

Supplementary information for Scientific Reports

Characterization of the polyphenol oxidase gene family reveals a novel microRNA involved in posttranscriptional regulation of *PPOs* in *Salvia miltiorrhiza*

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Supplementary Table S1 Primers used for 5'-RACE of *SmPPOs*.

Gene name	Primer sequence (5'-3')
<i>SmPPO13</i>	Nesting: TGGATTCCGGGTCATGCCGAGATCC
	Nested: GTCAGACATCGGTGGGCAGCAGTTCATG
<i>SmPPO14</i>	Nesting: AGATCCGTGGGAGCAGGCGGGAGAT
	Nested: TTGCATGAAACTGCGCGGGTCATCG
<i>SmPPO15</i>	Nesting: CGTACGTATTCCGGTGTTCAGCCTGTGCG
	Nested: TGCGATTGAGTGGGTTGCGGTTGCAG
<i>SmPPO16</i>	Nesting: CCTGTTTCGTAGCGGCCGGTACAGTAAGC
	Nested: AGCCGCCCTCCGACAATCCTCCCTT
<i>SmPPO17</i>	Nesting: TCTTCTACGCACATCACGCCAACG
	Nested: TCTTTCCTTACCCAGGGCCTCAATC
<i>SmPPO18</i>	Nesting: CCCGACGACTTCAACAACGCCTCC
	Nested: GAGGATTGTTCGGAGGGCGGCTGTCT
<i>SmPPO19</i>	Nesting: ATCTCTAGACGTCAACTGCTGCCCCG
	Nested: CGCTTCCGATTTTGCCTCCTGACCT

Supplementary Table S2 Primers used for 3'-RACE of *SmPPOs*.

Gene name	Primer sequence (5'-3')
<i>SmPPO17</i>	Nesting: CACGGAGTTTCGCTTTCTTCATAC
	Nested: ACTTCGTCGGCTTTGCCTTTGGTTG
<i>SmPPO18</i>	Nesting: ACTTCGTCGGCTTTGCCTTTGGT
	Nested: CGCTTCCGCAGTCCTAACCTTAT
<i>SmPPO19</i>	Nesting: TGGCGAGAACTTTGGCATTGACC
	Nested: CGTCGGCTTTGCCTTTGGTTGGT

Supplementary Table S3 Primers used for the PCR-amplification of coding sequences of *SmPPOs*

Gene name	Primer sequence (5'-3')
<i>SmPPO1</i>	Forward: ATGGCTTCGTCGATTCTCTCACAT Reverse: GTCATCAAGCTCGATCTTGATATCATG
<i>SmPPO2</i>	Forward: ATGTCATCTGTTCTCTTTCCACCG Reverse: CCACCAATGGTAATATCTTCTCCTCCA
<i>SmPPO3</i>	Forward: ATGGCATTTCCTCCACTTTCCTC Reverse: TGACACCACCAATGGTAATGTCTTC
<i>SmPPO4</i>	Forward: ATGGCCTCTCTTCAATCTTCATGC Reverse: CGCGACTTAACAATTTTGATACCA
<i>SmPPO5</i>	Forward: ATGGCTTCCCTTCAATCTTCTTTCTC Reverse: GCGGGATGATCTTAATACCACCAAT
<i>SmPPO6</i>	Forward: ATGGCTTCCCTCTACCTTCATTGCA Reverse: AGCTTTAGGTGGATTCTCAATGATCTTGA
<i>SmPPO7</i>	Forward: ATGGCTTCTCTTCCATCTTCACTCA Reverse: AGCAGCAACACGCGGGATAATCT
<i>SmPPO8</i>	Forward: ATGGCTTCCCTTCAATCTTCATTC Reverse: ACACGCGGGATGATCTTAATACC
<i>SmPPO9</i>	Forward: ATGGCTTCTTCCCTTCAGTCCTTAAG Reverse: TTTACGCTGCGCTGGGATGA
<i>SmPPO10</i>	Forward: ATGGCTTCCCTTCAATCTTCATTCAC Reverse: TCAAGCGGTAACACGCGGGAT
<i>SmPPO11</i>	Forward: ATGGCTTCTTCCCTTATTGCTTCATG Reverse: ACGCTGCGCTGGGACGATCTT
<i>SmPPO12</i>	Forward: ATGGCTTCCCTTCAATCTTCATTCA Reverse: AGCGGTAACACGCGGGATGATC
<i>SmPPO13</i>	Forward: ATGGCTTCCCTCTACCTTCATTGC Reverse: TCAAGATCTTCGGGGTGGAGAAG
<i>SmPPO14</i>	Forward: ATGGCATCACTTCCGTTTTCCAC Reverse: TTAAACGTAGATGATCTTGACCCAC
<i>SmPPO15</i>	Forward: ATGGCTTCCCTTCAATCTTCATTC Reverse: CACGCGGGATGATCTTAATACCA
<i>SmPPO16</i>	Forward: ATGGGGCACCAAACCATGGC Reverse: TCAAGCGGTAACACGAGGAATGATC
<i>SmPPO17</i>	Forward: ATGCTCCTCGGTCTTGGAGGGC Reverse: TCAAACGGGAACACGCGGGATG
<i>SmPPO18</i>	Forward: ATGGCGAAGATGAGGGAGCTG Reverse: TCAAGCGGTAACACGCGGGATG
<i>SmPPO19</i>	Forward: ATGGCTTCCCTTCAATCTTC Reverse: TCAAGCAGTAACACGCGGGATG

Supplementary Table S4 Gene names of *PPOs*.

Gene name	Accession no.
<i>Populus trichocarpa</i>	
<i>PtPPO1</i>	Potri.011G108300
<i>PtPPO2</i>	Potri.001G387900
<i>PtPPO3</i>	Potri.011G047300
<i>PtPPO4</i>	Potri.T062100
<i>PtPPO5</i>	Potri.001G388600
<i>PtPPO6</i>	Potri.001G388400
<i>PtPPO7</i>	Potri.001G388300
<i>PtPPO8</i>	Potri.T061900
<i>PtPPO9</i>	Potri.001G388100
<i>PtPPO10</i>	Potri.011G108200
<i>PtPPO11</i>	Potri.001G388900
<i>PtPPO12</i>	Potri.001G388800
<i>PtPPO13</i>	Potri.004G156500
<i>PtPPO14</i>	Potri.001G388200
<i>PtPPO15</i>	Potri.T062200
<i>PtPPO16</i>	Potri.001G388000
<i>Glycine max</i>	
GmPPO1	Glyma06g42170.1
GmPPO2	Glyma07g31270.1
GmPPO3	Glyma07g31280.1
GmPPO4	Glyma07g31300.1
GmPPO5	Glyma07g31310.1
GmPPO6	Glyma13g25150.1
GmPPO7	Glyma13g25180.1
GmPPO8	Glyma13g25260.1
GmPPO9	Glyma13g31590.1
GmPPO10	Glyma15g07710.1
GmPPO11	Glyma18g45900.1
<i>Zea mays</i>	
ZmPPO1	GRMZM2G108103_T01
ZmPPO2	GRMZM2G121605_T01
ZmPPO3	AC233851.1_FGT017
ZmPPO4	GRMZM2G319062_T01
ZmPPO5	GRMZM2G137909_T01
ZmPPO6	AC209206.3_FGT014
<i>Oryza sativa</i>	
OsaPPO1	LOC_Os01g58100.1
OsaPPO2	LOC_Os04g53300.1
<i>Phycomitrella patens</i>	
PpPPO1	AAX69084.1
PpPPO2	XP_001770109.1

PpPPO3	XP_001785624.1
PpPPO4	XP_001752158.1
PpPPO5	XP_001773770.1
PpPPO6	XP_001785949.1
PpPPO7	XP_001767174.1
PpPPO8	XP_001766337.1
PpPPO9	XP_001766259.1
PpPPO10	XP_001766553.1
PpPPO11	XP_001755077.1
PpPPO12	XP_001785245.1

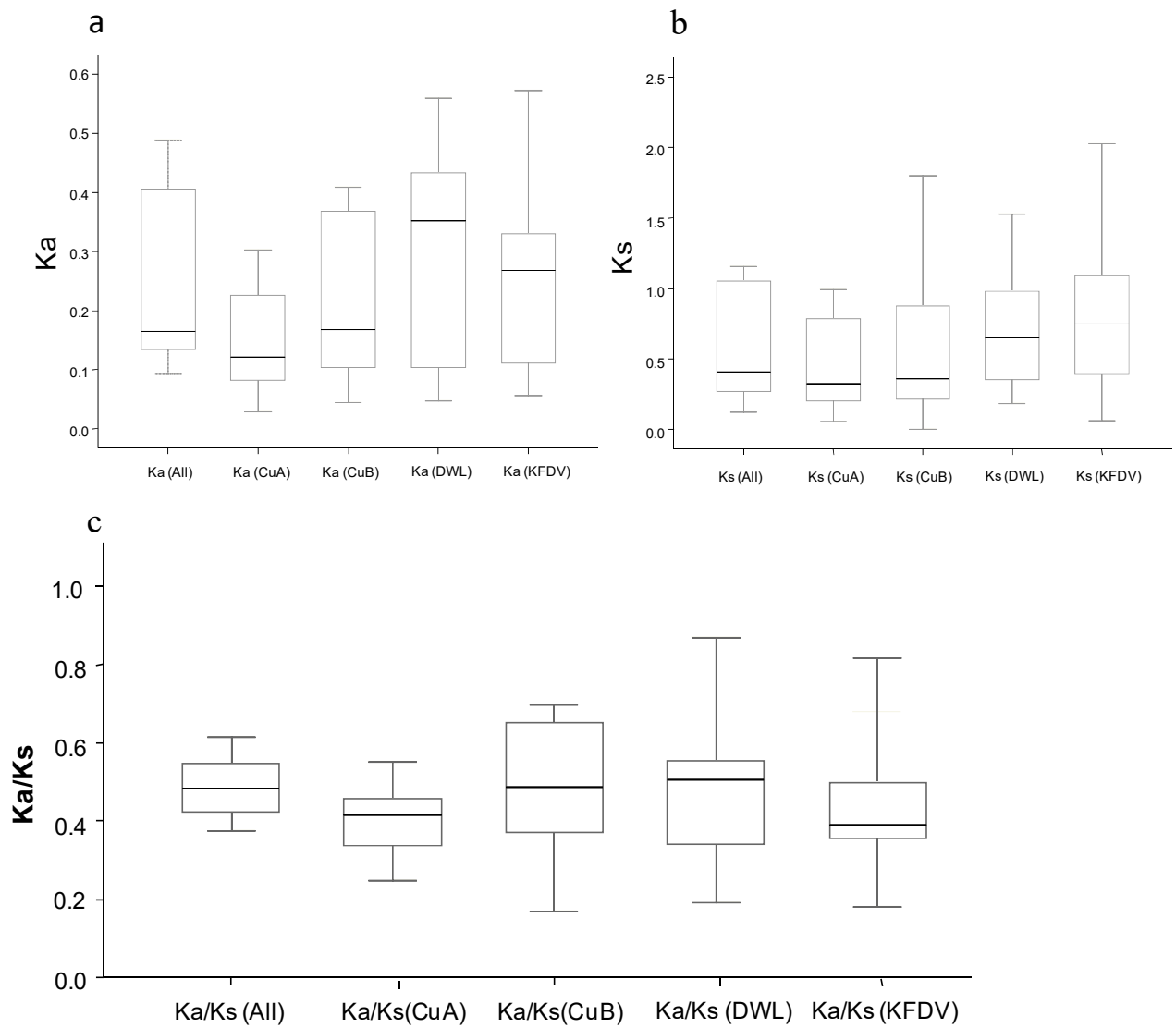
Supplementary Table S5 Primers used for quantitative real-time RT-PCR of *SmPPOs*

Gene name	Primer sequence (5'-3')
<i>SmPPO1</i>	Forward: GAAGAGGAGCAAGAAGGAGAAG
	Reverse: TCGTCCTCGTCGTTGATGTA
<i>SmPPO2</i>	Forward: TCGTGAGGTTTGAGGTGTTC
	Reverse: AGCCTGATCTTGCTCTTCAC
<i>SmPPO3</i>	Forward: GGTAAGGGTGTGGTGGATAG
	Reverse: TATTTGGCGGTGTTCGATCTC
<i>SmPPO4</i>	Forward: GCCTTCTACTCCCATCACTCTA
	Reverse: CGTTGAGGTAGTCAGGTTTGTT
<i>SmPPO5</i>	Forward: CGGTCAATCTCAAGGACATCT
	Reverse: CCTTTGGTTGGTTTGGACAC
<i>SmPPO6</i>	Forward: GCATGTGGACTCTATGGAAGG
	Reverse: GAAAGAGGCGTTGAGGTAGTC
<i>SmPPO7</i>	Forward: CACAGTGTTCCCTCTCAAAC
	Reverse: CCTCACGAGCTTGCTGTTAT
<i>SmPPO8</i>	Forward: CAAAGTGTTCCCTCTCACTCTC
	Reverse: AACTCACCAGCTTGGAACATC
<i>SmPPO9</i>	Forward: CTGAAGAAGGCGGTTAGAGTTT
	Reverse: TTGGTGCAATCGGTCTCTATG
<i>SmPPO10</i>	Forward: GGTGCATATCGAAGAGTGGAT
	Reverse: GAGAGTGAGAGGGAACAGATTG
<i>SmPPO11</i>	Forward: GGAGCTCTACGAGAACATTGAC
	Reverse: GACGATCTTGACACCACCAA
<i>SmPPO12</i>	Forward: CAATCTGTTCCCTCTCACTCTC
	Reverse: ACCTCACGACCTTCAAACATC
<i>SmPPO13</i>	Forward: TGTGGACTCTGTGGCAATC
	Reverse: GCGTCCTCGTCGTAGAATAAG
<i>SmPPO14</i>	Forward: CGACTACGGCTAAGGCAATTA
	Reverse: CTCCTTCTCCTTCTTGCTTCTC

<i>SmPPO15</i>	Forward: AACTCCATGGATGGCTTACAG Reverse: ACAGTGTTGGTGAGAGTGATG
<i>SmPPO16</i>	Forward: CGTGTATGAACACAGCGAAAC Reverse: GAGAGTGAGAGGGAACAGATTG
<i>SmPPO17</i>	Forward: CAATGTGTTCCCTCTCACTCTC Reverse: AACTCACCAGCTTGGAACATATC
<i>SmPPO18</i>	Forward: TCGTGAGGTTTCGATGTGTTC Reverse: CTTTCCGCAGTCCTAACCTTAT
<i>SmPPO19</i>	Forward: CCTCTCACTCTCACCAAGACTA Reverse: CCTCACGATCTTGGACCTTTC

Supplementary Table S6 Primers used for analysis of Smi-miR12112-directed cleavage of *SmPPOs*

Gene name	Primer sequence (5'-3')
<i>SmPPO3</i>	Nesting: GCAGCAAGCAGTACAAGTAGTG Nested: GTGTAGACGATGATTCACAGAAG
<i>SmPPO5</i>	Nesting: GTCGACAAATCTCAAGCAACATGGAG Nested: GAAACATTGGCGATGGATGCAG
<i>SmPPO9</i>	Nesting: CGGTTAACCGAACCATAAACCGG Nested: GATATGTAGATTTTATTGTGGAGC
<i>SmPPO11</i>	Nesting: GCTTCCCCTCTCTCACACACACAC Nested: CATTGGAGGAATTGGAACATGCG
<i>SmPPO13</i>	Nesting: GGTCTGCCTCCATGGAACGATTGTC Nested: GGAGAAATGCTAGCGTAGGGC



Supplementary Figure S1 Divergence of different classes in SmPPOs. (a) the Ka values of different classes in SmPPOs; (b) the Ks values of different classes in SmPPOs; (c) the Ka/Ks ratios of different classes in SmPPOs.

