

**Table S8.** Genes list corresponding to the GO enrichment analysis by the AgriGO program.

GO_acc	term_type	Term	query item	query total	bgitem	bgtotal	pvalue	FDR
GO:0006869	P	lipid transport	47	53	104	39203	#####	#####
GO:0010876	P	lipid localization	47	53	104	39203	#####	#####
GO:0033036	P	macromolecule localization	47	53	997	39203	6.50E-68	1.50E-66
GO:0051234	P	establishment of localization	47	53	3841	39203	6.20E-41	8.40E-40
GO:0006810	P	transport	47	53	3841	39203	6.20E-41	8.40E-40
GO:0051179	P	localization	47	53	3887	39203	1.10E-40	1.20E-39
GO:0048871	P	multicellular organismal homeostasis	12	53	1758	39203	3.20E-06	1.90E-05
GO:0042309	P	homiothermy	12	53	1758	39203	3.20E-06	1.90E-05
GO:0050826	P	response to freezing	12	53	1758	39203	3.20E-06	1.90E-05
GO:0009409	P	response to cold	12	53	1758	39203	3.20E-06	1.90E-05
GO:0001659	P	temperature homeostasis	12	53	1758	39203	3.20E-06	1.90E-05
GO:0009266	P	response to temperature stimulus	12	53	1765	39203	3.40E-06	1.90E-05
GO:0009628	P	response to abiotic stimulus	12	53	1856	39203	5.60E-06	2.90E-05
GO:0032501	P	multicellular organismal process	12	53	1983	39203	1.10E-05	5.30E-05
GO:0042592	P	homeostatic process	12	53	2047	39203	1.50E-05	6.80E-05
GO:0065008	P	regulation of biological quality	12	53	2139	39203	2.30E-05	9.80E-05
GO:0006950	P	response to stress	12	53	3059	39203	0.00066	0.0026
GO:0050896	P	response to stimulus	12	53	3421	39203	0.0018	0.0066
GO:0065007	P	biological regulation	12	53	6272	39203	0.13	0.46
GO:0008289	F	lipid binding	47	53	202	39203	1.10E-98	9.40E-98
GO:0005488	F	binding	52	53	24279	39203	3.20E-10	1.30E-09
GO:0050825	F	ice binding	12	53	1758	39203	3.20E-06	6.80E-06
GO:0050824	F	water binding	12	53	1758	39203	3.20E-06	6.80E-06

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## entries