

SUPPLEMENTARY Table 2 :
Primers used for the expression analysis of *AtESL* par qRT-PCR.

Genes	Sequences	Tm (°C)	Length (nt)	Taille de l'amplicon (bp)	Efficacité	Observations
<i>At1g19450</i>	F : TAAGCCTCGTGATTGTAGCA	58	20	121	90	-
	R : CAAGAAATAACCATAGCCACA	58	21			
<i>At1g75220</i>	F : GTGGAGGAACTTTCACCTGTGA	64	22	99	90	-
	R : GTTCTTCAAGAGTTTTCGCCTT	58	21			
<i>At2g48020</i>	F : CTGAGATATTTCCCATAAACAT	58	21	139	93	-
	R : AAGATTGCAATTTATGTCCCAG	58	21			
<i>At3g05150</i>	F : TTCTTTTGAAGGCTCATGGTT	58	21	200	-	-
	R : GACAAGCCATGAACTTAGCCA	62	19			
<i>At5g18840</i>	F : CTGAGATATTTCCGATAAACGTA	62	23	177	114	-
	R : TCTTCGCCACAAATATTATCGTC	64	23			
<i>At1g54730</i>	F : AATCCGGCAGGAACGTTTATATG	64	22	179	100	-
	R : CGTTATAATTAGGGACTGATAT	58	22			
<i>At4g04750</i>	F : TACAGGAACGTTTATGATGTTT	58	22	203	90	-
	R : CATGAAATATAAAGAGATCTGA	56	22			
<i>At4g04760</i>	F : CAGAGATTATCCAGTAGATGT	60	22	174	100	-
	R : TGGCTATGAACACAAATCCAA	58	21			
<i>At3g05400</i>	F : CTTCTTATGACTTCAGCGTTGG	64	22	148	100	-
	R : CAATTGCGAACATCATCGT	54	19			
<i>At1g08890</i>	F : TGGAGTGCTTTCGGAACATATT	60	22	150	109	-
	R : GAGTAAAGAATCACTTTGGAT	58	22			
<i>At1g08900</i>	F : GTAGCCTTGTAACGATGTCA	58	20	99	97	-
	R : ATATGAAATATGTTCCGGAAGC	60	22			
<i>At3g20460</i>	F : ACGCTATGCAATTTAACTAGCTG	64	23	89	97	-
	R : AACACACCAGAAGAGCTCCACT	66	22			
<i>At1g08920</i>	F : TCGGTCTCATTCCATGTGCG	62	20	131	97	-
	R : TGCATCATCTCCGCGAAGAC	62	20			
<i>At1g08930</i>	F : ATAATGGCTGAGATATTTCCG	58	21	176	95	-
	R : ATAAATACGATCGAACTGGC	60	20			
<i>At3g05155</i>	F : GTGGTACCAAAAGCTATATTGG	62	22	164	-	No amplification
	R : CCGATGCCTGCAAATTTGTACA	64	22			
<i>At3g05160</i>	F : GCACAAGGAACCTTTTACATAT	60	22	101	90	-
	R : AATTCTTCAAGCGACTGTCC	58	20			
<i>At3g05165</i>	F : AAGGAACATTTTACATATTTGC	56	22	167	-	Not enough specificity
	R : TTTATTATTGCAATGCGACT	54	21			
<i>At5g27350</i>	F : TATGTCGGAGATATTTCCAATA	58	22	269	97	-
	R : CAAATTAAGTTTGATTCTTTTC	54	22			
<i>At5g27360</i>	F : AGTGGAGCACTCAAGGAACATT	64	22	149	95	-
	R : TTGATTATTCTATCGGGCTC	60	22			