Appendix S2. Ray scaffold locations and top BLASTP hits for the additional 125 bait sequences.

|  |  |
| --- | --- |
| *A. camansi* gene | Top BLASTP hit |
| Name |  | Exon coordinates (strand) | GI | OHR | *E*-value | Title |
| mb001 | scaffold-1657 | 166–599 (+) | 519898277 | 1.00 | 4.00E-34 | K-box region and MADS-box transcription factor family protein, partial [*Glycine max*] |
| mb002 | scaffold-2344 | 2409–2595, 2687–3121 (-) | 587938012 | 0.26 | 8.00E-38 | MADS-box protein FBP24 [*Morus notabilis*] |
| mb003 | scaffold-2360 | 2966–3225, 3331–3412 (+) | 587834458 | 0.57 | 1.00E-52 | Agamous-like MADS-box protein AGL16 [*Morus notabilis*] |
| mb004 | scaffold-4283 | 1083–1414, 1584–1741, 2091–2132 (-) | 587899828 | 0.31 | 1.00E-29 | Agamous-like MADS-box protein AGL11 [*Morus notabilis*] |
| mb005 | scaffold-7327 | 1026–2050 (+) | 587942863 | 1.05 | 6.00E-105 | Agamous-like MADS-box protein AGL62 [*Morus notabilis*] |
| mb006 | scaffold-9167 | 74–409 (+) | 587939681 | 0.31 | 6.00E-10 | MADS-box protein SOC1 [*Morus notabilis*] |
| mb007 | scaffold-14352 | 2437–3711 (-) | 587868023 | 0.49 | 2.00E-66 | Agamous-like MADS-box protein AGL97 [*Morus notabilis*] |
| mb008 | scaffold-14947 | 1–395, 506–618, 1426–1510, 1631–1675, 2168–2411 (-) | 587900830 | 0.47 | 1.00E-44 | Agamous-like MADS-box protein AGL8-like protein [*Morus notabilis*] |
| mb009 | scaffold-17451 | 1505–3364, 3990–4051 (-) | 587933572 | 0.98 | 6.00E-150 | MADS-box transcription factor PHERES 1 [*Morus notabilis*] |
| mb010 | scaffold-19703 | 1556–2703 (+) | 587868182 | 1.00 | 3.00E-116 | Agamous-like MADS-box protein AGL80 [*Morus notabilis*] |
| mb011 | scaffold-20036 | 2346–2752 (+) | 587872298 | 0.28 | 2.00E-39 | Agamous-like MADS-box protein AGL62 [*Morus notabilis*] |
| mb012 | scaffold-23088 | 649–1429, 1515–2341 (-) | 587947051 | 1.01 | 2.00E-144 | MADS-box protein [*Morus notabilis*] |
| mb013 | scaffold-24802 | 1446–2773 (+) | 587930139 | 0.98 | 1.00E-66 | Agamous-like MADS-box protein AGL61 [*Morus notabilis*] |
| mb014 | scaffold-29024 | 330–665, 1064–1105, 1988–2152, 2292–2358, 2752–2781 (-) | 587938012 | 0.76 | 9.00E-76 | MADS-box protein FBP24 [*Morus notabilis*] |
| mb015 | scaffold-30246 | 1–622, 701–952, 2044–2241 (-) | 587924418 | 0.24 | 4.00E-34 | MADS-box transcription factor 6 [*Morus notabilis*] |
| mb016 | scaffold-31966 | 676–816, 922–1481 (+) | 225455667 | 0.29 | 1.00E-32 | PREDICTED: MADS-box protein SOC1-like [*Vitis vinifera*] |
| mb017 | scaffold-32402 | 1880–2731 (-) | 587942861 | 0.98 | 2.00E-91 | Agamous-like MADS-box protein AGL61 [*Morus notabilis*] |
| mb018 | scaffold-32402 | 4894–5413, 5630–5891 (-) | 587942863 | 0.31 | 4.00E-23 | Agamous-like MADS-box protein AGL62 [*Morus notabilis*] |
| mb019 | scaffold-35005 | 4912–5345, 5629–5841 (-) | 587913365 | 0.53 | 6.00E-36 | MADS-box transcription factor 27 [*Morus notabilis*] |
| mb020 | scaffold-35240 | 235–334, 935–976, 1801–1842, 1963–2120, 2738–2919 (+) | 587924418 | 0.58 | 7.00E-82 | MADS-box transcription factor 6 [*Morus notabilis*] |
| mb021 | scaffold-37879 | 3236–4351 (-) | 587881704 | 0.98 | 3.00E-105 | Agamous-like MADS-box protein AGL80 [*Morus notabilis*] |
| mb022 | scaffold-38271 | 2234–4121 (-) | 587925724 | 0.98 | 5.00E-173 | Agamous-like MADS-box protein AGL36 [*Morus notabilis*] |
| mb023 | scaffold-39068 | 621–662, 879–1195 (+) | 587929496 | 0.25 | 6.00E-13 | MADS-box protein SVP [*Morus notabilis*] |
| mb024 | scaffold-39170 | 66–94, 264–375, 1841–1934, 2978–3039, 3157–3274, 3988–4029, 4788–4829, 4954–5261 (+) | 587924418 | 0.67 | 4.00E-87 | MADS-box transcription factor 6 [*Morus notabilis*] |
| mb025 | scaffold-40947 | 541–723, 2920–3111 (-) | 225455667 | 0.29 | 3.00E-30 | PREDICTED: MADS-box protein SOC1-like [*Vitis vinifera*] |
| mb026 | scaffold-41218 | 966–1272, 1539–1729, 1963–2117, 2403–2450, 2535–2634 (-) | 587917239 | 0.61 | 2.00E-85 | MADS-box transcription factor 3 [*Morus notabilis*] |
| mb027 | scaffold-44908 | 1325–1968, 2071–2190 (-) | 587942863 | 0.43 | 2.00E-09 | Agamous-like MADS-box protein AGL62 [*Morus notabilis*] |
| mb028 | scaffold-45897 | 1023–1122, 1214–1255, 1355–1396, 1902–2014, 2250–2510 (+) | 587900830 | 0.44 | 4.00E-50 | Agamous-like MADS-box protein AGL8-like protein [*Morus notabilis*] |
| mb029 | scaffold-47193 | 1270–1460, 1799–1971, 2071–2112, 2220–2471 (-) | 587907291 | 0.39 | 3.00E-60 | Agamous-like MADS-box protein AGL15 [*Morus notabilis*] |
| mb030 | scaffold-47324 | 1785–2741 (-) | 587917454 | 1.04 | 5.00E-59 | Agamous-like MADS-box protein AGL93 [*Morus notabilis*] |
| mb031 | scaffold-48806 | 1–401 (-) | 587945785 | 0.26 | 8.00E-33 | MADS-box transcription factor 6 [*Morus notabilis*] |
| mb032 | scaffold-49670 | 295–394, 772–833, 932–959, 1078–1139, 1304–1322, 1432–1485, 1961–2271 (-) | 359480834 | 0.93 | 5.00E-94 | PREDICTED: MADS-box transcription factor 18-like [*Vitis vinifera*] |
| mb033 | scaffold-56115 | 33–181, 585–841 (-) | 587942863 | 0.21 | 2.00E-22 | Agamous-like MADS-box protein AGL62 [*Morus notabilis*] |
| mb034 | scaffold-58008 | 4132–4231, 4337–4378, 4467–4550, 4698–4879, 4987–5172 (+) | 587946686 | 0.50 | 1.00E-63 | MADS-box transcription factor 27 [*Morus notabilis*] |
| mb035 | scaffold-63607 | 309–961 (-) | 587930681 | 0.82 | 8.00E-37 | MADS-box protein SOC1 [*Morus notabilis*] |
| mb036 | scaffold-64679 | 1–291 (-) | 316890746 | 0.50 | 1.00E-30 | MADS-box protein AGL2 subfamily [*Coffea arabica*] |
| mb037 | scaffold-65472 | 2762–3871 (-) | 587847952 | 1.09 | 5.00E-71 | Agamous-like MADS-box protein AGL62 [*Morus notabilis*] |
| mb038 | scaffold-71228 | 278–1451 (-) | 587881704 | 0.97 | 4.00E-110 | Agamous-like MADS-box protein AGL80 [*Morus notabilis*] |
| mb039 | scaffold-71532 | 273–322, 433–530, 818–1006 (+) | 587919217 | 0.33 | 4.00E-35 | MADS-box transcription factor 13 [*Morus notabilis*] |
| mb040 | scaffold-73395 | 101–438, 1062–1073, 1151–1491 (-) | 449531683 | 0.98 | 3.00E-36 | PREDICTED: agamous-like MADS-box protein AGL8-like [*Cucumis sativus*] |
| mb041 | scaffold-73661 | 646–2245 (+) | 587947051 | 1.04 | 1.00E-173 | MADS-box protein [*Morus notabilis*] |
| mb042 | scaffold-73790 | 1051–1591 (-) | 587872297 | 0.56 | 8.00E-39 | Agamous-like MADS-box protein AGL62 [*Morus notabilis*] |
| mb043 | scaffold-74914 | 785–846, 996–1095, 1506–1547, 1682–1723, 2471–2928 (+) | 587924420 | 0.91 | 3.00E-74 | MADS-box protein SOC1 [*Morus notabilis*] |
| mb044 | scaffold-76861 | 849–1281 (+) | 357446033 | 0.85 | 2.00E-35 | MADS-box family transcription factor [*Medicago truncatula*] |
| mb045 | scaffold-81123 | 2306–3061 (+) | 587888968 | 0.87 | 9.00E-35 | Agamous-like MADS-box protein AGL8-like protein [*Morus notabilis*] |
| mb046 | scaffold-83177 | 1670–1905, 2156–2232, 2678–2719, 2858–2957, 3054–3201 (-) | 587834458 | 0.53 | 1.00E-30 | Agamous-like MADS-box protein AGL16 [*Morus notabilis*] |
| mb047 | scaffold-84607 | 3684–4156, 4314–4471, 4725–4766, 4886–4981 (-) | 27763670g | 0.35 | 3.00E-37 | MADS-box transcription factor [*Momordica charantia*] |
| mb048 | scaffold-88291 | 1–611 (-) | 587942861 | 0.98 | 9.00E-93 | Agamous-like MADS-box protein AGL61 [*Morus notabilis*] |
| mb049 | scaffold-90628 | 132–455, 1501–1637, 1869–1910, 2020–2101 (-) | 587850123 | 0.40 | 2.00E-58 | Agamous-like MADS-box protein AGL9-like protein [*Morus notabilis*] |
| mb050 | scaffold-92003 | 1–1131 (-) | 440548641 | 0.95 | 3.00E-32 | MADS-box transcription factor [*Larix sibirica*] |
| mb051 | scaffold-92143 | 401–1531 (-) | 587881704 | 0.95 | 1.00E-107 | Agamous-like MADS-box protein AGL80 [*Morus notabilis*] |
| mb052 | scaffold-92850 | 1–611 (-) | 587866411 | 0.26 | 8.00E-26 | Agamous-like MADS-box protein AGL62 [*Morus notabilis*] |
| mb053 | scaffold-93337 | 211–1951 (-) | 587925724 | 0.89 | 2.00E-149 | Agamous-like MADS-box protein AGL36 [*Morus notabilis*] |
| mb054 | scaffold-99714 | 686–767, 866–1071 (-) | 587834458 | 0.57 | 5.00E-53 | Agamous-like MADS-box protein AGL16 [*Morus notabilis*] |
| mb055 | scaffold-107282 | 1–1069 (+) | 587868182 | 0.95 | 1.00E-89 | Agamous-like MADS-box protein AGL80 [*Morus notabilis*] |
| mb056 | scaffold-110809 | 71–217, 306–497 (+) | 587867915 | 0.21 | 3.00E-13 | MADS-box protein JOINTLESS [*Morus notabilis*] |
| mb057 | scaffold-115399 | 816–2318 (+) | 449531683 | 0.97 | 5.00E-36 | PREDICTED: agamous-like MADS-box protein AGL8-like [*Cucumis sativus*] |
| mb058 | scaffold-119292 | 146–312, 1044–1114, 1710–1809, 1921–1962, 2475–2516, 2605–2732, 3156–3367 (+) | 32478023g | 0.75 | 3.00E-50 | FRUITFULL-like MADS-box [*Heuchera americana*] |
| mb059 | scaffold-120326 | 82–268, 398–495, 608–658, 729–828, 965–1026, 1150–1177, 1292–1353, 1453–1471, 1594–1771 (-) | 587919217 | 0.54 | 9.00E-96 | MADS-box transcription factor 13 [*Morus notabilis*] |
| mb060 | scaffold-124343 | 434–512, 2203–2270, 2359–2716 (+) | 587850123 | 0.34 | 1.00E-44 | Agamous-like MADS-box protein AGL9-like protein [*Morus notabilis*] |
| mb061 | scaffold-145035 | 126–523 (+) | 587919217 | 0.15 | 5.00E-11 | MADS-box transcription factor 13 [*Morus notabilis*] |
| mb062 | scaffold-146750 | 56–137, 234–554 (+) | 645234937 | 0.97 | 0.046 | PREDICTED: MADS-box protein FLOWERING LOCUS C-like [*Prunus mume*] |
| mb063 | scaffold-150563 | 1313–1367, 1454–1544, 2279–2581 (-) | 587892925 | 0.34 | 3.00E-56 | Agamous-like MADS-box protein AGL12 [*Morus notabilis*] |
| mb064 | scaffold-151659 | 54–193, 1317–1626 (+) | 587850123 | 0.29 | 2.00E-35 | Agamous-like MADS-box protein AGL9-like protein [*Morus notabilis*] |
| mb065 | scaffold-153683 | 461–766, 842–1016 (+) | 587942862 | 0.68 | 6.00E-22 | Agamous-like MADS-box protein AGL62 [*Morus notabilis*] |
| mb066 | scaffold-154414 | 244–661 (-) | 587837684 | 0.33 | 1.00E-26 | Floral homeotic protein PMADS 2 [*Morus notabilis*] |
| mb067 | scaffold-160214 | 356–1013 (+) | 587872559 | 0.43 | 3.00E-32 | MADS-box transcription factor 1 [*Morus notabilis*] |
| mb068 | scaffold-160753 | 621–1027, 1453–1531 (-) | 587867914 | 0.34 | 1.00E-33 | MADS-box protein AGL24 [*Morus notabilis*] |
| mb069 | scaffold-162676 | 269–921 (-) | 587924421 | 0.67 | 2.00E-35 | MADS-box transcription factor 50 [*Morus notabilis*] |
| mb070 | scaffold-165255 | 209–681 (-) | 587942863 | 0.60 | 7.00E-42 | Agamous-like MADS-box protein AGL62 [*Morus notabilis*] |
| mb071 | scaffold-165372 | 74–265, 550–660, 1137–1193, 1309–1327, 1481–1545, 1641–1656, 1742–1803, 1899–1995, 2267–2311, 2402–2529, 2948–3317, 3409–3576, 3817–4043 (+) | 587932994 | 1.06 | 0 | MADS-box transcription factor 16 [*Morus notabilis*] |
| mb072 | scaffold-171384 | 685–881 (-) | 587889184 | 0.22 | 5.00E-24 | Agamous-like MADS-box protein AGL8 [*Morus notabilis*] |
| mb073 | scaffold-173381 | 506–1936 (+) | 587872298 | 1.01 | 1.00E-107 | Agamous-like MADS-box protein AGL62 [*Morus notabilis*] |
| mb074 | scaffold-174915 | 47–146, 241–305, 955–1027 (-) | 587907291 | 0.32 | 4.00E-41 | Agamous-like MADS-box protein AGL15 [*Morus notabilis*] |
| mb075 | scaffold-176584 | 1135–1631 (-) | 81238288g | 0.26 | 7.00E-35 | MADS-box protein [*Taihangia rupestris*] |
| mb076 | scaffold-184191 | 1–681 (-) | 587917454 | 0.96 | 3.00E-56 | Agamous-like MADS-box protein AGL93 [*Morus notabilis*] |
| mb077 | scaffold-185687 | 566–1589 (+) | 587847952 | 1.09 | 2.00E-75 | Agamous-like MADS-box protein AGL62 [*Morus notabilis*] |
| mb078 | scaffold-189357 | 713–766, 956–1241 (-) | 568837954 | 0.72 | 2.00E-47 | PREDICTED: MADS-box transcription factor 15-like [*Citrus sinensis*] |
| mb079 | scaffold-193767 | 274–315, 460–559, 918–1081 (-) | 587929496 | 0.23 | 4.00E-26 | MADS-box protein SVP [*Morus notabilis*] |
| mb080 | scaffold-205335 | 1–79, 177–276, 372–441, 1102–1403 (+) | 587907291 | 0.31 | 2.00E-26 | Agamous-like MADS-box protein AGL15 [*Morus notabilis*] |
| mb081 | scaffold-217873 | 1–831 (-) | 587866408 | 1.02 | 4.00E-88 | Agamous-like MADS-box protein AGL62 [*Morus notabilis*] |
| mb082 | scaffold-219955 | 494–687, 903–1054, 1337–1384, 1514–1613, 1759–1820, 1927–1954, 2052–2121 (-) | 587917239 | 0.63 | 3.00E-81 | MADS-box transcription factor 3 [*Morus notabilis*] |
| mb083 | scaffold-226833 | 1–713 (-) | 587868182 | 0.94 | 2.00E-102 | Agamous-like MADS-box protein AGL80 [*Morus notabilis*] |
| mb084 | scaffold-240128 | 1577–2062 (+) | 587938012 | 0.26 | 2.00E-20 | MADS-box protein FBP24 [*Morus notabilis*] |
| mb085 | scaffold-244092 | 176–484 (+) | 316890746 | 0.50 | 1.00E-30 | MADS-box protein AGL2 subfamily [*Coffea arabica*] |
| mb086 | scaffold-244979 | 219–461 (-) | 316890746 | 0.56 | 2.00E-34 | MADS-box protein AGL2 subfamily [*Coffea arabica*] |
| mb087 | scaffold-257216 | 1197–1810 (-) | 587866407 | 0.65 | 7.00E-45 | Agamous-like MADS-box protein AGL62 [*Morus notabilis*] |
| mb088 | scaffold-282378 | 169–351, 517–641 (-) | 225455667 | 0.29 | 1.00E-30 | PREDICTED: MADS-box protein SOC1-like [*Vitis vinifera*] |
| mb089 | scaffold-351522 | 3186–4047 (+) | 587867914 | 0.34 | 4.00E-34 | MADS-box protein AGL24 [*Morus notabilis*] |
| mb090 | scaffold-351577 | 54–284 (+) | 52219460g | 0.24 | 6.00E-24 | MADS4 [*Prunus persica*] |
| mb091 | scaffold-366542 | 321–907 (-) | 587942862 | 0.51 | 1.00E-40 | Agamous-like MADS-box protein AGL62 [*Morus notabilis*] |
| mb092 | scaffold-368038 | 1767–2009, 2802–2858, 2940–2958, 3087–3151, 3242–3257, 3344–3405, 3500–3596, 3877–3921, 4012–4139, 4565–4937, 5036–5375 (+) | 587932994 | 0.96 | 0 | MADS-box transcription factor 16 [*Morus notabilis*] |
| mb093 | scaffold-369186 | 1–551 (-) | 587942862 | 0.30 | 2.00E-36 | Agamous-like MADS-box protein AGL62 [*Morus notabilis*] |
| mb094 | scaffold-369186 | 1780–2711 (-) | 587942863 | 0.53 | 2.00E-39 | Agamous-like MADS-box protein AGL62 [*Morus notabilis*] |
| mb095 | scaffold-376598 | 926–1362 (+) | 449529148 | 0.72 | 5.00E-37 | PREDICTED: agamous-like MADS-box protein AGL12-like, partial [*Cucumis sativus*] |
| mb096 | scaffold-383686 | 76–117, 205–288, 494–663, 772–975 (+) | 587946686 | 0.36 | 1.00E-42 | MADS-box transcription factor 27 [*Morus notabilis*] |
| mb097 | scaffold-385393 | 1371–1470, 1732–2161 (-) | 587889184 | 0.22 | 2.00E-26 | Agamous-like MADS-box protein AGL8 [*Morus notabilis*] |
| mb098 | scaffold-389621 | 4936–5234, 5417–5558, 5665–5961 (+) | 587917454 | 0.30 | 7.00E-12 | Agamous-like MADS-box protein AGL93 [*Morus notabilis*] |
| v001 | scaffold-16353 | 188–389, 631–886, 969–1350, 1504–1719, 1791–1929, 2537–2785, 2997–3546 (+) | 587848860 | 0.97 | 0 | Myrcene synthase [*Morus notabilis*] |
| v002 | scaffold-48735 | 736–1063, 1185–1470, 1710–2088, 2199–2556, 2647–2895 (+) | 587936332 | 0.83 | 0 | R-linalool synthase [*Morus notabilis*] |
| v003 | scaffold-360968 | 1–316, 424–672, 797–935, 1121–1339, 1696–1981, 3957–4211 (-) | 587936327 | 0.83 | 0 | Isoprene synthase [*Morus notabilis*] |
| v004 | scaffold-3299 | 187–435, 816–951, 1198–1416, 2031–2412, 2815–3088, 3833–4031 (-) | 645240864 | 0.81 | 0 | PREDICTED: (-)-alpha-pinene synthase-like [*Prunus mume*] |
| v005 | scaffold-70158 | 183–401, 537–672, 790–1005, 1953–2366 (+) | 587886521 | 0.63 | 4.00E-136 | (-)-germacrene D synthase [*Morus notabilis*] |
| v006 | scaffold-17096 | 487–705, 801–939, 1067–1315, 3290–3919 (+) | 587886433 | 0.51 | 0 | (3S,6E)-nerolidol synthase 1 [*Morus notabilis*] |
| v007 | scaffold-13624 | 26–306, 400–783, 1028–1128, 1301–1444, 1530–1659 (-) | 627761272 | 0.60 | 1.00E-130 | flavonoid 3′-hydroxylase 2 [*Morus notabilis*] |
| v008 | scaffold-4480 | 1176–1971 (-) | 627761270 | 0.41 | 2.00E-131 | flavonoid 3′-hydroxylase 1 [*Morus notabilis*] |
| v009 | scaffold-21853 | 498–1182, 2870–3058, 4308–4863 (+) | 568883573 | 0.52 | 1.00E-91 | PREDICTED: geraniol 8-hydroxylase-like [*Citrus sinensis*] |
| v010 | scaffold-18340 | 1030–1712 (+) | 470132082 | 0.77 | 5.00E-93 | PREDICTED: flavonoid 3′,5′-hydroxylase-like [*Fragaria vesca* subsp. *vesca*] |
| v011 | scaffold-384884 | 737–1537, 2306–3251 (-) | 587947840 | 0.97 | 0 | (*S*)-*N*-methylcoclaurine 3′-hydroxylase isozyme 2 [*Morus notabilis*] |
| v012 | scaffold-192645 | 866–1686, 1765–1898, 2555–3125 (+) | 587908602 | 0.94 | 0 | Trans-cinnamate 4-monooxygenase [*Morus notabilis*] |
| v013 | scaffold-183437 | 4577–5466, 5811–6831 (-) | 587893901 | 1.01 | 0 | Putative carboxylesterase 11 [*Morus notabilis*] |
| v014 | scaffold-343381 | 1746–1931, 2665–2709, 2789–3545 (+) | 587892232 | 0.20 | 2.00E-159 | E3 ubiquitin-protein ligase RKP [*Morus notabilis*] |
| v015 | scaffold-55579 | 1–931 (-) | 657972761 | 0.57 | 8.00E-146 | PREDICTED: isoflavone 2′-hydroxylase-like [*Malus domestica*] |
| v016 | scaffold-128260 | 251–632, 781–999, 1101–1239, 1350–1803 (+) | 590625479 | 0.56 | 1.00E-138 | 3S-linalool/(E)-nerolidol /(E,E)-geranyl linalool synthase, putative [*Theobroma cacao*] |
| v017 | scaffold-39613 | 116–1533, 2196–2572 (+) | 645239530 | 1.00 | 2.00E-154 | PREDICTED: BAHD acyltransferase At5g47980-like [*Prunus mume*] |
| v018 | scaffold-161447 | 1586–2918, 3959–4048, 4139–4679 (+) | 587893893 | 0.99 | 0 | Salutaridinol 7-*O*-acetyltransferase [*Morus notabilis*] |
| v019 | scaffold-110427 | 1–1493 (-) | 587893888 | 0.99 | 0 | Deacetylvindoline *O*-acetyltransferase [*Morus notabilis*] |
| v020 | scaffold-387356 | 2936–4659 (+) | 587893898 | 1.00 | 0 | Vinorine synthase [*Morus notabilis*] |
| v021 | scaffold-78066 | 1646–3620 (+) | 587917112 | 1.01 | 0 | Taxadien-5-alpha-ol *O*-acetyltransferase [*Morus notabilis*] |
| v022 | scaffold-44341 | 1–615, 690–2161 (-) | 587930142 | 1.00 | 0 | Putative acetyltransferase [*Morus notabilis*] |
| v023 | scaffold-5070 | 279–1528 (-) | 587835357 | 0.70 | 4.00E-147 | Hydroxycinnamoyl-Coenzyme A shikimate/quinate hydroxycinnamoyltransferase [*Morus notabilis*] |
| v024 | scaffold-60973 | 1926–3662 (+) | 587868836 | 0.99 | 0 | BAHD acyltransferase DCR [*Morus notabilis*] |
| v025 | scaffold-29980 | 1–859, 1566–1671 (-) | 52139953 | 0.56 | 4.00E-90 | Alcohol acyl transferase [*Malus domestica*] |
| v026 | scaffold-302676 | 1381–2635 (-) | 587939571 | 0.64 | 1.00E-156 | Benzyl alcohol *O*-benzoyltransferase [*Morus notabilis*] |
| v027 | scaffold-88804 | 1–203, 571–799, 1649–1783, 1867–2111 (-) | 587873318 | 0.80 | 2.00E-150 | Hypothetical protein L484\_008312 [*Morus notabilis*] |