Table S2. Unigene classification according to their GO "biological process" term assigned to the most similar *Arabidopsis* protein

Biological process	Induced unigenes	Repressed unigenes
	I6; I13; I15; I16; I17; I19;	R5; R6; R8; R12; R13; R19;
Unknown	I20; I21; I29; I38; I47; I50;	R21; R27; R31; R32; R33;
	I56; I59	R34; R37; R39; R41; R44
Protein modification	I9; I26; I27; I31; I36; I49	R4; R9; R11; R17; R23; R42
Transcription and RNA	I1; I2; I24; I44; I52; I55;	D7. D14. D16. D20
metabolism	I57;	R7; R14; R16; R38
Translation	I3; I21; I35	R10; R15
Carbohydrate	120, 122, 140	D20: D20
metabolic process	I30; I32; I40	R20; R28
Vesicle-mediated	142	D20 D24 D25
transport	I43	R30; R34; R35
Metabolic process	I4; I11; I58	R25
DNA structure	I22; I48	R22
Biosynthetic process	I8	R2
Defense response	I53	R3
Response to cold,	I12; I34; I39; I42; I46; I51;	
feezing and heat	I60	none
Lipid formation and	I37; I41; I45	none
metabolism		
Response to auxin	70 75 700	
stimulus	12; 15; 123	none
Electron transport	110 114	
chain	I10; I14	none
Tricarboxylic acid	110	
cycle	I18	none
Photosynthesis	I28	none
Response to Pi	T.C. 4	
starvation	I54	none
H ₂ O ₂ catabolic process	I7	none
Fruit development	I33	none
Cell death	I25	none
Cell growth	none	R1; R29
Response to		,
carbohydrate stimulus	none	R39
Response to nitrate	none	R18
Response to cadmium		
ion	none	R26
Signalling	none	R24