

Supplementary Materials: Carotenoid Cleavage Oxygenases from Microbes and Photosynthetic Organisms: Features and Functions

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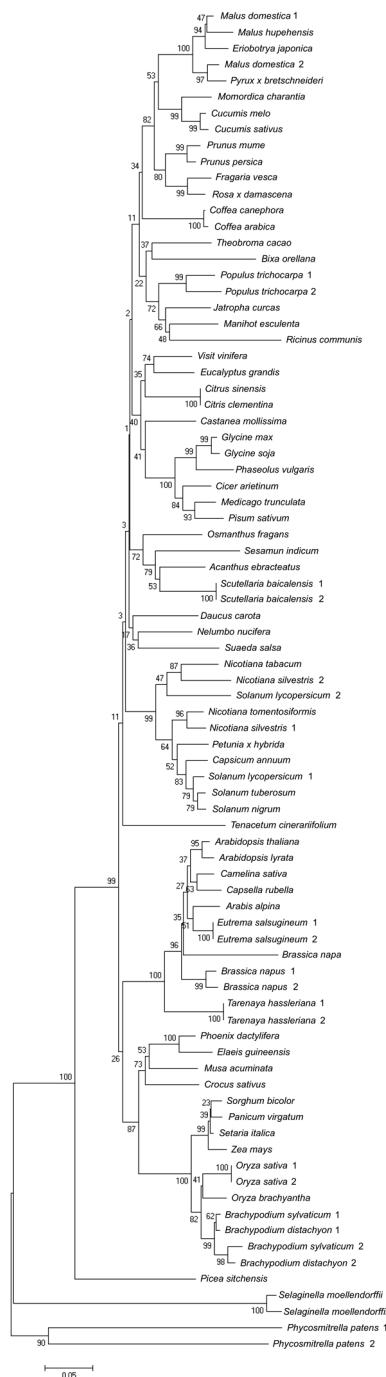


Figure S1. Phylogenetic tree of CCD1 enzymes. The analysed proteins are those presented in Table S1. Proteins were aligned using ClustalW and tree was generated using the Neighbor-Joining method. The optimal tree is represented and the percentages of replicated trees in which the associated proteins clustered together in the bootstrap test (2500 replicates) are shown next to the branches. The tree is drawn to scale with branch lengths in the same units as those of the evolutionary distances used to infer the phylogenetic tree. Analysis was conducted in MEGA6.

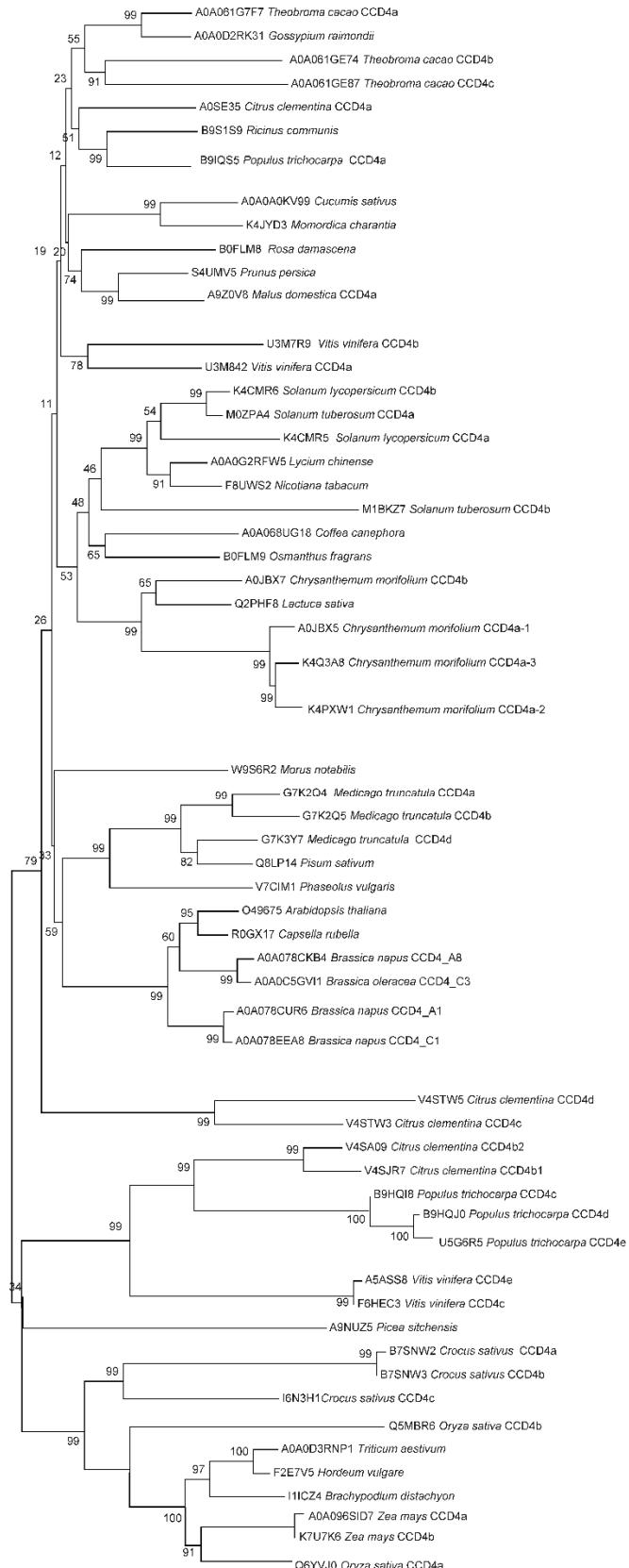


Figure S2. Phylogenetic tree of CCD4 enzymes. A total of 59 proteins were selected from the UniProtKB based on sequence similarity to *Arabidopsis* CCD4 and used for the analysis. Methods for proteins alignment and tree generation are described in the legend of Figure S1. UniProtKB accession numbers are indicated.

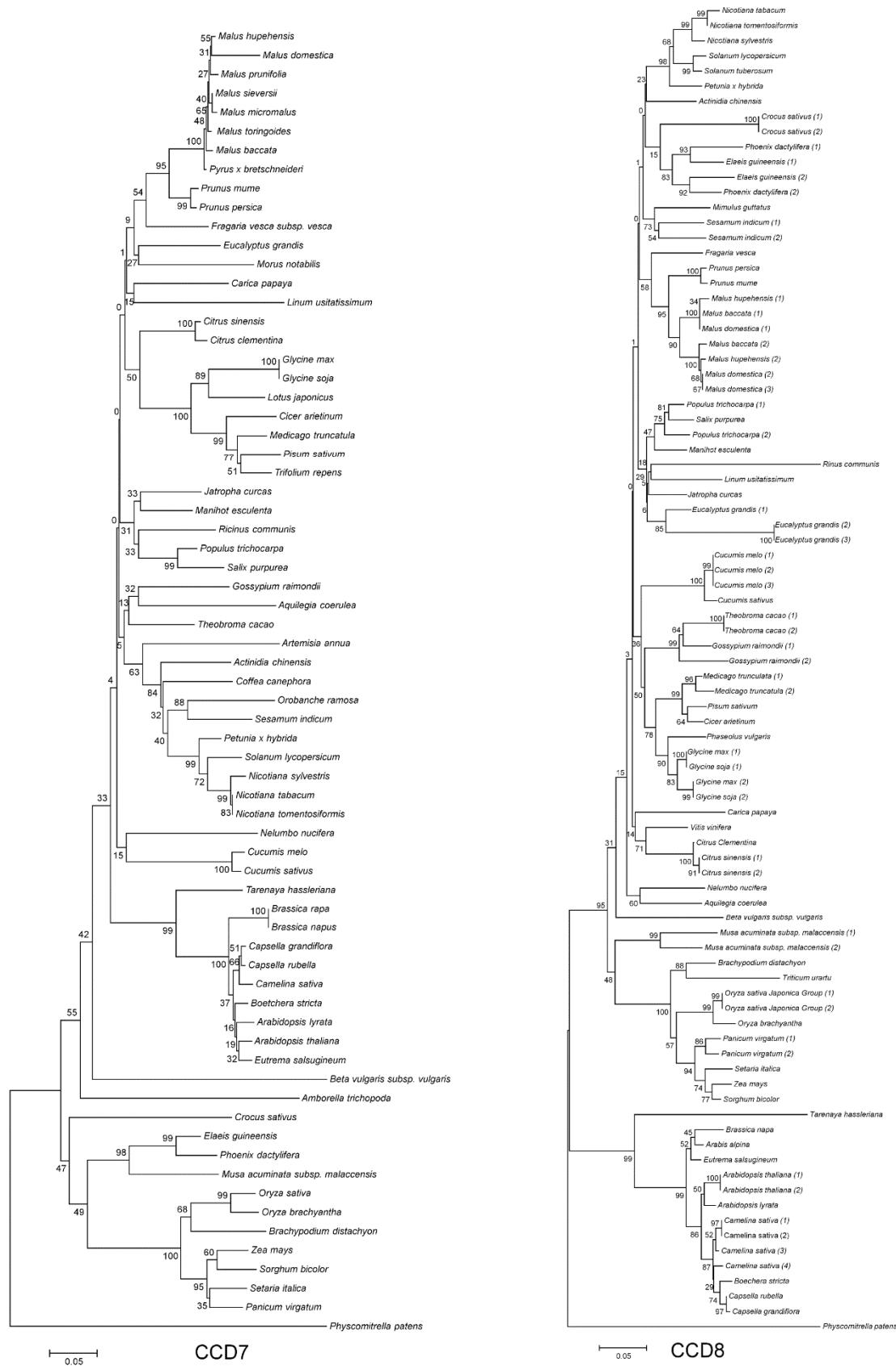


Figure S3. Phylogenetic tree of CCD7 and CCD8 enzymes. The analysed proteins are those presented in Tables S2 and S3. Methods for proteins alignment and tree generation are described in the legend of Figure S1.

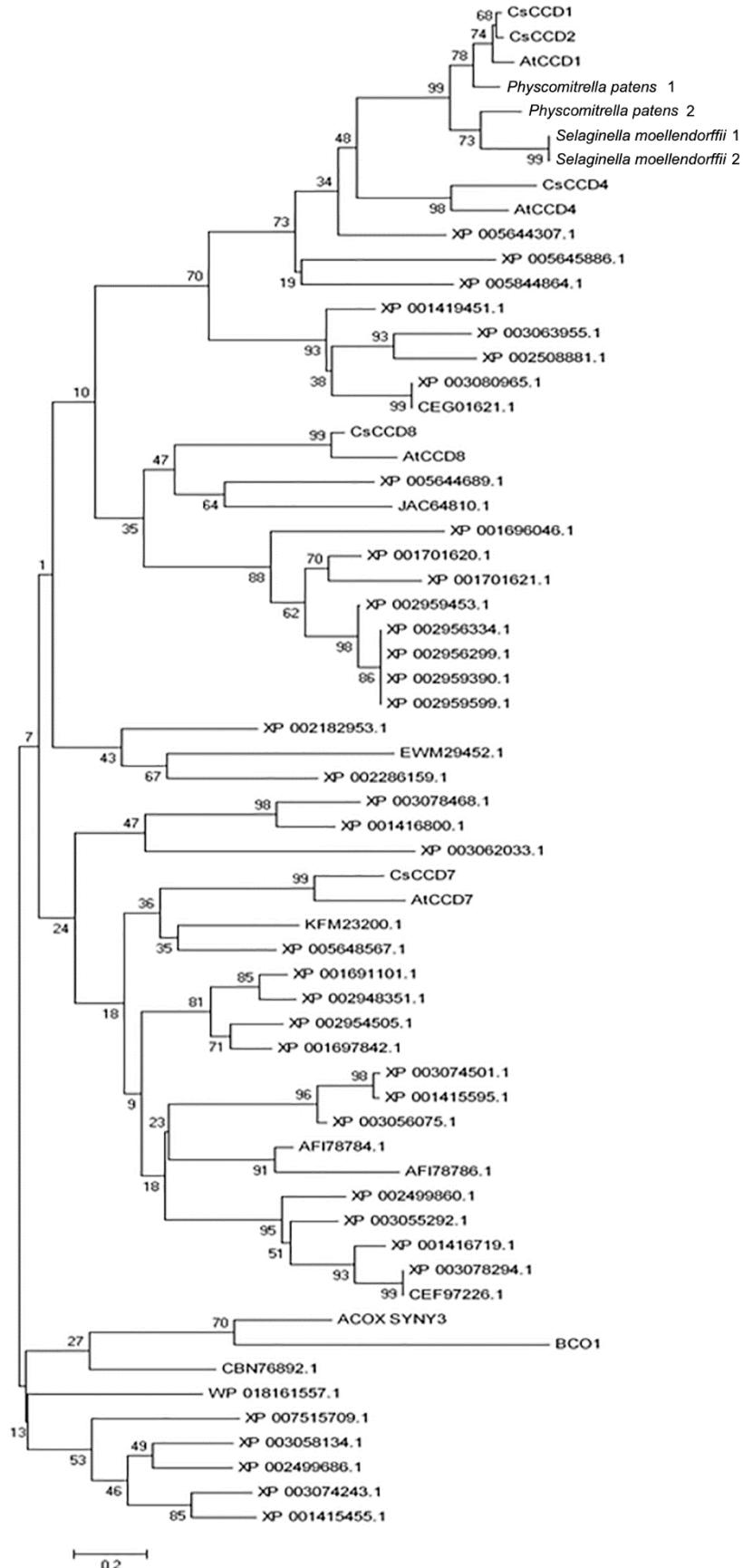


Figure S4. Phylogenetic tree of algal CCD enzymes. Methods for proteins alignment and tree generation are described in the legend of Figure S1.

Table S1. CCD1 genes and proteins present in different plant species.

Species	Protein Code	Intron Number	Gene Bank	Chromosome Number
<i>Acanthus ebracteatus</i>	AFC88468.			
<i>Arabidopsis lyrata</i>	XP_002876727.1	14	NW_003302551.1	3
<i>Arabidopsis thaliana</i>	NP_191911.1	13	AT3G63520.1	3
<i>Arabis alpina</i>	KFK35504.1	13	CM002873.1	
<i>Beta vulgaris</i>	XP_010695196.1	13	NC_025813.1	2
<i>Brachypodium distachyon</i>	XP_003577313.1	13	NC_016134.1	4 tandem
	XP_003577349.1	13	NC_016134.1	4
<i>Brachypodium sylvaticum</i>	ACO87672.1	13	FJ234838.1	
	ACO87673.1	13	FJ234838.1	tandem
<i>Brassica rapa</i>	XP_009117091	13	NC_024803	A9
	XP_010512783.1	13	NC_026891	5
<i>Camellia sativa</i>	XP_010413131.1	19	NC_025691.1	7
	XP_010413132.1			
<i>Capsicum annuum</i>	CAC79644.1			
<i>Castanea mollissima</i>	ABO69703.1			
<i>Cicer arietinum</i>	XP_004510991.1			
	XP_004510990.1	13	NC_021166.1	Ca7
<i>Citrus clementina</i>	XP_006439107.1			
	XP_006439108.1	14	NW_006262274.1	
<i>Citrus sinensis</i>	XP_006482868.1			
	XP_006482869.1	13	NC_023052.1	7
	XP_006482870.1			
<i>Coffea arabica</i>	ABA43904.1			
<i>Coffea canephora</i>	ABA43900.1	13	GSCOC_T00025171001	2
<i>Crocus sativus</i>	Q84KG5.1			
<i>Cryptomeria japonica</i>	BAF31905			
<i>Cucumis melo</i>	XP_008461300.1	13	NW_007546327.1	
<i>Cucumis sativus</i>				
<i>Daucus carota</i>	ABB52081.1			
<i>Elaeis guineensis</i>	XP_010931015.1	13	NC_026001	9
<i>Eriobotrya japonica</i>	AFG30025.1			
<i>Eualiptus grandis</i>	XP_010067867.1	13	GO3374	
	KCW66101.1	13	GO3362	tandem
<i>Eutrema salsugineum</i>	XP_006402261.1			
	XP_006402262.1			
	XP_006402263.1	13	NW_006256838.1	
	XP_006402264.1			
<i>Fragaria vesca</i>	XP_004306539.1	13	NC_020497.1	LG7
<i>Glycine max</i>	XP_003542847.1	13	NC_0161001	13
<i>Glycine soja</i>	KHN19457	13	KN659065.1	
<i>Jatropha curcas</i>	KDP43480.1	13	KK914267.1	
<i>Lactuca sativa</i>	BAE72095.1			
<i>Malus domestica</i>	XP_008375177.1	13	NC_024245.1	7
	XP_008363317.1	13	NW_007545925.1	2
	XP_008349836.1	9	NC_024240	2
<i>Malus hupehensis</i>	ACF75911.1			
<i>Manihot sculenta</i>	ADN65332.1			
<i>Medicago truncatula</i>	CAR57918.1			
<i>Momordica charantia</i>	AFU91489.1			
<i>Morus notabilis</i>	XP_010088226.1	13	NW_010357396	
<i>Musa acuminata</i>	XP_009394395.1	14	NC_025204.1	3
<i>Nelumbo nucifera</i>	XP_010241260.1	14	NW_010729221.1	
	XP_010241261.1			
<i>Nicotiana sylvestris</i>	XP_009760607.1	13	NW_009359434.1	
	XP_009784260	13	NW_009344641.1	
	XP_009784266			
<i>Nicotiana tabacum</i>	AIL30506.1			

Table S1. Cont.

Species	Protein Code	Intron Number	Gene Bank	Chromosome Number
<i>Nicotiana tomentosiformis</i>	XP_009589876.1			
	XP_009589877.1			
	XP_009589878.1	13	NW_008927330.1	
	XP_009589879.1			
	XP_009589880.1			
	XP_009589881.1			
<i>Oryza sativa</i>	ABA99624	13		
	ABA99623	13		
<i>Oryza brachiiata</i>	XP_006661114.1	12	NC_023171.1	9
<i>Osmanthus fragans</i>	BAJ05401.1			
<i>Panicum virgatum</i>	Pavirv00030688m	12	Pavirv00030688m	
<i>Petunia x hybrida</i>	AAT68189.1			
<i>Phaseolus vulgaris</i>	Q94IR2			
<i>Phoenix dactylifera</i>	XP_008801952.1	13	NW_008246719.1	
<i>Physcomitrella patens</i>	XP_001771028.1	12	Pp1s132_26V6	
	XP_001754114.1	14	Pp1s12_317V6	
<i>Picea glauca</i>	BT115844.1			
<i>Picea sitchensis</i>	ABK24523.1			
<i>Pirux x bretschneideri</i>	XP_009365069.1	12	NW_008988158.1	
	XP_009367500.1	12	NW_008988186.1	
<i>Pisum sativum</i>	Q8LP17			
<i>Populus trichocarpa</i>	XP_006379094.1	13	NC_008475.2	LGIX
	XP_002298429.2	13	NC_008467.2	LGI tandem
	XP_002298430.2			
<i>Populus euphratica</i>	XP_011019809	12	NW_011499853.1	
	XP_011012510.1	12	NW_011500894.1	
	XP_011005228.1	12	NW_011500182.1	
	XP_011019620.1	12	NW_011499853.1	
<i>Prunus mume</i>	XP_008231253.1	13	NC_024130.1	LG5
<i>Prunus persica</i>	XP_007220659.1	13	NW_006760324.1	
<i>Ricinus communis</i>	XP_002517824.1	11	NW_002994325.1	
<i>Rosa damascena</i>	ABY47994.1			
<i>Saccharum hybrid cultivar R570</i>	AGT17077.1	12	SHCRBa_185_C01_R_60	
<i>Scutellaria baicalensis</i>	AGN03859.1			
<i>Selaginella moellendorffii</i>	XP_002962877.1	13	165469	
	XP_002989454.1			
<i>Setaria italica</i>	XP_004963348.1	13	NW_004675963.1	
<i>Solanum lycopersicum</i>	XP_010324092	13	NC_015438.2	1
	NP_001234542			
<i>Solanum nigrum</i>	AHA43413.1			
<i>Solanum tuberosum</i>	XP_006345441.1	14	NW_006238994.1	
	XP_006345442.1			
<i>Sorghum bicolor</i>	XP_002465882.1	12	Sb01g047540.1	1
<i>Suaeda salsa</i>	AAY21819.1			
<i>Tanacetum cinerariifolium</i>	AFV15390.1			
<i>Tarenaya hassleriana</i>	XP_010552739.1			
	XP_010552740.1	15	NW_010966822.1	
	XP_010552741.1			
<i>Theobroma cacao</i>	XP_007052518.1	13	NC_023625.1	1
	XP_007052519.1			
<i>Vitis vinifera</i>	AFJ94680	12	JQ712833	13
<i>Zea mays</i>		13	GRMZM2G057243_T01	9

In bold, presence of spliced variants.

Table S2. CCD7 genes and proteins present in different plant species.

Species	Protein Code	Intron Number	Gene Bank	Chromosome Number
<i>Actinidia chinensis</i>	ADP37985.1		ADP37985.1	
<i>Amborella trichopoda</i>	XP_011621212.1		XM_011622910.1	
<i>Aquilegia coerulea</i>	Aqua_037_00073.1 (Phytozome code)	6		
<i>Arabidopsis lyrata</i>	XP_002880149.1	5	XM_002880103.1	
<i>Arabidopsis thaliana</i>	NP_182026.4	6	NC_003071.7	2
<i>Artemisia annua</i>	ADB64459.1		GQ996728.1	
<i>Beta vulgaris</i>	XP_010692115.1		XM_010693813.1	
<i>Boechera stricta</i>	Bostr.25993s0273.1 (Phytozome code)	5		
<i>Brachypodium distachyon</i>	XP_003581501.1	7	XM_003581453.2	5
<i>Brassica napus</i>	CDY37022.1		LK032390.1	
<i>Brassica rapa</i>	XP_009142313.1	6	XM_009144065.1	
<i>Camelina sativa</i>	XP_010518109.1		XM_010519807.1	
<i>Capsella grandiflora</i>	Cagra.0239s0052.1 (Phytozome code)	5		
<i>Capsella rubella</i>	XP_006295645.1	5	XM_006295583.1	
<i>Carica papaya</i>	evm.model.supercontig_22.7 (Phytozome code)	6		
<i>Cicer arietinum</i>	XP_004513935.1		XM_004513878.1	
<i>Citrus clementina</i>	XP_006425527.1	6	XM_006425464.1	
<i>Citrus sinensis</i>	XP_006467157.1	6	XM_006467094.1	
<i>Coffea canephora</i>	CDP09468.1		HG739122.1	
<i>Crocus sativus</i>	AIF27228.1		KJ361477.1	
<i>Cucumis melo</i>	NP_001284427.1		NM_001297498.1	
<i>Cucumis sativus</i>	XP_004140624.1		XM_004140576.2	6
<i>Elaeis guineensis</i>	XP_010934132.1		XM_010935830.1	
<i>Eucalyptus grandis</i>	KCW79101.1	6	KK198755.1	
<i>Eucalyptus grandis</i>	XP_010056416.1		XM_010058114.1	
<i>Eutrema salsugineum</i>	XP_006397680.1	6	XM_006397617.1	
<i>Fragaria vesca</i>	XP_004306976.2	4	XM_004306928.2	
<i>Glycine max</i>	ADK26570.1	6	HM366150.1	
<i>Glycine soja</i>	KHN22759.1		KN656920.1	
<i>Gossypium raimondii</i>	KJB49230.1	5	CM001747.1	8
<i>Jatropha curcas</i>	XP_012083689.1		XM_012228299.1	
<i>Linum usitatissimum</i>	Lus10021241 (Phytozome code)	5		
<i>Lotus japonicus</i>	ADM88552.1		GU441766.1	
<i>Malus baccata</i>	AHJ78579.1		KF887978.1	
<i>Malus domestica</i>	XP_008363204.1	7	XM_008364982.1	2
<i>Malus hupehensis</i>	AHJ78580.1		KF887979.1	
<i>Malus micromalus</i>	AHJ78576.1		KF887975.1	
<i>Malus prunifolia</i>	AHJ78582.1		KF887981.1	
<i>Malus sieversii</i>	AHJ78581.1		KF887980.1	
<i>Malus toringoides</i>	AHJ78577.1		KF887976.1	
<i>Manihot esculenta</i>	cassava4.1_032749m (Phytozome code)	6		
<i>Medicago truncatula</i>	XP_003622555.1	6	XM_003622507.1	
<i>Mimulus guttatus</i>	EYU20441.1	6	KI632284.1	
<i>Morus notabilis</i>	XP_010089272.1		XM_010090970.1	
<i>Musa acuminata</i>	XP_009384463.1		XM_009386188.1	
<i>Nelumbo nucifera</i>	XP_010264426.1		XM_010266124.1	
<i>Nicotiana tabacum</i>	AFU10970.1		JQ034523.1	
<i>Nicotiana tomentosiformis</i>	XP_009592649.1		XM_009594354.1	
<i>Orobanche ramosa</i>	AEQ30075.1		JN412814.1	
<i>Oryza brachyantha</i>	XP_006652609.1		XM_006652546.1	

Table S2. Cont.

Species	Protein Code	Intron Number	Gene Bank	Chromosome Number
<i>Oryza sativa</i>	Q7XU29.2	6		4
<i>Panicum virgatum</i>	Pavir.Gb00928.1 (Phytozome code)	5		
<i>Petunia x hybrida</i>	ACY01408.1		FJ790878.1	
<i>Phoenix dactylifera</i>	XP_008796153.1		XM_008797931.1	
<i>Physcomitrella patens</i>	ADK36680.1		HM007802.1	
<i>Pisum sativum</i>	ABD67496.2		DQ403160.1	
<i>Populus trichocarpa</i>	XP_006375244.1	6	XM_006375182.1	14
<i>Prunus mume</i>	XP_008232618.1		XM_008234396.1	
<i>Prunus persica</i>	XP_007221108.1	5	XM_007221046.1	
<i>Pyrus x bretschneideri</i>	XP_009373202.1		XM_009374927.1	
<i>Ricinus communis</i>	XP_002511629.1	6	XM_002511583.1	
<i>Salix purpurea</i>	SapurV1A.0382s0050.1 (Phytozome code)	6		
<i>Sesamum indicum</i>	XP_011094476.1		XM_011096174.1	
<i>Setaria italica</i>	XP_004976437.1	5	XM_004976380.1	7
<i>Solanum lycopersicum</i>	ACY39883.1	6	GQ468556.1	1
<i>Solanum tuberosum</i>	XP_006359777.1	6	XM_006359715.1	1
<i>Sorghum bicolor</i>	XP_002446902.1	6	XM_002446857.1	6
<i>Tarenaya hassleriana</i>	XP_010544591.1		XM_010546289.1	
<i>Theobroma cacao</i>	EOX96379.1	6	CM001879.1	
<i>Trifolium repens</i>	AHN65153.1		KJ127512.1	
<i>Vitis vinifera</i>	XP_010648499.1	5	XM_010650197.1	15
<i>Zea mays</i>	NP_001183928.1	6	NM_001196999.1	2

In bold, presence of spliced variants.

Table S3. CCD8 genes and proteins present in different plant species.

Species	Protein Code	Intron Number	Gene Bank	Chromosome Number
<i>Actinidia chinensis</i>	ADP37984.1		GU206812.1	
<i>Aquilegia coerulea</i>	Aquca_001_00324.1 (Phytozome code)	5		
<i>Arabidopsis lyrata</i>	XP_002869241.1	5	XM_002869195.1	
<i>Arabidopsis thaliana</i>	NP_195007.2	5	NM_119434.3	4
<i>Arabidopsis thaliana</i>	Q8VY26.1		NM_119434.3	
<i>Arabis alpina</i>	KFK29890.1		CM002875.1	7
<i>Beta vulgaris</i>	XP_010668385.1		XM_010670083.1	2
<i>Boechera stricta</i>	Bostr.7867s1131.1 (Phytozome code)	5		
<i>Brachypodium distachyon</i>	XP_003569798.1	3	XM_003569750.2	
<i>Brassica rapa</i>	XP_009125426.1	5	XM_009127178.1	1
<i>Camelina sativa</i>	XP_010432658.1		XM_010434356.1	10
<i>Camelina sativa</i>	XP_010437851.1		XM_010439549.1	11
<i>Camelina sativa</i>	XP_010447340.1		XM_010449038.1	12
<i>Camelina sativa</i>	XP_010447342.1		XM_010449040.1	12
<i>Capsella grandiflora</i>	Cagra.4093s0010.1	5		
<i>Capsella rubella</i>	XP_006285526.1	5	XM_006285464.1	
<i>Carica papaya</i>	evm.TU.contig_24654.4	5		
<i>Cicer arietinum</i>	XP_004501157.1		XM_004501100.1	5
<i>Citrus clementina</i>	XP_006450690.1	5	XM_006450627.1	
<i>Citrus sinensis</i>	KDO79823.1	5	KK784877.1	
<i>Citrus sinensis</i>	XP_006476130.1		XM_006476067.1	4
<i>Crocus sativus</i>	AIF27229.1	5	KJ361478.1	
<i>Crocus sativus</i>	AIF27230.1	5	KJ361479.1	
<i>Cucumis melo</i>	NP_001284455.1		NM_001297526.1	
<i>Cucumis melo</i>	AHY18725.1		KJ473491.1	
<i>Cucumis melo</i>	XP_008445014.1		XM_008446792.1	
<i>Cucumis sativus</i>	XP_004148387.1	5	XM_004148339.2	2
<i>Elaeis guineensis</i>	XP_010917340.1		XM_010919038.1	1
<i>Elaeis guineensis</i>	XP_010924674.1		XM_010926372.1	6
<i>Eucaliptus grandis</i>	KCW81574.1		KK198755.1	
<i>Eucalyptus grandis</i>	XP_010049109.1	5	XM_010050807.1	
<i>Eucalyptus grandis</i>	XP_010051229.1		XM_010052927.1	
<i>Eutrema salsugineum</i>	XP_006412429.1	5	XM_006412366.1	
<i>Fragaria vesca</i>	XP_011458989.1	4	XM_011460687.1	
<i>Glycine max</i>	XP_003522713.2	5	XM_003522665.2	4
<i>Glycine max</i>	NP_001242715.1	5	NM_001255786.1	6
<i>Glycine soja</i>	KHN10194.1		KN664752.1	
<i>Glycine soja</i>	KHN18168.1		KN659929.1	
<i>Gossypium raimondii</i>	KJB07492.1	5	CM001740.1	1
<i>Gossypium raimondii</i>	KJB65669.1	5	CM001749.1	10
<i>Jatropha curcas</i>	XP_012077359.1		XM_012221969.1	
<i>Linum usitatissimum</i>	Lus10001599	5		
<i>Malus baccata</i>	AIN41153.1		KF887984.1	
<i>Malus baccata</i>	AIN41152.1		KF887983.1	
<i>Malus domestica</i>	XP_008352014.1	6	XM_008353792.1	15
<i>Malus domestica</i>	XP_008361152.1	4	XM_008362930.1	15
<i>Malus domestica</i>	XP_008378214.1	5	XM_008379992.1	8
<i>Malus hupehensis</i>	AIN41154.1		KF887985.1	
<i>Malus hupehensis</i>	AIN41155.1		KF927168.1	
<i>Manihot esculenta</i>	cassava4.1_005134m	5		
<i>Medicago truncatula</i>	XP_003603610.1	5	XM_003603562.1	3
<i>Medicago truncatula</i>	KEH23011.1	5	CM001223.2	7
<i>Mimulus guttatus</i>	EYU20435.1	5	KI632284.1	
<i>Musa acuminata</i>	XP_009389368.1		XM_009391093.1	2
<i>Musa acuminata</i>	XP_009407022.1		XM_009408747.1	6
<i>Nelumbo nucifera</i>	XP_010249576.1		XM_010251274.1	
<i>Nicotiana sylvestris</i>	XP_009779163.1		XM_009780861.1	

Table S3. *Cont.*

Species	Protein Code	Intron Number	Gene Bank	Chromosome Number
<i>Nicotiana tabacum</i>	AGO64766.1		KC795555.1	
<i>Nicotiana tomentosiformis</i>	XP_009626361.1		XM_009628066.1	
<i>Oryza brachyantha</i>	XP_006446317.1		XM_006446254.1	1
<i>Oryza sativa</i>	NP_001044229.2		NM_001050764.2	
<i>Oryza sativa</i>	Q8LIY8.1		OSJNBa0014K08.38	
<i>Panicum virgatum</i>	Pavir.Ea02892.1 (Phytozome code)	3		
<i>Panicum virgatum</i>	Pavir.Eb03110.1 (Phytozome code)	3		5
<i>Petunia hybrida</i>	AAW33596.1		AY746977.1	
<i>Phaseolus vulgaris</i>	XP_007137221.1	5	XM_007137159.1	9
<i>Phoenix dactylifera</i>	XP_008810664.1		XM_008812442.1	
<i>Phoenix dactylifera</i>	XP_008804630.1		XM_008806408.1	
<i>Physcomitrella patens</i>	ADK36681.1		HM007803.1	
<i>Pisum sativum</i>	AAS66907.1		AY557342.1	3
<i>Populus trichocarpa</i>	XP_002309543.1	5	XM_002309507.1	
<i>Populus trichocarpa</i>	XP_002324797.1	5	XM_002324761.1	
<i>Prunus mume</i>	XP_008220105.1		XM_008221883.1	
<i>Prunus persica</i>	XP_007222386.1	5	XM_007222324.1	
<i>Ricinus communis</i>	XP_002516503.1	4	XM_002516457.1	
<i>Salix purpurea</i>	SapurV1A.0272s0100.1	5		
<i>Sesamum indicum</i>	XP_011070469.1		XM_011072167.1	
<i>Sesamum indicum</i>	XP_011082234.1		XM_011083932.1	
<i>Setaria italica</i>	XP_004972173.1		XM_004972116.1	
<i>Solanum lycopersicum</i>	NP_001266276.1	5	NM_001279347.1	
<i>Solanum tuberosum</i>	XP_006359761.1		XM_006359699.1	
<i>Sorghum bicolor</i>	XP_002458477.1		XM_002458432.1	
<i>Tarenaya hassleriana</i>	XP_010548752.1		XM_010550450.1	
<i>Theobroma cacao</i>	XP_007012130.1	5	XM_007012068.1	
<i>Theobroma cacao</i>	EOY29749.1		CM001887.1	
<i>Triticum urartu</i>	EMS51709.1		KD216346.1	
<i>Vitis vinifera</i>	XP_002281239.1	4	XM_002281203.2	4
<i>Zea mays</i>	NP_001183929.1	2	NM_001197000.1	3

In bold, presence of spliced variants.