Erratum

Initiation and processing of *atp6*, *T-urf13* and *ORF221* transcripts from mitochondria of T cytoplasm maize

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The final sequence and the footnote to Table 1 should read as follows:

Table 1. Sequences proximal to primer extension bands

Band length		Downstream site		Transcript (kb)		Sequence
20-mer	18-mer	<i>atp6</i>	<i>T-urf13</i>	<i>atp6</i>	<i>urf13/ORF221</i>	
480	581 ± 5	–807	–432	1.8	2.0	TATATAAGCGTAGTTAT
470	571 ± 5	–797	–422	1.8	2.0	TATGTTAAGTTCTCT
391		–718		?		
297	398 ± 3	–624	–249	1.6	1.85	CATAGAGAAAGATGTTCTGATTCAGTT
290	391 ± 3	–617	–242	1.6	1.85	
279	380 ± 3	–606	–231	1.6	1.85	
241	342	–568	–193	1.55	1.8	AGACGGGGCCCTTAGCTTAGGGA
250	351	–577	–202	1.55	1.8	
252	353	–579	–204	1.55	1.8	
254	—	–581	—	1.55	—	
195		–522		?		TGGTAGCCTCTCTGCTG
123	225	–450	–76	?	?	TGTGTGGGTGTTTCAGT
	141		+10		1.6	ATCACTACTTTCTTAAAC
	136		+14		1.6	
	107		+43		1.5	TTTGATCAAGGTTTGGTAT

↓ denotes mapped transcript termini. Italicized sequences are not in N mitochondrial DNAs; mapped termini only detected in T cytoplasm in this region. 20-mer specific to *atp6*; 18-mer specific to *T-urf13/ORF221*