Arabidopsis sterol carrier protein-2 is required for normal development of seeds and seedlings.

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## **Supplementary material**

Table S1. Ranking of genes with significantly altered expression in *Atscp2-1* relative to wt.

Consa	Description <sup>b</sup>	۸ <sup>C</sup>	R statistic
<u>A+2~20720</u>	Expressed protein	11/1	$\frac{D-SIGHISHC}{14.20}$
AlogoU/20	Expressed protein	4.30	14.20
ALSG14210	myrosinase-associated protein	-4.03	13.28
At1g21400	2-oxoisovalerate denydrogenase	3.04	12.79
At3g4/340	Glu-dependent Asn synthetase	3.05	12.65
(ASNI)		2.02	10.07
At1g10090	Expressed protein	2.92	12.37
At3g45140	lipoxygenase	-2.80	12.22
(LOX2)			
At4g01870	tolB protein-related	3.38	12.01
At3g24420	hydrolase, alpha/beta fold	2.84	12.01
At1g52060	jacalin lectin	-3.72	11.89
At5g04960	pectinesterase family protein	-2.40	11.68
At2g18193	AAA-type ATPase	-2.51	11.59
At2g47770	benzodiazepine receptor-related	2.69	11.37
At4g30650	hydrophobic protein	-2.25	11.05
At3g15670	late embryogenesis abundant (LEA) protein	2.67	10.97
At1g48130	peroxiredoxin	2.45	10.85
(PER1)			
At5g20230	plastocyanin-like	2.60	10.82
At1g11260	glucose transporter	2.70	10.77
(STP1)			
At1g79360	transporter-related	2.29	10.75
At3g21720	isocitrate lyase	2.16	10.64
At5g50600	short-chain dehydrogenase/reductase (SDR)	2.71	10.63
C	family protein		
At3g28300	integrin-like protein At14a	-1.98	10.31
At1g08630	Thr aldolase	3.58	10.06
(THA1)			
At2g35300	LEA group 1 protein	2.55	9.98
At4g368500	PQ-loop repeat family protein	2.19	9.93
At5g14470	GHMP kinase-related protein	2.95	9.91
At1g06570	4-hydroxyphenylpyruvate dioxygenase	2.07	9.84
(PDS1)	, ,, ,, ,, ,, ,, ,, ,, ,, ,, ,, ,, ,, ,		-
At1g05240	peroxidase	-1.91	9.82
At4g15530	pyruvate.orthophosphate dikinase	2.34	9.77
At5g53970	aminotransferase	1.89	9.70
At5g54010	glycosyltransferase family protein	2.10	9.68

At4g39940	adenosine-5'-phosphosulfate-kinase	-1.84	9.60
(AKN2)			
At1g52050	jacalin lectin	-2.77	9.55
At3g48940	remorin	-2.48	9.53
At4g35770	senescence-associated protein	1.95	9.55
(SEN1)	1		
At5g55730	fasciclin-like arabinogalactan-protein	-1.81	9.44
(FLA1)			
At3g20340	expressed protein	1.77	9.45
At3g28720	expressed protein	2.16	9.38
At1g73120	expressed protein	1.77	9.30
At2g20520	fasciclin-like arabinogalactan-protein	-2.49	9.27
(FLA6)	raserenn nite araemogaraetan protein	2.19	, <b>.</b>
At1954100	aldehyde dehydrogenase	1 69	9 17
(ALDH7R4)	and only do doing drogonaso	1.07	<i></i>
$At5\sigma64570$	beta-xylosidase	1 94	916
(XYI4)	beta Aylosidase	1.94	7.10
$\Delta t/a25820$	vyloglucan endotransglycosylase	-1.92	9.04
(YTR0)	xylogidean endotransgryeosylase	-1.72	7.04
$(\Lambda I \Lambda \mathcal{F})$	ABA responsive protein AtHVA22h	2.02	8 0 8
A13g02490 (A+ $UVA22b$ )	ABA-responsive protein AttivA220	2.02	0.90
(AIIIVA220) At1 $a52400$	alvoosul hydroloso family 1 protain	2 75	8 02
At1g32400	bete vulosidese	-2.75	0.92
(DVI 1)	beta-xylosidase	1.67	0.09
(BALI)	multi compan avidada tuma I mutain	1 70	0 07
At4g5/100	multi-copper oxidase type I protein	-1.79	ð.ð/ 0.07
At4g25580	stress-responsive protein	2.10	8.8/
At3g2/630	expressed protein	-2.16	8.80
At3g15280	expressed protein	1.89	8./1
At3g07700	ABC1 family protein	1.64	8.60
At2g05540	glycine-rich protein	1.63	8.58
At2g43820	UDP-glucoronosyl/UDP-glucosyl transferase	2.22	8.57
	family		
At3g49960	peroxidase	-2.05	8.47
At5g10130	pollen Ole e 1 allergen and extensin family	-1.66	8.47
	protein		
At1g15330	CBS domain-containing protein	2.63	8.46
At5g14180	lipase	1.79	8.42
At1g15380	lactoylglutathione lyase family protein	1.63	8.41
At5g05500	pollen Ole e 1 allergen and extensin family	-1.56	8.41
	protein		
At1g73330	protease inhibitor	-2.98	8.35
(ATDR4)	-		
At1g72680	cinnamyl-alcohol dehydrogenase	1.53	8.25
At1g33110	MATE efflux family protein	1.50	8.20
At5g54370	LEA protein	-2.98	8.14
At5g38200	expressed protein	1.63	8.06
At3g13750	beta-galactosidase	1.71	8.05
At1g05240	peroxidase ATP11a	-1.92	8.04
At4g15760	monooxvgenase	1.50	8.00
At1g29660	GDSL-motif lipase/hydrolase family	-1.73	7.99
U		-	

At3g14990	4-methyl-5(b-hydroxyethyl)-thiazole	1.71	7.95
	monophosphate biosynthesis protein		
At4g02270	pollen Ole e 1 allergen and extensin family	-1.96	7.88
	protein		
At3g26960	expressed protein	-1.89	7.88
At3g06850	branched chain alpha-keto acid	1.59	7.83
( <i>BCE2</i> )	dehydrogenase E2 subunit (din3)		
At3g07800	thymidine kinase	-1.55	7.78
At4g40090	arabinogalactan-protein	1.70	7.71
(AGP3)			
At5g05410	DREB2A is a member of the DREB	1.40	7.70
(DREB2A)	subfamily A-2 of ERF/AP2 transcription		
	factor family		
At2g33790	pollen Ole e 1 allergen protein	-2.36	7.70
At4g26490	hypothetical protein	-1.65	7.68
At2g44080	expressed protein	1.45	7.61
At4g02850	phenazine biosynthesis PhzC/PhzF family	-1.49	7.59
At2g37770	aldo/keto reductase family protein	1.79	7.56
At3g15620	6-4 photolyase	1.45	7.50
(UVR3)			
At1g61580	60S ribosomal protein L3	-1.48	7.50
(ARP1)			
At3g14900	expressed protein	-1.60	7.49
At1g48600	phosphoethanolamine N-methyltransferase 2	-1.38	7.46
At3g30775	proline oxidase	2.10	7.42
At2g31980	cysteine proteinase inhibitor-related	1.82	7.41
At5g60660	major intrinsic family protein	-1.78	7.35
At5g02440	expressed protein	1.39	7.35
At1g22880	glycosyl hydrolase family 9 protein	-1.43	7.32
At5g56750	Ndr family protein	1.45	7.30
At2g21880	Ras-related GTP-binding protein	-1.46	7.27
At4g39950	cytochrome P450 79B2	-1.49	7.21
( <i>CYP79B2</i> )			

<sup>a</sup> AtSCP2 was excluded from the ranking list.
<sup>b</sup> Descriptions are according to The Arabidopsis Information Resource (TAIR) at www.arabidopsis.org.
<sup>c</sup> M=log2 differential expression relative to Col-0.

Gene	<i>Description</i> <sup>a</sup>	<i>Microarray<sup>b</sup></i>	$qRT$ - $PCR^{b}$
At3g14210	myrosinase-associated protein	-	-
At1g21400	2-oxoisovalerate dehydrogenase	+	+
At3g47340	Glu-dependent Asn synthetase	+	+
(ASN1)			
At3g45140	lipoxygenase	-	-
(LOX2)			
At5g04960	pectinesterase family protein	-	-
At3g21720	isocitrate lyase	+	+
At1g08630	Thr aldolase	+	+
At1g06570	4-hydroxyphenylpyruvate	+	+
(PDS1)	dioxygenase		
At1g52050	jacalin lectin	-	-
At5g55730 (FLA1)	fasciclin-like arabinogalactan-protein	-	-

Table S2. Comparison of results from microarray and quantitative real-time reverse transcriptasePCR (qRT-PCR) analysis of gene expression in wt and *Atscp2-1* seedlings.

<sup>a</sup> Descriptions are according to The Arabidopsis Information Resource (TAIR) at www.arabidopsis.org

<sup>b</sup> Relative to the expression levels in A. thaliana Col-0, + indicates that the gene is upregulated in *Atscp2-1*, while - indicates that the gene is down-regulated in *Atscp2-1*.