

Supplementary Materials: Morphological, Genome and Gene Expression Changes in Newly Induced Autopolyploid *Chrysanthemum lavandulifolium* (Fisch. ex Trautv.) Makino

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1. Primer Combination of SRAP (Sequence-Related Amplified Polymorphism) (77)

em #4 plus me #4 (abbreviated "E4+M4")

M4+E4, M4+E6, M4+E11, M4+E12, M4+E13, M4+E18, M6+E2, M6+E4, M6+E5, M6+E9, M6+E10, M6+E12, M6+E14, M6+E15, M6+E17.

M7+E2, M7+E3, M7+E5, M7+E12, M7+E15, M7+E18, M8+E2, M8+E3, M8+E4, M8+E9, M8+E10, M8+E11, M8+E12, M8+E14, M8+E18.

M10+E2, M10+E3, M10+E4, M10+E5, M10+E9, M10+E10, M10+E11, M10+E12, M10+E17, M10+E18, M21+E2, M21+E3, M21+E4, M21+E5, M21+E10, M21+E11, M21+E12, M21+E14, M21+E15, M21+E17, M21+E18.

M22+E2, M22+E10, M22+E12, M22+E18, M23+E2, M23+E3, M23+E5, M23+E6, M23+E10, M23+E11, M23+E12, M23+E15, M23+E18, M25+E2, M25+E3, M25+E4, M25+E5, M25+E6, M25+E9, M25+E10, M25+E11, M25+E12, M25+E14, M25+E15, M25+E17, M25+E18.

Table S1. Sequences of primers for SRAP analysis.

Primer	Sequence (5'-3')	Primer	Sequence (5'-3')
em2	GACTGCGTACGAATTTGC	me4	TGAGTCCAAACCGGACC
em3	GACTGCGTACGAATTGAC	me6	TGAGTCCAAACCGGTAG
em4	GACTGCGTACGAATTTGA	me7	TGAGTCCAAACCGGTTG
em5	GACTGCGTACGAATTAAC	me8	TGAGTCCAAACCGGTGT
em6	GACTGCGTACGAATTGCA	me10	TGAGTCCAAACCGGATG
em9	GACTGCGTACGAATTACG	me21	GTACATAGAACCGGAGT
em10	GACTGCGTACGAATTTAG	me22	TACGACGAATCCGGACT
em11	GACTGCGTACGAATTTTCG	me23	CACAGTCATGCCGGAAT
em12	GACTGCGTACGAATTGTC	me25	CAGGACTAAACCGGATA
em14	GACTGCGTACGAATTCAG		
em15	GACTGCGTACGAATTCTG		
em16	GACTGCGTACGAATTCGG		
em17	GACTGCGTACGAATTCCA		
em18	GACTGCGTACGAATTGAT		

"em#" and "me#" indicates name of forward primers and reverse primers.

2. Primer Combination of AFLP (Amplified Fragment Length Polymorphism) (56)

*Eco*RI selective primer #2 plus *Mse*I selective primer #5 (abbreviated "E2+M5")

E2+M5, E2+M6, E2+M3, E2+M9, E2+M7, E2+M2, E7+M4, E7+M5, E7+M6, E7+M3, E7+M7, E7+M8, E4+M9, E4+M6, E4+M5, E4+M4, E4+M2, E4+M7, E4+M8.

E9+M8, E3+M8, E3+M7, E3+M2, E3+M4, E3+M5, E3+M6, E3+M3, E3+M9, E5+M7, E5+M4, E5+M5, E5+M6, E5+M3, E5+M9, E9+M7, E9+M2, E9+M4, E9+M5, E9+M6, E9+M3.

E8+M2, E8+M9, E8+M3, E8+M6, E8+M5, E8+M4, E8+M7, E6+M9, E6+M3, E6+M6, E6+M5, E6+M4, E6+M2, E6+M7, E6+M8, E8+M2.

3. Primer Combination of MSAP (Methylation Sensitive Amplified Polymorphism)

EcoRI selective primer #9 plus *HpaII/MspI* selective primer #2 (abbreviated "E9+M2")

E9+H2, E9+H4, E9+H5, E9+H6, E9+H7, E9+H8, E9+H9

E6+H6, E6+H8, E6+H9, E8+H5, E8+H6, E8+H7, E8+H8, E8+H9,

E8+H2, E8+H3, E8+H4, E6+H2, E6+H3, E6+H4,

E4+H3, E4+H5, E4+H6, E4+H7, E4+H8

Table S2. Sequences of adaptors and primers used for pre-amplification and selective amplification in AFLP and MSAP analysis.

Primer	Sequence (5'-3')
<i>EcoRI</i> adaptor-F	CTCGTAGACTGCGTACC
<i>EcoRI</i> adaptor-R	AATTGGTACGCAGTCTAC
<i>MseI</i> adaptor-F	GACGATGAGTCCTGAG
<i>MseI</i> adaptor-R	TACTCAGGACTCAT
<i>HpaII/MspI</i> adaptor-F	GATCATGAGTCCTGCT
<i>HpaII/MspI</i> adaptor-R	CGAGCAGGACTCATGA
<i>EcoRI</i> pre-selective primer-1	GACTGCGTACCAATTCA
<i>MseI</i> pre-selective primer-1	GATGAGTCCTGAGTAAC
<i>HpaII/MspI</i> pre-selective primer-1	ATCATGAGTCCTGCTCGG
<i>EcoRI</i> selective primer-2	GACTGCGTACCAATTCAAC
<i>EcoRI</i> selective primer-3	GACTGCGTACCAATTCAAG
<i>EcoRI</i> selective primer-4	GACTGCGTACCAATTCACA
<i>EcoRI</i> selective primer-5	GACTGCGTACCAATTCACT
<i>EcoRI</i> selective primer-6	GACTGCGTACCAATTCACC
<i>EcoRI</i> selective primer-7	GACTGCGTACCAATTCACG
<i>EcoRI</i> selective primer-8	GACTGCGTACCAATTCAGC
<i>EcoRI</i> selective primer-9	GACTGCGTACCAATTCAGG
<i>MseI</i> selective primer-2	GATGAGTCCTGAGTAACAA
<i>MseI</i> selective primer-3	GATGAGTCCTGAGTAACAC
<i>MseI</i> selective primer-4	GATGAGTCCTGAGTAACAG
<i>MseI</i> selective primer-5	GATGAGTCCTGAGTAACAT
<i>MseI</i> selective primer-6	GATGAGTCCTGAGTAACATA
<i>MseI</i> selective primer-7	GATGAGTCCTGAGTAACATC
<i>MseI</i> selective primer-8	GATGAGTCCTGAGTAACATG
<i>MseI</i> selective primer-9	GATGAGTCCTGAGTAACATT
<i>HpaII/MspI</i> selective primer-2	ATCATGAGTCCTGCTCGGTAA
<i>HpaII/MspI</i> selective primer-3	ATCATGAGTCCTGCTCGGTCC
<i>HpaII/MspI</i> selective primer-4	ATCATGAGTCCTGCTCGGTTC
<i>HpaII/MspI</i> selective primer-5	ATCATGAGTCCTGCTCGGTAC
<i>HpaII/MspI</i> selective primer-6	ATCATGAGTCCTGCTCGGTGC
<i>HpaII/MspI</i> selective primer-7	ATCATGAGTCCTGCTCGGTAG
<i>HpaII/MspI</i> selective primer-8	ATCATGAGTCCTGCTCGGTTG
<i>HpaII/MspI</i> selective primer-9	ATCATGAGTCCTGCTCGGTCA

4. AFLP Marker

4.1. Protein AE7 (*Daucus carota* subsp. *Sativus*)

GACTGCGTACCAATTCAACAATTCTATTCACCTGAGCTTATTGATCAGCTTGAGATATTTGA
 ACATATTAGAGATATTAAGGATCCTGAGCATCCTTTTCTCTAGAAGATTTGAAAGTTATTAC
 AAAGATGCTATTGAGGTAATGATAAGCGCGTTATGTGCGTGTACTTTTACCCAACAGTTGA
 GCACTCAGTATGGCAACGATTATTGGCTTTGTTTGC GGGTTAACTTATGCGTAGCCTTCCTC

CCGTTTCAAGGTTGACATCAGGGTAGCGCCTGGGACTCATGCAACTGAAGATGCAGTGAAC
AAACAGTTGAACGATAATATGTTACTCAG

4.2. *Myb Domain Protein 73, Putative (Theobroma cacao)*

GACTGCGTACCAATTCACAAATTCACGACCCGCCACCTCGTTGACTTTGTCGCTTCCTGGA
GCGGAGTCGAGTGAGGCTTCCTCGGCGGCTGTCACAGTGCCTGCGCCTCCGCTGCCGTGG
CGGCGCCACCAGCTTTTGTGCGGAGTTTTGGCGGTGATGCAAGAGATGATTAGAATGGA
AGTGAGGAATTATATTAGTGGTGTGGGTTTAGGGTTGGCGGCTGTGGGAAATAGGGAGAG
TCGTGGGTTTAGTTTATTTATTTTTGTTTATGGTAAACTAAATTAATTTGTTGATGGTGTGAT
GATCAACATTTAGTTACTCAGGACTCATC

4.3. *No Apical Meristem (NAM) Protein (Cynara cardunculus var. scolymus)*

GATGAGTCCTGAGTAACTTTTCTCTATTTTCTTTGCTTTGAGGATACACAAATTATCCATTTT
GGGATGAGCATGAAACCAAACACAAACAACAGAAATGACATGGTGTAAACGATGAAAACG
ATACCCTCCGAACCTCTCAACTTCTTCAAGGAATCTCAAGAATAGTATCACAAGTTCATGT
CCTTCATGTGGACATGACATACAACTTCAAGCTAAGGGTTTGTTCATGATTTGCCGGGATT
ACCAGCTGGAGTGAAGTTCGACCCGACTGATCAAGAAATTCTTGAGCATTGAAAGCCAAG
GTTGGATGTGATGCAGAAGAGCTTCATCCCCTTATCGATGAATTCATTCCAACCTATTGAAGG
TGAAACTGGAATATGTTATACTCATCCTGAAAAATTACCTGGAGTTAGCAACGATGGCCAG
ATTCGCCATTTCTTTCACCGTCCATCCAAGGCTTACACCACAGGCACACGAAAGCGAAGAA
AGGTGCACAAATTCGACTGCGTACCAATTCAAC

4.4. *Transcription Factor, Teosint-Like1, Cycloidea and Proliferating Cell Factor(TCP) (Cynara cardunculus var. scolymu)*

GACTGCGTACCAATTCACAAATTCGCCGCATCACTTTCATGGCGCTTTGGCGCTTGCTCCTC
ATCATCATCCTTTTGAAGAAGGGTTTTCTCAAATGATGGGCTTTCATCAGGGCCATGGTCAG
GGCAGGTTTCATGGTCAAGGGCAGCATATTATGACTGCTGACCAGATAGCTGAGGCGATTT
CGAGTGGTGGGGTAATAATGGTGGAAATGGTGTGGGAGTGAGAATTATATGAGGAAAA
GGTATAGGGAGGATTTGTTTAAGGATGAGGGTGGTCAGAACAACGAGGTGGCGGTTGGAT
CACCGTCACCTTCGCCGTCTAACAAGGAGGGGAATACTGGTGCCGGAGCAGGGAATAGTA
TGCTCCGGCACCATGGGGGTATGATGCTTTGTTACTCAGGACTCATC

4.5. *Pol-Polyprotein (Silene Latifolia)*

TGAGTCCAAACCGGTGTTGGTAAAAAAGAGTGACGGAGGATGGCGCATGTGCGTGGATTTT
ACTGACATCAACAAAGCATGCCCTAAGGATTGCTACCCCTACCCGAATTCGTACGCAGTC