Supplementary Materials: Genome-Wide Identification and Characterization of Long
Non-Coding RNAs from Mulberry
(*Morus notabilis*) RNA-*seq* Data

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**Table S1.** Primers used in this study.

|  |  |  |
| --- | --- | --- |
| **NO** | **Name** | **Sequence(5′-3′)** |
| **1** | **MnACTIN3-F** | GCATGAAGATCAAGGTGGTG |
| **2** | **MnACTIN3-R** | CATCTGCTGGAAGGTGCTAA |
| **3** | **XLOC\_000003F** | CCATGTGAGATCATCCACC |
| **4** | **XLOC\_000003R** | ACTTGTGTTTAGTTCTGGGAAT |
| **5** | **XLOC\_022903F** | CTAGAAGTTGAGATTTGCCG |
| **6** | **XLOC\_022903R** | GAGAGAAGGAGGGAAACCA |
| **7** | **XLOC\_019631F** | TGCTGAGACAACATTAGACG |
| **8** | **XLOC\_019631R** | AATAACTCTCTCGCCACC |
| **9** | **XLOC\_003002F** | GAGGAACGGGCTAAGATG |
| **10** | **XLOC\_003002R** | ATTTCACTGATTCTGATTGCG |
| **11** | **XLOC\_005792F** | AGCAACATTGATCTGTTCATAG |
| **12** | **XLOC\_005792R** | CCAAGAACCGACTTCACC |
| **13** | **XLOC\_000207F** | CATCTTCTAATCTCCATGCACT |
| **14** | **XLOC\_000207R** | TTTATCCTTAGGCAGGCTTG |
| **15** | **XLOC\_017186F** | TCACACAAATCCAAGATGGAAC |
| **16** | **XLOC\_017186R** | CTTTGTCTGTCCTCTTAACTGA |
| **17** | **XLOC\_009108F** | TGATCTGCACCAACCATTC |
| **18** | **XLOC\_009108R** | ATAGCTGCAACAAGAGCGA |
| **19** | **XLOC\_022049F** | TATGCACATAAGACTTAGCAGC |
| **20** | **XLOC\_022049R** | TCAAGTTGTAGAAAGCACGG |
| **21** | **XLOC\_003519F** | GAGCTAAGCCAACCTAGTAAC |
| **22** | **XLOC\_003519R** | GAGAATCTTACCTGTCCGTG |
| **23** | **XLOC\_013694F** | GTTAGTCGACGACCCATTC |
| **24** | **XLOC\_013694R** | ATTTAGTTGACGCGGGTT |
| **25** | **XLOC\_009918F** | GCTCACTAGCAAGGGAATGATA |
| **26** | **XLOC\_009918R** | CTCACAAGAAGCCGGTAA |
| **27** | **XLOC\_009911F** | TGAGGAATAACTGCTGGGAAA |
| **28** | **XLOC\_009911R** | GACCCACACGTAGAATAATACT |
| **29** | **XLOC\_003878F** | TCTGTATTCTTGTATGGAGGCA |
| **30** | **XLOC\_003878R** | AAGGGACTTCAACTCAATGAAA |
| **31** | **XLOC\_010969F** | ATTTGTCCAAGAGCAGGCA |
| **32** | **XLOC\_010969R** | TCTCCACTGTTCTCGTCG |
| **33** | **XLOC\_022840F** | TGTGAAGTCAGTCAGCTCC |
| **34** | **XLOC\_022840R** | GACTAAGGTTGTGGGTCTTATT |
| **35** | **XLOC\_000033F** | AGAACCTAAACCTGCTGG |
| **36** | **XLOC\_000033R** | CCTTGTGCTTTGGAACTAC |

**Table S2.** Result of conservation analysis.

|  |  |  |  |  |
| --- | --- | --- | --- | --- |
| **LncRNA ID** | **Mapped P.tri LncRNA NO** | **Mapped P.tri LncRNA List** | **Mapped Tair LncRNA NO** | **Mapped Tair LncRNA List** |
| Mn\_lnc-0001 | 3 | lincRNA1509 lincRNA2088 lincRNA2100 | 1 | NONATHT000414 |
| Mn\_lnc-0002 | 1 | lincRNA1270 | 66 | NONATHT000064,NONATHT000068…… |
| Mn\_lnc-0003 | 1 | lincRNA2062 | 7 | NONATHT000970,NONATHT001685…… |
| Mn\_lnc-0004 | 1 | lincRNA2062 | 7 | NONATHT000970,NONATHT001685…… |
| Mn\_lnc-0005 | 1 | lincRNA3306 | 5 | NONATHT000140,NONATHT003062…… |
| Mn\_lnc-0006 | 1 | lincRNA1209 | 3 | NONATHT000984,NONATHT001209…… |
| Mn\_lnc-0010 | 0 | — | 27 | NONATHT000043,NONATHT000052…… |
| Mn\_lnc-0011 | 0 | — | 23 | NONATHT000028,NONATHT000045…… |
| Mn\_lnc-0012 | 0 | — | 8 | NONATHT000436,NONATHT000970, …… |
| Mn\_lnc-0013 | 0 | — | 8 | NONATHT000436,NONATHT000970…… |
| Mn\_lnc-0014 | 0 | — | 7 | NONATHT000815,NONATHT000984…… |
| Mn\_lnc-0015 | 0 | — | 6 | NONATHT000984,NONATHT001685…… |
| Mn\_lnc-0016 | 0 | — | 5 | NONATHT000414,NONATHT000815…… |
| Mn\_lnc-0017 | 0 | — | 5 | NONATHT000436,NONATHT000984…… |
| Mn\_lnc-0018 | 0 | — | 4 | NONATHT002060,NONATHT003599…… |
| Mn\_lnc-0019 | 0 | — | 3 | NONATHT001309,NONATHT002122…… |
| Mn\_lnc-0020 | 0 | — | 3 | NONATHT000970,NONATHT000984, …… |
| Mn\_lnc-0021 | 0 | — | 3 | NONATHT000436,NONATHT001311…… |
| Mn\_lnc-0022 | 0 | — | 3 | NONATHT000436,NONATHT001311…… |
| Mn\_lnc-0023 | 0 | — | 3 | NONATHT000984,NONATHT001685…… |
| Mn\_lnc-0024 | 0 | — | 3 | NONATHT000984,NONATHT001685…… |
| Mn\_lnc-0025 | 0 | — | 3 | NONATHT000984,NONATHT002122…… |
| Mn\_lnc-0026 | 0 | — | 3 | NONATHT000970,NONATHT001309…… |
| Mn\_lnc-0027 | 0 | — | 3 | NONATHT001730,NONATHT002060…… |
| Mn\_lnc-0028 | 0 | — | 3 | NONATHT000970,NONATHT000984…… |
| Mn\_lnc-0029 | 0 | — | 2 | NONATHT001730,NONATHT002010 |
| Mn\_lnc-0030 | 0 | — | 2 | NONATHT000436,NONATHT003487 |
| Mn\_lnc-0031 | 0 | — | 2 | NONATHT000984,NONATHT002122 |
| Mn\_lnc-0032 | 0 | — | 2 | NONATHT000970,NONATHT003487 |
| Mn\_lnc-0033 | 0 | — | 2 | NONATHT002060,NONATHT003295 |
| Mn\_lnc-0034 | 0 | — | 2 | NONATHT002060,NONATHT003295 |
| Mn\_lnc-0035 | 0 | — | 2 | NONATHT001730,NONATHT002060 |
| Mn\_lnc-0036 | 0 | — | 2 | NONATHT000970,NONATHT001690 |
| Mn\_lnc-0037 | 0 | — | 2 | NONATHT001685,NONATHT002060 |
| Mn\_lnc-0038 | 0 | — | 2 | NONATHT001685,NONATHT002060 |
| Mn\_lnc-0039 | 0 | — | 2 | NONATHT002060,NONATHT003295 |
| Mn\_lnc-0040 | 0 | — | 2 | NONATHT003541,NONATHT003542 |
| Mn\_lnc-0041 | 0 | — | 2 | NONATHT000815,NONATHT001689 |
| Mn\_lnc-0042 | 0 | — | 2 | NONATHT001685,NONATHT002010 |
| Mn\_lnc-0043 | 0 | — | 2 | NONATHT000414,NONATHT001309 |
| Mn\_lnc-0044 | 0 | — | 2 | NONATHT000414,NONATHT001690 |
| Mn\_lnc-0045 | 0 | — | 2 | NONATHT002122,NONATHT003487 |
| Mn\_lnc-0046 | 0 | — | 2 | NONATHT002392,NONATHT003576 |
| Mn\_lnc-0047 | 0 | — | 2 | NONATHT003836,NONATHT003850 |
| Mn\_lnc-0048 | 0 | — | 2 | NONATHT000414,NONATHT003610 |
| Mn\_lnc-0049 | 0 | — | 2 | NONATHT001617,NONATHT003295 |
| Mn\_lnc-0050 | 0 | — | 2 | NONATHT001685,NONATHT003487 |
| Mn\_lnc-0051 | 0 | — | 2 | NONATHT003541,NONATHT003542 |
| Mn\_lnc-0052 | 0 | — | 2 | NONATHT003839,NONATHT003847 |
| Mn\_lnc-0053 | 0 | — | 2 | NONATHT000970,NONATHT000984 |
| Mn\_lnc-0054 | 0 | — | 2 | NONATHT000436,NONATHT002122 |
| Mn\_lnc-0055 | 0 | — | 2 | NONATHT003045,NONATHT003046 |
| Mn\_lnc-0056 | 0 | — | 2 | NONATHT002060,NONATHT003295 |
| Mn\_lnc-0057 | 0 | — | 1 | NONATHT003295 |
| Mn\_lnc-0058 | 0 | — | 1 | NONATHT003144 |
| Mn\_lnc-0059 | 0 | — | 1 | NONATHT001311 |
| Mn\_lnc-0060 | 0 | — | 1 | NONATHT001730 |
| Mn\_lnc-0061 | 0 | — | 1 | NONATHT002060 |
| Mn\_lnc-0062 | 0 | — | 1 | NONATHT002122 |
| Mn\_lnc-0063 | 0 | — | 1 | NONATHT003295 |
| Mn\_lnc-0064 | 0 | — | 1 | NONATHT002779 |
| Mn\_lnc-0065 | 0 | — | 1 | NONATHT000436 |
| Mn\_lnc-0066 | 0 | — | 1 | NONATHT003487 |
| Mn\_lnc-0067 | 0 | — | 1 | NONATHT003295 |
| Mn\_lnc-0068 | 0 | — | 1 | NONATHT001730 |
| Mn\_lnc-0069 | 0 | — | 1 | NONATHT001721 |
| Mn\_lnc-0070 | 0 | — | 1 | NONATHT001721 |
| Mn\_lnc-0071 | 0 | — | 1 | NONATHT003610 |
| Mn\_lnc-0072 | 0 | — | 1 | NONATHT001311 |
| Mn\_lnc-0073 | 0 | — | 1 | NONATHT003295 |
| Mn\_lnc-0074 | 0 | — | 1 | NONATHT001824 |
| Mn\_lnc-0075 | 0 | — | 1 | NONATHT003487 |
| Mn\_lnc-0076 | 0 | — | 1 | NONATHT001311 |
| Mn\_lnc-0077 | 0 | — | 1 | NONATHT001685 |
| Mn\_lnc-0078 | 0 | — | 1 | NONATHT001082 |
| Mn\_lnc-0079 | 0 | — | 1 | NONATHT003295 |
| Mn\_lnc-0080 | 0 | — | 1 | NONATHT000984 |
| Mn\_lnc-0081 | 0 | — | 1 | NONATHT003295 |
| Mn\_lnc-0082 | 0 | — | 1 | NONATHT000984 |
| Mn\_lnc-0083 | 0 | — | 1 | NONATHT002060 |
| Mn\_lnc-0084 | 0 | — | 1 | NONATHT000970 |
| Mn\_lnc-0085 | 0 | — | 1 | NONATHT001721 |
| Mn\_lnc-0086 | 0 | — | 1 | NONATHT002060 |
| Mn\_lnc-0087 | 0 | — | 1 | NONATHT001311 |
| Mn\_lnc-0088 | 0 | — | 1 | NONATHT001311 |
| Mn\_lnc-0089 | 0 | — | 1 | NONATHT001311 |
| Mn\_lnc-0090 | 0 | — | 1 | NONATHT001311 |
| Mn\_lnc-0091 | 0 | — | 1 | NONATHT003295 |
| Mn\_lnc-0092 | 0 | — | 1 | NONATHT001311 |
| Mn\_lnc-0093 | 0 | — | 1 | NONATHT001309 |
| Mn\_lnc-0094 | 0 | — | 1 | NONATHT003732 |
| Mn\_lnc-0095 | 0 | — | 1 | NONATHT001418 |
| Mn\_lnc-0096 | 0 | — | 1 | NONATHT001309 |
| Mn\_lnc-0097 | 0 | — | 1 | NONATHT000984 |
| Mn\_lnc-0098 | 0 | — | 1 | NONATHT000984 |
| Mn\_lnc-0099 | 0 | — | 1 | NONATHT000984 |
| Mn\_lnc-0100 | 0 | — | 1 | NONATHT003295 |
| Mn\_lnc-0101 | 0 | — | 1 | NONATHT001309 |
| Mn\_lnc-0102 | 0 | — | 1 | NONATHT000984 |
| Mn\_lnc-0103 | 0 | — | 1 | NONATHT001800 |
| Mn\_lnc-0104 | 0 | — | 1 | NONATHT000984 |
| Mn\_lnc-0105 | 0 | — | 1 | NONATHT002073 |
| Mn\_lnc-0106 | 0 | — | 1 | NONATHT000984 |
| Mn\_lnc-0107 | 0 | — | 1 | NONATHT000984 |
| Mn\_lnc-0108 | 0 | — | 1 | NONATHT003295 |
| Mn\_lnc-0109 | 0 | — | 1 | NONATHT001721 |
| Mn\_lnc-0110 | 0 | — | 1 | NONATHT002122 |
| Mn\_lnc-0111 | 0 | — | 1 | NONATHT003295 |
| Mn\_lnc-0112 | 0 | — | 1 | NONATHT002122 |
| Mn\_lnc-0113 | 0 | — | 1 | NONATHT002060 |
| Mn\_lnc-0114 | 0 | — | 1 | NONATHT000730 |
| Mn\_lnc-0115 | 0 | — | 1 | NONATHT003295 |

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**Table S3.** Summary of tissue specific genes in root with neighbor coding genes and functions.

|  |  |  |
| --- | --- | --- |
| **LncRNA ID** | **Neighbor Coding Gene** | **Function** |
| Mn\_lnc-0041 | EXB66525.1, EXB66527.1, EXB66528.1 | Phosphatidylinositol-glycan biosynthesis class X protein Mitochondrial inner membrane protease ATP23，hypothetical protein Mitogen-activated protein kinase kinase 5 |
| Mn\_lnc-0042 | EXB72255.1 | hypothetical protein |
| Mn\_lnc-0050 | EXB87346.1 | putative amino acid permease YfnA |
| Mn\_lnc-0053 | EXB93293.1 | hypothetical protein |
| Mn\_lnc-0056 | EXB94111.1 | putative NADH dehydrogenase |
| Mn\_lnc-0057 | EXB95198.1 | hypothetical protein |
| Mn\_lnc-0059 | EXB97271.1 | Non-specific lipid-transfer protein-like protein |
| Mn\_lnc-0065 | EXC06697.1 | Sulfate transporter 1.3 |
| Mn\_lnc-0087 | EXC30709.1 | Werner Syndrome-like exonuclease |
| Mn\_lnc-0091 | EXC31783.1 | 4-coumarate--CoA ligase-like 7 |

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**Table S4.** Summary of tissue specific genes in bark with neighbor coding genes and functions.

|  |  |  |
| --- | --- | --- |
| **LncRNA ID** | **Neighbor Coding Gene** | **Function** |
| Mn\_lnc-0001 | EXB19758.1 | Ethylene-responsive transcription factor |
| Mn\_lnc-0002 | EXB20730.1 | hypothetical protein |
| Mn\_lnc-0003 | EXB23830.1 | Auxin-binding protein ABP19a |
| Mn\_lnc-0011 | EXB38292.1, EXB38293.1 | 60S ribosomal protein L6 Phosphatidylinositol-4-phosphate 5-kinase 4 |
| Mn\_lnc-0014 | EXB39149.1 | hypothetical protein |
| Mn\_lnc-0015 | EXB40022.1 | hypothetical protein |
| Mn\_lnc-0024 | EXB51228.1 | putative LRR receptor-like serine/threonine-protein kinase |
| Mn\_lnc-0037 | EXB62622.1 | UDP-glycosyltransferase 85A |
| Mn\_lnc-0039 | EXB65570.1 | hypothetical protein |
| Mn\_lnc-0043 | EXB72448.1 | Curved DNA-binding protein |
| Mn\_lnc-0049 | EXB82808.1 | Polyphenol oxidase |
| Mn\_lnc-0066 | EXC10952.1 | DNA polymerase III subunit |
| Mn\_lnc-0075 | EXC21500.1 | hypothetical protein |
| Mn\_lnc-0080 | EXC25147.1 | hypothetical protein |
| Mn\_lnc-0082 | EXC25309.1, EXC25310.1 | Isoleucine N-monooxygenase 2 Type II inositol-1%2C4%2C5-trisphosphate 5-phosphatase FRA3 |

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**Table S5.** Summary of tissue specific genes in winter bud with neighbor coding genes and functions.

|  |  |  |
| --- | --- | --- |
| **LncRNA ID** | **Neighbor Coding Gene** | **Function** |
| Mn\_lnc-0004 | EXB26546.1 | hypothetical protein |
| Mn\_lnc-0005 | EXB27067.1, EXB27068.1, EXB27070.1 | putative glutathione S-transferase parC putative glutathione S-transferase putative glutathione S-transferase |
| Mn\_lnc-0009 | EXB36838.1, EXB36842.1 | Dihydrolipoyllysine-residue acetyltransferase component 2 of pyruvate dehydrogenase complex ER lumen protein retaining receptor |
| Mn\_lnc-0012 | EXB38871.1 | Casein kinase I isoform delta-like protein |
| Mn\_lnc-0017 | EXB41586.1 | F-box protein |
| Mn\_lnc-0025 | EXB51430.1 | hypothetical protein |
| Mn\_lnc-0028 | EXB53993.1 | GC-rich sequence DNA-binding factor 1 |
| Mn\_lnc-0030 | EXB54658.1 | hypothetical protein |
| Mn\_lnc-0031 | EXB54658.1 | hypothetical protein |
| Mn\_lnc-0032 | EXB54869.1 | hypothetical protein |
| Mn\_lnc-0033 | EXB54966.1 | hypothetical protein |
| Mn\_lnc-0034 | EXB56221.1 | NAD(P)H-quinone oxidoreductase subunit 3 |
| Mn\_lnc-0035 | EXB56922.1 | hypothetical protein |
| Mn\_lnc-0051 | EXB88542.1 | Glutamate receptor 2.7 |
| Mn\_lnc-0052 | EXB93127.1,EXB93128.1 | GDSL esterase/lipase, G-type lectin S-receptor-like serine/threonine-protein kinase |
| Mn\_lnc-0063 | EXC01148.1 | MATE efflux family protein 5 |
| Mn\_lnc-0067 | EXC11337.1 | Wall-associated receptor kinase-like 20 |
| Mn\_lnc-0070 | EXC16242.1 | Ubiquitin-40S ribosomal protein S27a |
| Mn\_lnc-0071 | EXC19487.1, EXC19488.1 | 40S ribosomal protein S23 RNA-binding protein 38 |
| Mn\_lnc-0073 | EXC20544.1 | MATE efflux family protein DTX1 |
| Mn\_lnc-0078 | EXC24413.1 | hypothetical protein |
| Mn\_lnc-0085 | EXC28022.1 | Protein ycf2 |
| Mn\_lnc-0086 | EXC28324.1 | hypothetical protein |
| Mn\_lnc-0099 | EXC43199.1 | Legumin A |

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**Table S6.** Summary of tissue specific genes in leaf with neighbor coding genes and functions.

|  |  |  |
| --- | --- | --- |
| **LncRNA ID** | **Neighbor Coding Gene** | **Function** |
| Mn\_lnc-0021 | EXB50291.1 | hypothetical protein |
| Mn\_lnc-0022 | EXB50426.1 | putative beta-1%2C3-galactosyltransferase 2 |
| Mn\_lnc-0023 | EXB50920.1, EXB50928.1, EXB50929.1 | (-)-germacrene D synthase (-)-germacrene D synthase Pinene synthase |
| Mn\_lnc-0027 | EXB53001.1 | Trafficking protein particle complex subunit 4 |
| Mn\_lnc-0036 | EXB58188.1 | ,60S acidic ribosomal protein P0 |
| Mn\_lnc-0038 | EXB65076.1 | Heparanase-like protein 2 |
| Mn\_lnc-0044 | ,EXB72462.1 | Cation/H(+) antiporter 4 |
| Mn\_lnc-0045 | EXB75948.1 | Wall-associated receptor kinase-like 20 |
| Mn\_lnc-0047 | EXB80599.1 | Zinc finger protein MAGPIE |
| Mn\_lnc-0058 | EXB96347.1 | hypothetical protein |
| Mn\_lnc-0060 | EXB97676.1, EXB97677.1 | 3-oxo-5-alpha-steroid 4-dehydrogenase 2; 3-oxo-5-alpha-steroid 4-dehydrogenase 2 |
| Mn\_lnc-0074 | EXC21270.1 | hypothetical protein |
| Mn\_lnc-0077 | EXC23523.1 | hypothetical protein |
| Mn\_lnc-0081 | EXC25272.1 | hypothetical protein |
| Mn\_lnc-0088 | EXC30765.1 | hypothetical protein |
| Mn\_lnc-0089 | EXC30689.1 | hypothetical protein |
| Mn\_lnc-0090 | EXC31504.1, | putative receptor-like protein kinase |
| Mn\_lnc-0094 | EXC35008.1 | hypothetical protein |

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**Table S7**. Summary of tissue specific genes in male flower with neighbor coding genes and functions.

|  |  |  |
| --- | --- | --- |
| **LncRNA ID** | **Neighbor Coding Gene** | **Function** |
| Mn\_lnc-0006 | EXB28594.1, EXB28598.1 | Protein PROLIFERA DNA mismatch repair protein Msh6-1 |
| Mn\_lnc-0007 | EXB29133.1 | DnAJ-like protein |
| Mn\_lnc-0008 | EXB29871.1 | hypothetical protein |
| Mn\_lnc-0013 | EXB38817.1 | hypothetical protein |
| Mn\_lnc-0019 | EXB48303.1 | Puromycin-sensitive aminopeptidase |
| Mn\_lnc-0048 | EXB81017.1 | Serine/threonine-protein phosphatase PP1 |
| Mn\_lnc-0054 | EXB93909.1 | ,Protein lap1 |
| Mn\_lnc-0061 | EXB97657.1 | Cysteine-rich receptor-like protein kinase 25 |
| Mn\_lnc-0068 | EXC13800.1 | hypothetical protein; |
| Mn\_lnc-0069 | EXC14254.1 | hypothetical protein; |
| Mn\_lnc-0072 | EXC20310.1, EXC20311.1, EXC20313.1 | Phosphoenolpyruvate/phosphate translocator 2 tRNA (adenine(58)-N(1))-methyltransferase catalytic subunit Isocitrate dehydrogenase [NADP] |
| Mn\_lnc-0079 | EXC24965.1 | GDSL esterase/lipase |
| Mn\_lnc-0084 | EXC25828.1 | ,Katanin p60 ATPase-containing subunit |
| Mn\_lnc-0092 | EXC32126.1 | Elongation factor 1-delta;,Cation/H(+) antiporter 15 |
| Mn\_lnc-0095 | EXC35507.1 | Purple acid phosphatase 25 |

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