ctc / L				Cta / L				gaG / E	Tgc / C			cGc / R	
GID_1	A	G	A	G	A	G	A	A	G	C	A	G	C	C	1647
GID_2	A	G	A	A	A	G	A	G	G	C	A	G	A	C	63
GID_3	G	G	A	A	A	C	A	G	G	C	C	C	C	T	24
GID_4	A	G	A	G	c	G	A	A	G	C	A	G	C	C	21
GID_5	A	A	A	A	A	G	A	G	G	G	A	G	A	C	21
GID_6	A	G	A	A	A	G	G	G	A	C	A	G	A	C	15
GID_7	A	G	c	A	A	G	A	G	G	C	A	G	A	C	9
GID_8	A	A	A	A	A	G	A	G	G	C	A	G	A	C	7
GID_9	A	G	A	A	A	G	G	G	A	C	A	G	A	C	6

LOC_Os02g29190



H1_Lv	C	C	C	T	nd	7
	cGg/ R	cgG/ R	gaG/ E	gAg/ E	nd	
H2_Hv	T	T	T	C	nd	6
	cAg/ Q	cgA/ R	gaA/ E	gGg/ G	nd	
Position	17309	17310	17313	17314		
	380	406	820	930	672	
NIP	C	C	C	C	G	1
	cGg/ R	cgG/ R	gaG/ E	gGg/ G		
GID_1	T	T	T	C	G	411
GID_2	T	T	C	C	G	409
GID_3	T	T	C	C	G	251
GID_4	C	C	C	T	G	219
GID_5	C	C	C	C	G	118
GID_6	T	T	C	C	T	34
GID_7	C	C	C	T	G	13
GID_8	T	T	T	C	G	3
GID_9	T	T	t	C	G	3
GID_10	T	T	C	C	G	3
GID_11	T	T	T	C	G	3

LOC_Os02g29210



H1_Lv	T	C	nd	nd	T	nd	nd	nd	nd	nd	T	nd	7
	gaA/ E	aGc/ S	nd	nd	Aat/ N	nd	nd	nd	nd	nd	cat/ N	nd	
H2_Hv	C	T	nd	nd	C	nd	nd	nd	nd	nd	G	nd	6
	gaG/ E	aAc/ N	nd	nd	Gat/ D	nd	nd	nd	nd	nd	aCt/ T	nd	
Position	17328				17331	17333	17334				17335		
	82	89	99	110	926	979	575	422	456	481	532	549	
NIP	C	C	C	G	C	G	A	G	A	G	G		1
	gaG/ E	aGc/ S			Gat/ D						aCt/ T		
GID_1	C	T	T	T	C	C	A	A	G	A	G	G	321
GID_2	C	C	T	T	C	C	A	A	G	A	G	G	264
GID_3	T	C	T	T	T	C	A	A	G	A	T	G	194
GID_4	C	C	T	T	C	T	A	C	T	G	G	G	184
GID_5	C	C	C	G	C	C	G	A	G	A	G	G	97
GID_6	C	C	T	T	C	T	A	C	T	G	G	A	15
GID_7	C	C	T	T	C	T	A	C	T	G	G	G	10
GID_8	T	C	T	T	T	C	A	A	G	A	T	G	7
GID_9	T	C	T	T	T	C	A	A	G	A	T	G	4