

Additional file 5. Average Ct values measured for the four T-cell stimulatory epitopes involved in CD and for the four most stable references genes in 11 spelt accessions and 3 diploid accessions representative of the ancestral genomes of spelt and bread wheat.

Species	Accession name	TaqMan probes targeting epitopes				TaqMan probes targeting reference genes			
		A20	A1	A9	A2	ARF	RLI	VAS	DUF52
<i>Triticum aestivum ssp. spelta</i>	BEL08	19.62	21.90	16.59	15.15	27.64	29.92	32.03	29.36
	DK01	18.88	21.80	16.44	15.37	26.70	29.55	31.49	28.60
	SPA03	18.30	20.47	15.80	14.68	28.01	29.89	31.32	30.16
	BUL04	21.05	24.14	18.76	17.99	29.27	32.54	34.32	31.36
	GER11	18.83	20.75	16.43	15.18	27.06	30.39	32.49	29.52
	GER12	19.60	21.19	17.08	15.91	27.82	30.34	31.96	29.61
	TAD06	19.85	22.67	17.77	16.16	27.27	29.68	31.10	28.86
	SWI23	18.68	21.23	16.58	15.32	27.25	29.76	31.66	28.94
	US06	19.74	18.91	17.37	15.91	29.00	31.58	33.21	30.77
	Iran77d	21.03	22.79	18.72	17.11	28.59	30.66	32.02	30.19
IRA03	19.19	21.28	17.03	15.38	27.36	30.14	31.58	29.03	
<i>Triticum urartu</i>	LB01	19.97	35.08	18.83		27.90	32.00	33.58	29.35
<i>Aegilops speltoides</i>	TR08		22.09			28.07	31.41	34.68	31.83
<i>Aegilops tauschii</i>	TR10	22.61	24.97	20.53	19.40	29.73	33.45	34.75	31.25