

Locus	Classification	Direction	Sequence(5'-3')
At2g23170	GH3.3	Forward	TCCACTAAGGACGTGAAGGCTCTAAG
		Reverse	TGCTGGTAATCCACCGGGAGTCTTCG
At1g59500	GH3.4	Forward	ATGGCTGTTGATTTCGCTTCTTCAATC
		Reverse	CGACTTCACAAATAAGAAGTATAAAC
At4g27260	GH3.5	Forward	AGCCCTAACGAGACCATCCT
		Reverse	AAGCCATGGATGGTATGAGC
At4g36110	SAUR9	Forward	GACGTGCCAAAAGGTCACTT
		Reverse	AGTGAGACCCATCTCGTGCT
At4g38850	SAUR15	Forward	ATGGCTTTTTTGGAGGAGTTTCTTGGG
		Reverse	TCATTGTATCTGAGATGTGACTGTG
At5g18060	SAUR23	Forward	ATGGCTTTGGTGAGAAGTCTATTGGT
		Reverse	TCAATGGAGCCGAGAAGTCACATTGA
At1g29430	SAUR62	Forward	CATTTCGATTCCGGCCTTTTTTA
		Reverse	TGGAAGGCATTTTTGAATTTTT
At1g29490	SAUR66	Forward	CACAAAGAACTCATGAAGATGG
		Reverse	GAATCGAATCGAATGGCAAC
At4g14560	IAA1	Forward	ATGGAAGTCACCAATGGGCTTAACCT
		Reverse	TCATAAGGCAGTAGGAGCTTCGGATC
At1g15580	IAA5	Forward	GGAACATCTCCAGCAAGCAT
		Reverse	GGGCTTCTCCGGAAGTAGAG
At1g52830	IAA6	Forward	GGAAGATTGACCTTGGTTTCG
		Reverse	TTCACGATCCTCAGCCTCTT
At1g04250	IAA17	Forward	GGTACC ATGATGGGCAGTGTCCGAG
		Reverse	CCATGG AGCTCTGCTCTTGCACCTT
At3g15540	IAA19	Forward	ATGGAGAAGGAAGGACTCGGGCTTG
		Reverse	GTCTTCGTATATGGTAACGTATTCCG
At4g32280	IAA29	Forward	AAGATGGATGGTGTGGCAAT
		Reverse	ATCCCCTCGAAGTAGCCAGT
At5g47370	HAT2	Forward	ATGATGATGGGCAAAGAAGATCTAG
		Reverse	TCCACGTGAGTACCCTCGATCCGGAG
At3g58190	LBD29	Forward	CAACAAGCAGCACAAAGCAT
		Reverse	CTGAGGAGGTTTTCGTTGTGG
At1g78100	F-box	Forward	GATGCGATTCCAGATCCTGT
		Reverse	TCGGGTTAGGATTTTCGTCAG
At5g02760	PP2C6	Forward	GGCCGCTTACATTCAACAAT
		Reverse	TGCATATCACTCCTGCCAAG
At2g26710	BAS1	Forward	TTCAGGCAAAGAATGTGACG
		Reverse	GCCACATGGGATTTTTGTACC
At5g62280	DUF1442	Forward	AACGTGCGCGAATTCATATC
		Reverse	TCCATGACCCTCTCGAAAAC
At5g44200	CBP20	Forward	ACTGGCTCATTGGGAGCTTA
		Reverse	CATCTTCACCATCGTCATCG
AT1g20010	TUB5	Forward	AACCATTTGATCTCCGCAAC
		Reverse	TGGAGTCCCACATTTGTTGA
At5g66240	WD-40 repeat	Forward	CCGTTGGTGGTGTATCTTTCT
		Reverse	TGTTTTCCGCTCCTTACACC
	DR5:GUS	Forward	GCATTCAGTCTGGATCGCGAAAAC TG
		Reverse	ATTACGCTGCGATGGATTCCGGCATA