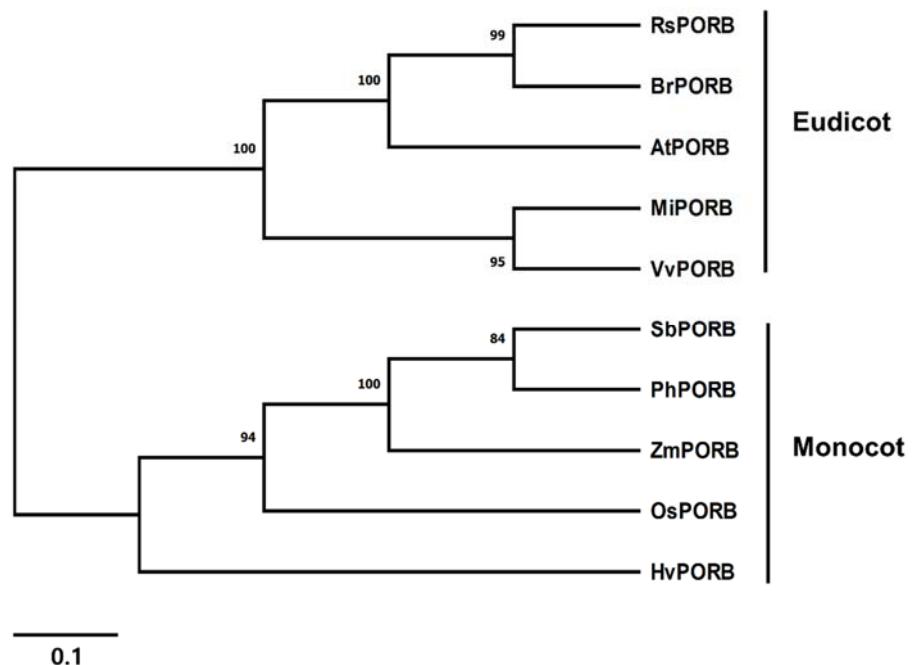


**Table S1.** Gene specific primers used in this study

Usage	Primer name	Primer sequence (5'-3')
Gene cloning	RsPORB-promoter-F	AATGTTCATGGACAATCCTCCTGATTGT
	RsPORB-promoter-R	TTGAAGGCCATTTGGACAAACAGAGA
	RsPORB-F	ATGGCCCTCAAGCCGCTTGGTC
	RsPORB-R	TTAGGCCAAGCCAACCAGCTTCAC
Gene expression	RsPORA-F	ACGGTGAAAAGGCATATAAA
	RsPORA-R	GCTCTCTGAACAAACCTGTC
	RsPORB-F	GACAGAGTCTGGCAAAAGAC
	RsPORB-R	TGATCTCCCAAACTTACG
	RsPORC-F	CAGAGAACACATACCCCTGT
	RsPORC-R	CACCTGTGCTAGTCTTTCC
	RsRPII-F	ATCACGCTAAATGGTCTCCT
	RsRPII-R	GCTGCTCTCAATCAAGTCAATC
Virus-induced gene silencing	pTRV2-RsPORB-F	CGACGACAAGACCCTAAAGATGTAAAAGTGA
	pTRV2-RsPORB-R	GAGGAGAACAGCCCTAGCTTCGGTGGTACGT
	pTRV2-RsPORs-F	CGACGACAAGACCCTAAATGGCACGTGAT
	pTRV2-RsPORs-R	GAGGAGAACAGCCCTACATGTTGCACACTT
Promoter activation assay	RsPORB-promoter-F	AATGTTCATGGACAATCCTCCTGATTGT
	RsPORB-dP1-R	GTTGGATAATTCTTGGTTATAATTNTTA
	RsPORB-dP1-F	TAAAAATTATAAACCAAGAAAATTATCCAAAC
	RsPORB-dP2-R	CTTGGCATGGAGAAGAAGAAGATATCCTACAA
	RsPORB-dP2-F	TTGTAGGATATCTTCTTCTCCCAGCCAAAG
	RsPORB-promoter-R	TTGAAGGCCATTTGGACAAACAGAGA
	pUC-RsPORB-fLUC-F	AGGCTCTAGAGGATCCAATGTCATGGACAATCC
	pUC-RsPORB-fLUC-R	TTGGCGTCTTCATGGTTGGACAAACAGA



**Figure S1. Phylogenetic relationships among PORB proteins from radish and other plants.** The phylogenetic tree was constructed using the neighbor-joining method with MEGAX software. Numbers next to the nodes are bootstrap values from 1000 replications.

**A***proRsPORB-G1*

-1000 AATGTTCATG GACAATCCTC CTGATTGTCA AACTCTTCGT  
 -960 AAGGGTGTCT TCGCTGCAGG TGCCCTCTTT GTCTTCTTCA ACGCCATTGT TTCTCAGTTC  
 -900 TATTATTTCT TCTACTCCTC TGCTGCTGCC GCCTCCCTCT CGCCTTACTA GAGGATTGAGA  
 -840 ACCAATAAAA ATAAGTTGT TTATTTCTTA TGTTATGT GATATCATA TAATGTCTA  
 -780 TCTTGGGCCT GTTGCTTGTAA AAAGTATATG ACTGCAAAT TACTTGAAGT TGAGGGTACG  
 -720 ACTATGAGTT TTGTTAGATG GATTGTATG GATATTGAAT TGAAATGTT TTCTTAAAAA  
 -660 GCTGATCTAA TGTGGTTTTT ACCGAACAAA AAAAACCCAG TGTGCTAGAC TGCTGTGCTA  
 -600 TATGTTATAT ATTGTTAAT TCAACATG TTAGAAGTT TATATATGCA ACTAATAGTT  
 -540 GTAAGATGAT GGTAGAGCAT GAACAATATG ATAAATAATA TATGCTTTA TTACCTTAAA  
 -480 TTTGAAAAT TGCTTGTAA ATTCAGTTA CACTTTAAA TTACTTCGAA TTATATACAC  
 -420 AAAAATTATA AACCAAGAAA ATTATCCAAA CATATCAGAG TAACCATGAC GAATAGTTG  
 -360 TAAGATATCT TCTTCTCCC TGCCAAAGT TGGAATCCCA TGAACACCAA TACCATCCT  
 -300 TTATGCAAT GTTACCAAAA CAGCCAATAA AATAAAATAT CTACTGAGAG GAAAATCTAG  
 -240 CCCTCGAATT CTCATTGGAT ATCTAATGAT GATGTGGCAG ATACTTTCTT TAAGATAATG  
 -180 TTATCCACAC ATCTTCTCTG TGATTTCTCT GGGGACCACT CTCTCCCTG TCCGAACCTC  
 -120 TCACCTCCCTC CACTCCCTCG CGAGTATACA TAAACCGGTC TACACTCTCG CAAAATATCA  
 -60 ACTTCAGTAG TCTTCTCTT CATTGCTATT GCTATCTCTC ACTCTCTGTT TTGTCCAAA

CCAAT motif Dof CAACA motif

**B***proRsPORB-W1*

-1038 ATGTTCA TGGACAATCC  
 -1020 TCCTGATTGT CAAACTCTTC GTAAGGGTGT CTTCGCTGCG GGTGCCCTCT TTGCTTCTT  
 -960 CAACGCCATT GTTTCTCACT TCTATTATTT CTTCTACTCC TCTGCTGCTG CGCCTCCCT  
 -900 CTCGCCTTAC TAGAGGATTT GAACCAACAA AAATAAGTT GTTATTTCT TGTGTTATGT  
 -840 GATACCATAAC AAATGTCATA TTTTGGGCCT GTTGCTTGTAA AAAGTATATG ACTGCAAAT  
 -780 TACTTGAAGT TGAGGGTACG ACTATGAGTT TTGTCAGATG GATTGTATG GATATTGAAT  
 -720 TGAAATGTTG TTCTTAAGAA GCTGATCTAA TGTTGGTTTT ACCAACAAA AAAAACCTCAG  
 -660 TGTGCTAGAC TGCTATATGT TATTTGTT AATTCAACAC ATGTTACAAG TTTTATAAAT  
 -600 GCAACTAATA GTTGTAAAGAT GATGGCAGAG CATGAACAAAT ATGATAGATA ATAATATATG  
 -540 TCTTTATTAC CTTGAAAATA GAAAATTGCT TGTTAAATT CAGTTACACT TAAAATTAC  
 -480 TTTGAGTTAT ATGAATAAAA ATTATAAAACAAACAAATGG GAAAATTATC CAAACATATC  
 -420 AGACTAACCA TGACGAATAG TTTGAGGAT ATCTTCTGT CAAACAAAG AAAACAAAAAG  
 -360 AAACAAAT CTTCTCTCC CATGCCAAAG TATGGAATCC CATAAAACACC AATCACCATC  
 -300 CTTTATGCA ATGTTACCAA AACAGCCAAT AAAATAAAAT ATCTACTGAG AGGGAAATCT  
 -240 AGCCCTCGAA TTCTCATTGG ATATCTAATG ATGATGTGGC AGATACTTTC TTTAAGATAAA  
 -180 TGTTATCCAC ACATCTCTC TCTGATTTCT CTGGGGACCA CTCTCTCCCT TGTCCGAACCT  
 -120 TCACCTCCCTC CACTCCCTCA CGAGTATACA TAAACCGGTA CACACTCTCG CAAAATATC  
 -60 AACTTCAGTA GTCTTCTCTA TCGCTATTGC TATCTCTCTC ACACCTCTGTT TTGTCCAAA

CCAAT motif Dof CAACA motif

**C**

Symbols of cis-element: CCAAT Dof CAACA motif

**Figure S2. Nucleotide sequences of *RsPORB* promoters from radish cultivars G1 and W1.** Partial promoter of *RsPORB-G1* (A) and *RsPORB-G1* (B). Putative several *cis*-elements are indicated by different colors. The CCAAT motif is indicated by a yellow box, Dof motif by dark grey boxes, and CAACA motif by pink boxes. (C) Schematic of the pro*RsPORB-G1* and pro*RsPORB-W1* showing putative *cis*-elements involved in chlorophyll biosynthesis. The *cis*-elements are indicated by different symbols.