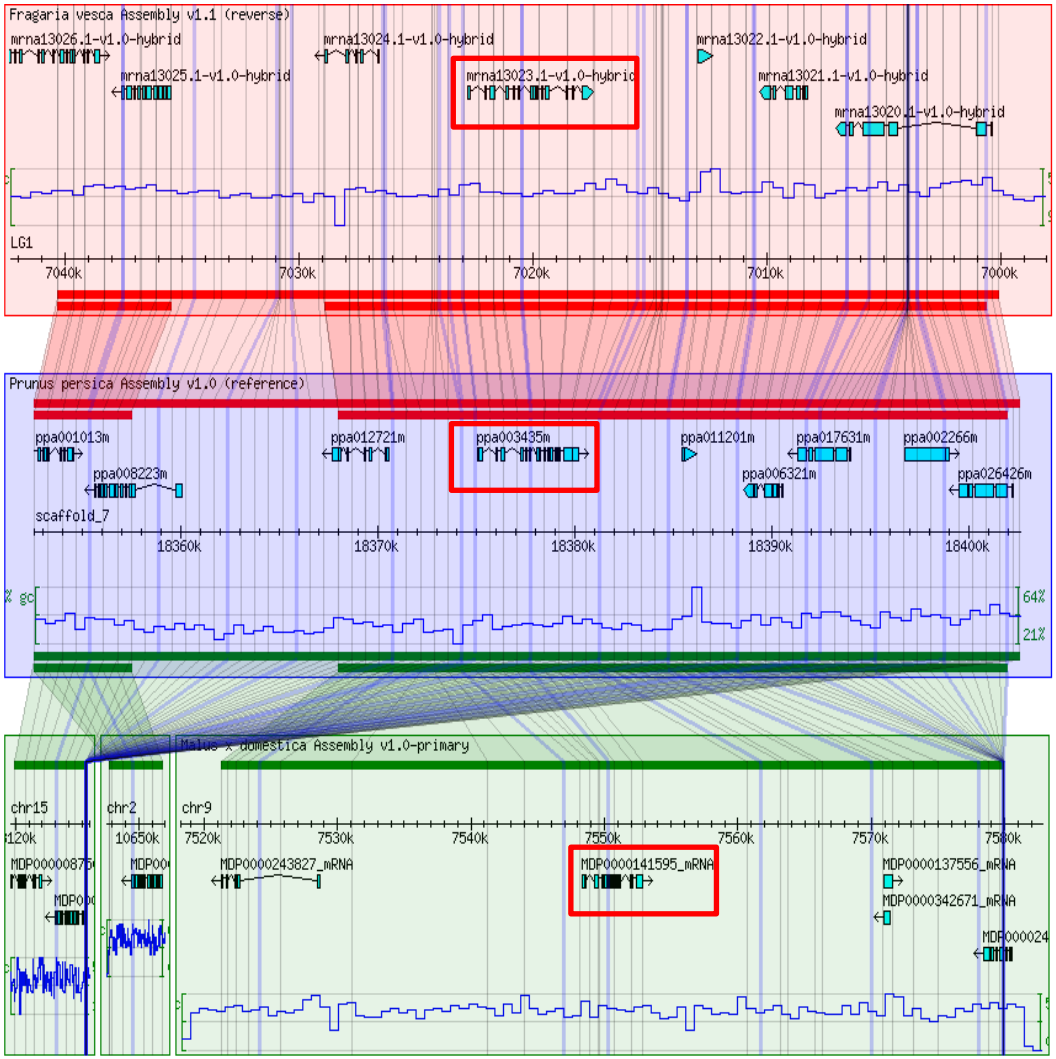
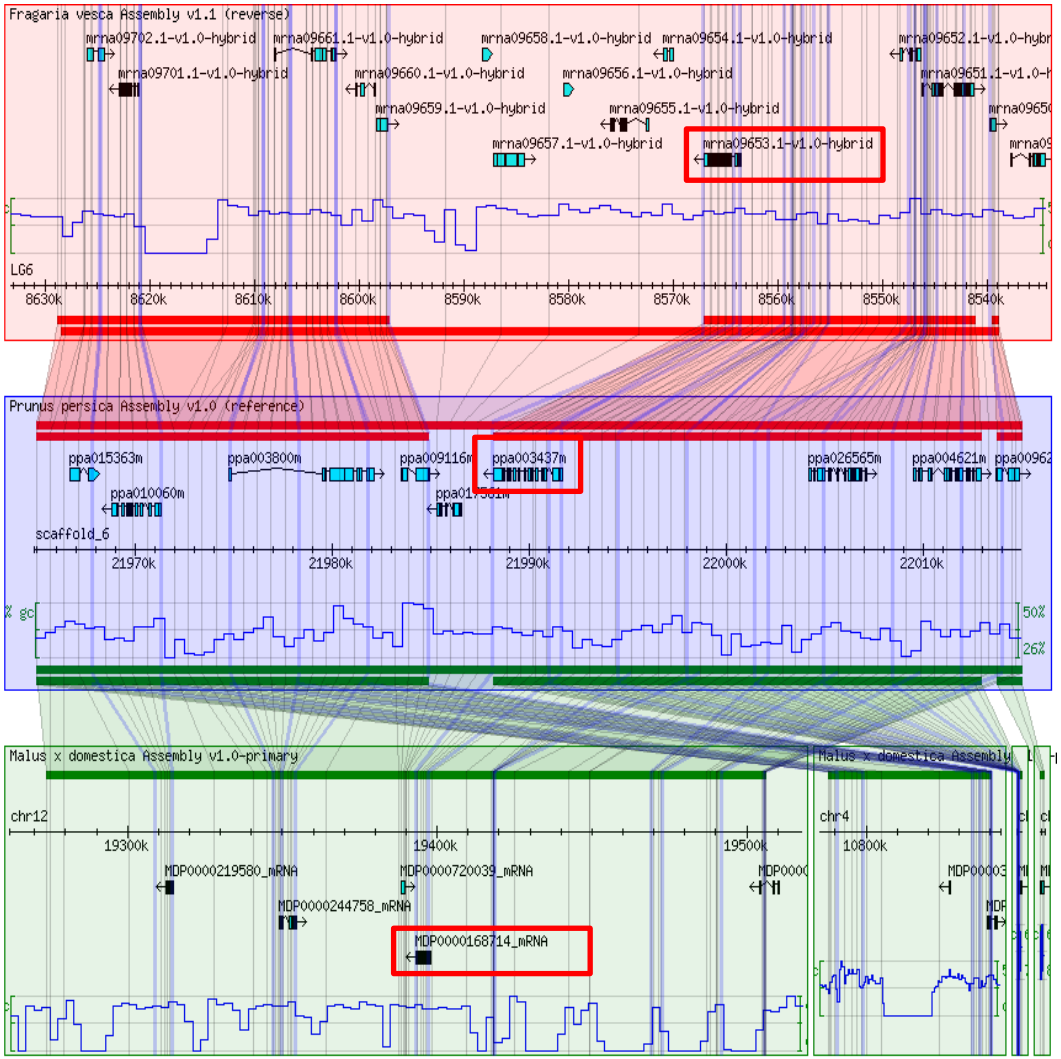
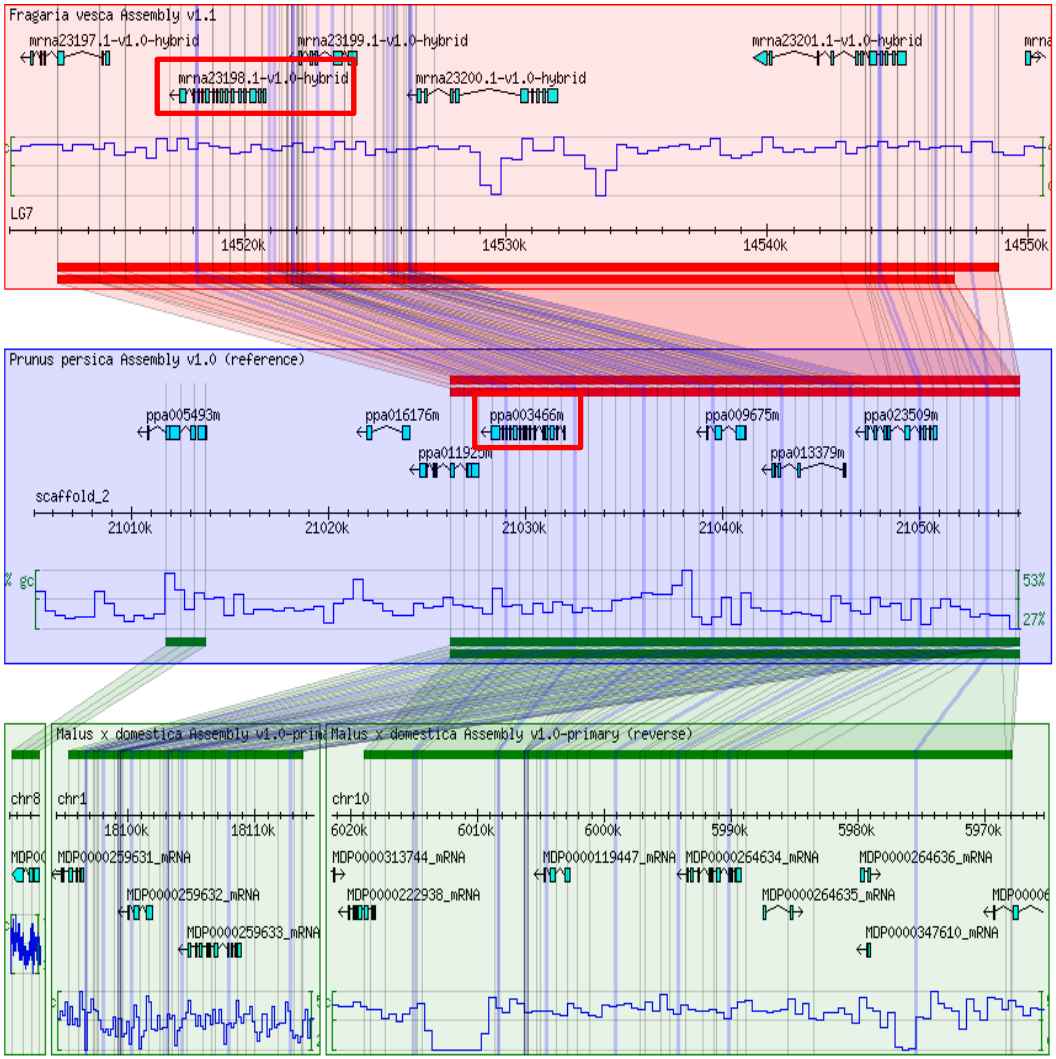
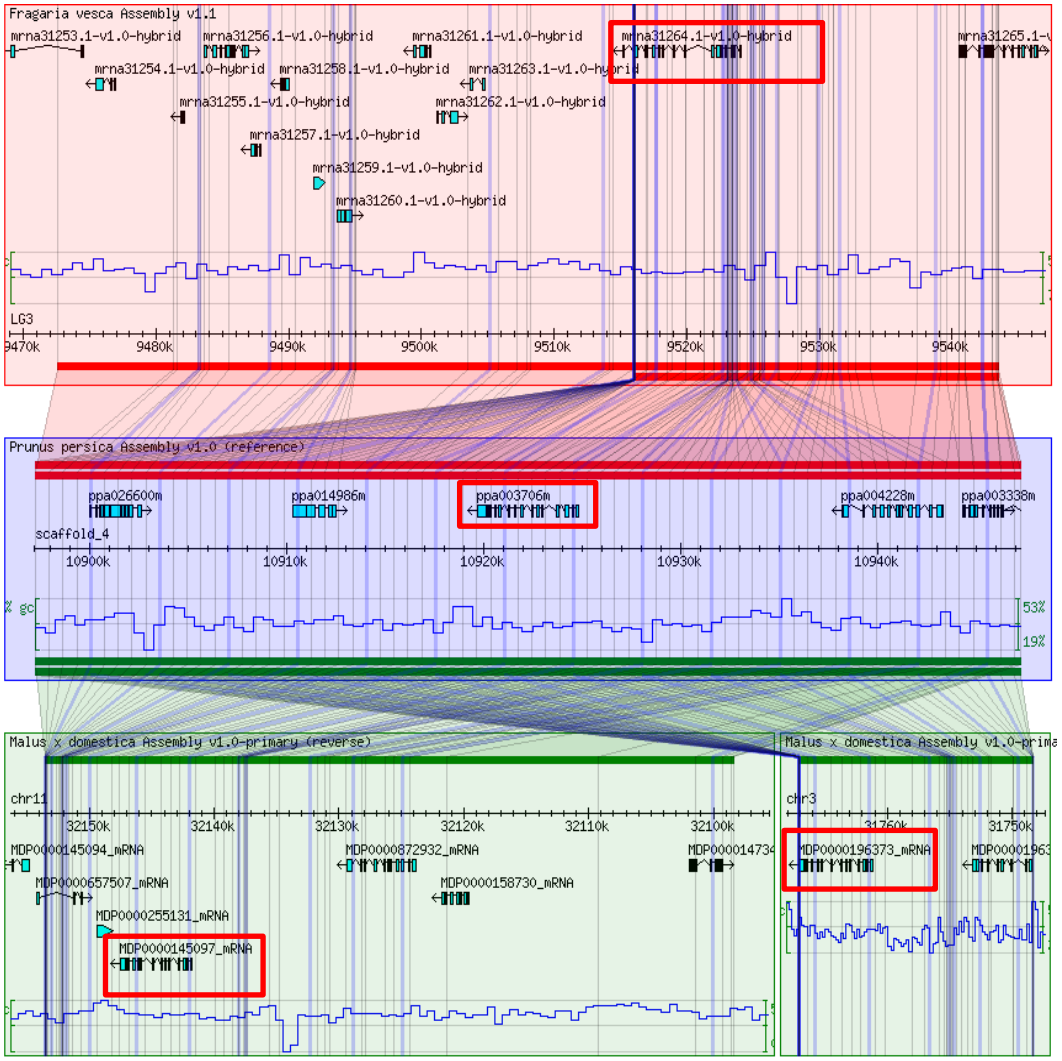
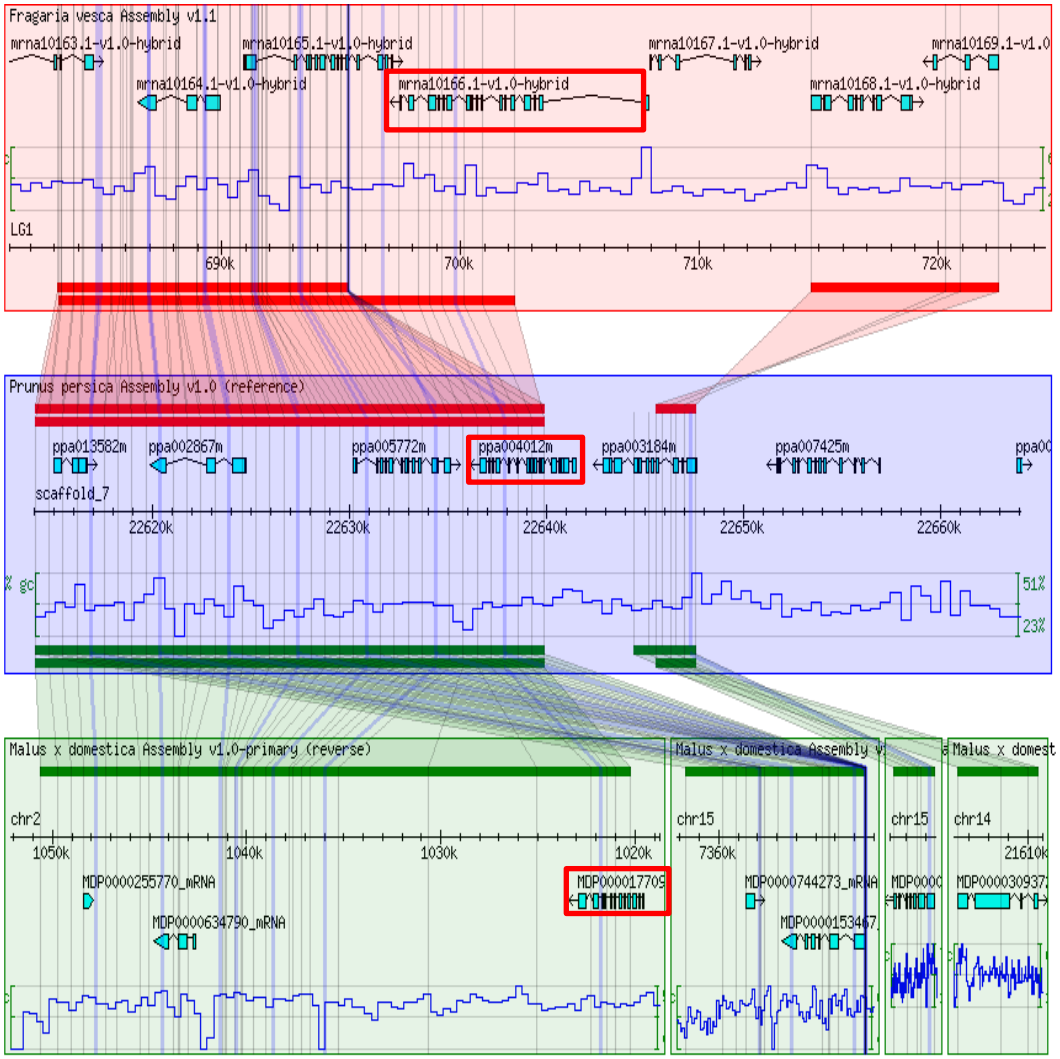
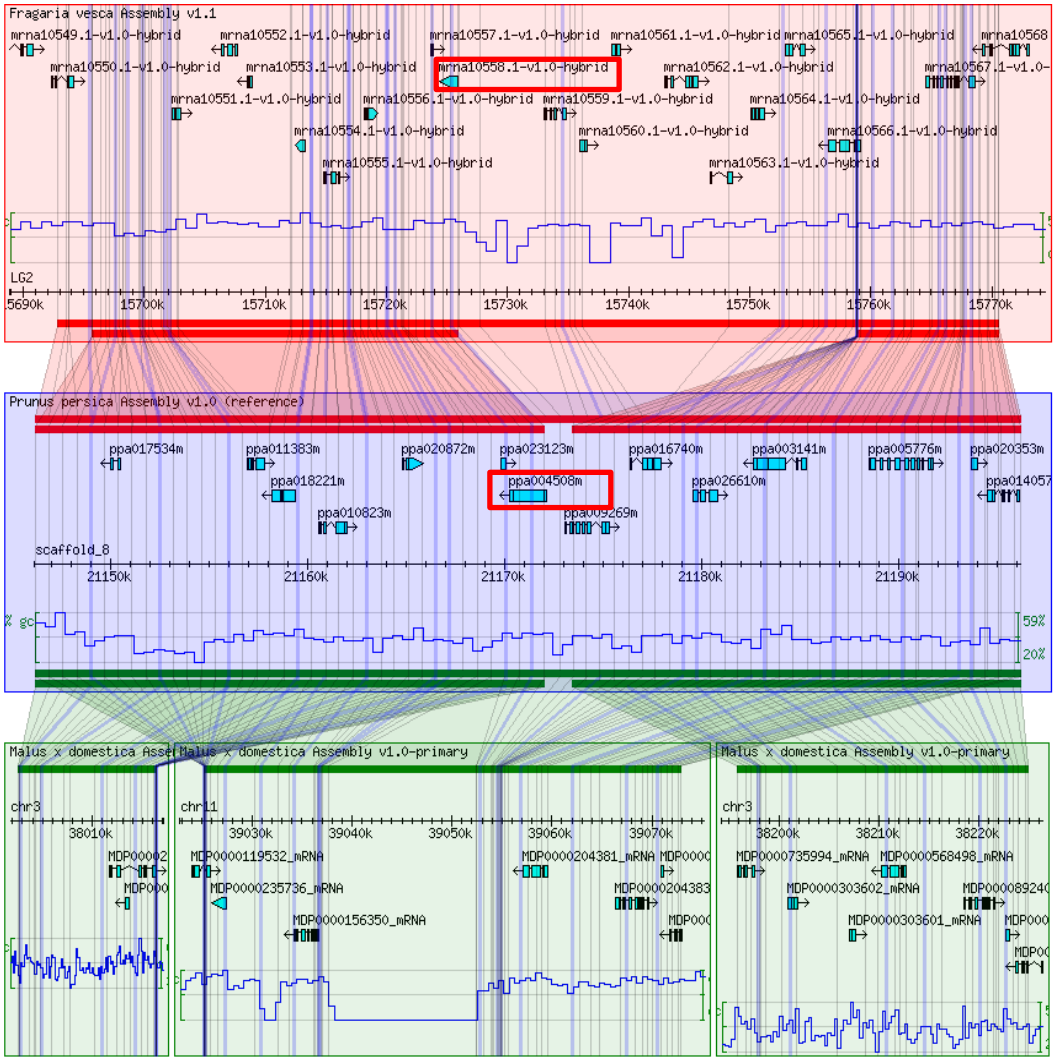

Fragaria vesca Assembly v1.1 (reverse)
mrna13026.1-v1.0-hybrid
mrna13024.1-v1.0-hybrid
mrna13022.1-v1.0-hybrid
mrna13025.1-v1.0-hybrid
mrna13023.1-v1.0-hybrid
mrna13021.1-v1.0-hybrid
mrna13020.1-v1.0-hybrid
LG1
7040k
7030k
7020k
7010k
7000k
Prunus persica Assembly v1.0 (reference)
ppa001013m
ppa012721m
ppa003435m
ppa011201m
ppa017631m
ppa002266m
ppa008223m
ppa006321m
ppa026426m
scaffold_7
18360k
18370k
18380k
18390k
18400k
% gc
64%
21%
Malus x domestica Assembly v1.0-primary
chr15
chr2
chr9
10650k
7520k
7530k
7540k
7550k
7560k
7570k
MDP0000243827_mRNA
MDP0000141595_mRNA
MDP0000137556_mRNA
MDP0000342671_mRNA

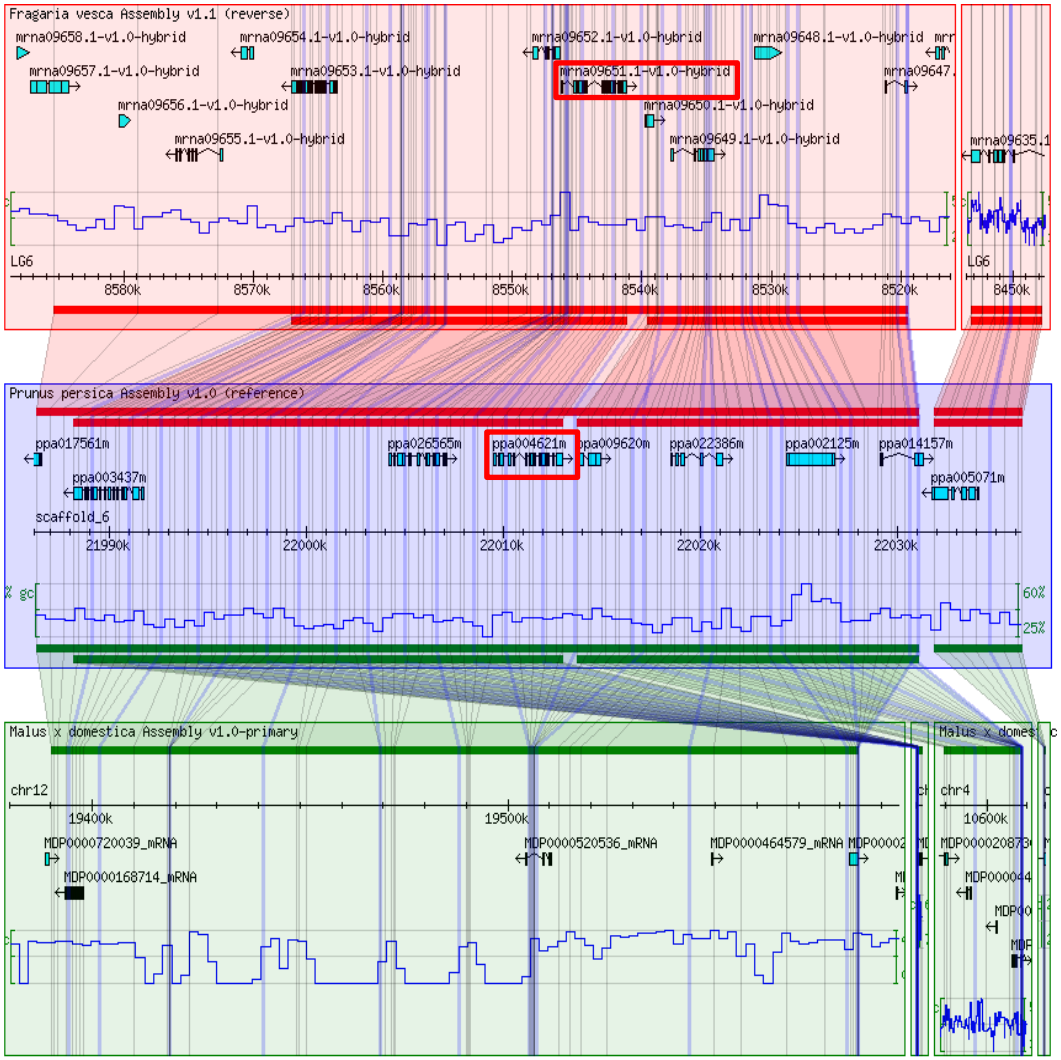

Fragaria vesca Assembly v1.1 (reverse)
mrna09702.1-v1.0-hybrid
mrna09661.1-v1.0-hybrid
mrna09658.1-v1.0-hybrid
mrna09654.1-v1.0-hybrid
mrna09652.1-v1.0-hybr
mrna09701.1-v1.0-hybrid
mrna09660.1-v1.0-hybrid
mrna09656.1-v1.0-hybrid
mrna09651.1-v1.0-h
mrna09659.1-v1.0-hybrid
mrna09655.1-v1.0-hybrid
mrna09657.1-v1.0-hybrid
mrna09653.1-v1.0-hybrid
LG6
8630k
8620k
8610k
8600k
8590k
8580k
8570k
8560k
8550k
8540k
Prunus persica Assembly v1.0 (reference)
ppa015363m
ppa003800m
ppa009116m
ppa003437m
ppa026565m
ppa004621m
ppa010060m
scaffold_6
21970k
21980k
21990k
22000k
22010k
% gc
50%
26%
Malus x domestica Assembly v1.0-primary
Malus x domestica Assembly
chr12
19300k
19400k
19500k
MDP0000219580_mRNA
MDP0000720039_mRNA
MDP0000244758_mRNA
MDP0000168714_mRNA
chr4
10800k

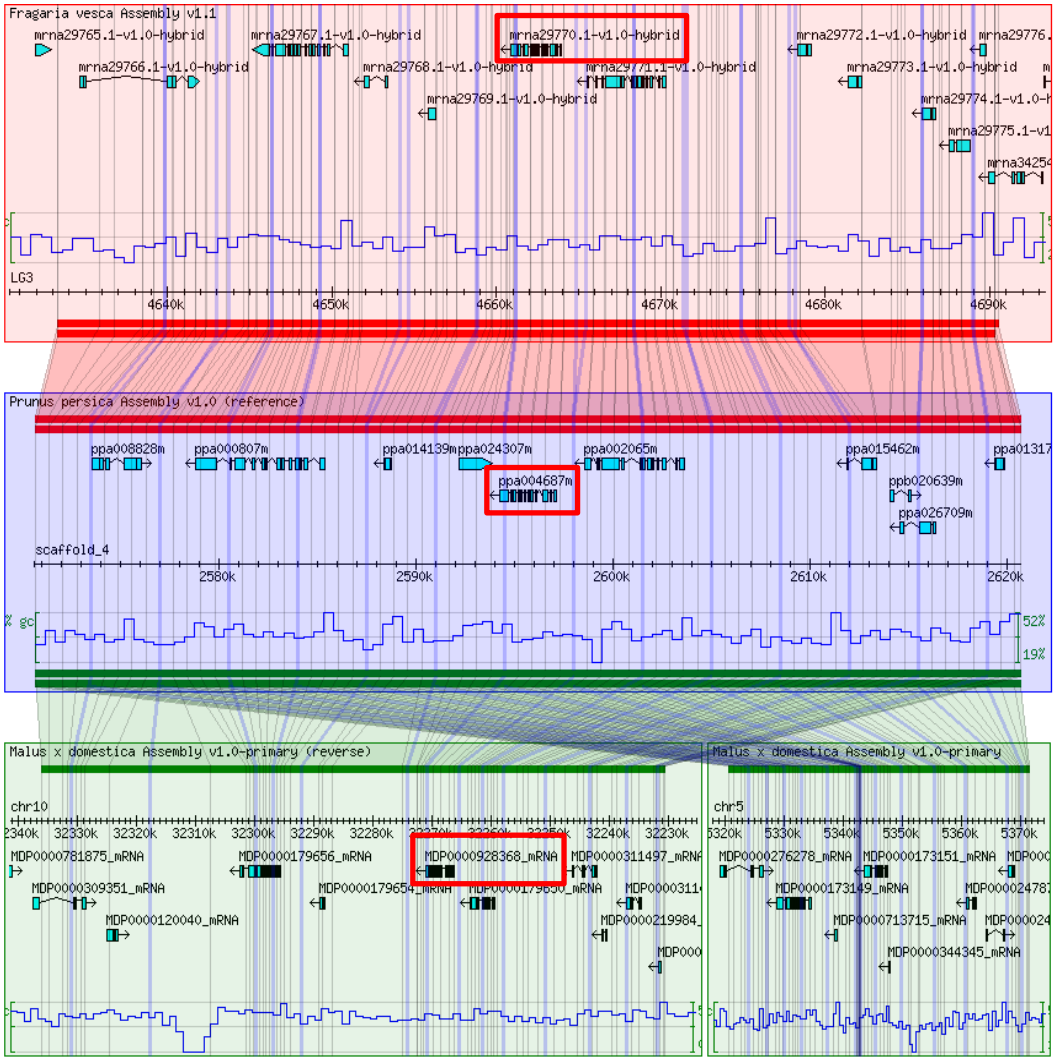

Fragaria vesca Assembly v1.1
mrna23197.1-v1.0-hybrid
mrna23199.1-v1.0-hybrid
mrna23201.1-v1.0-hybrid
mrna23198.1-v1.0-hybrid
mrna23200.1-v1.0-hybrid
LG7
14520k
14530k
14540k
14550k
Prunus persica Assembly v1.0 (reference)
ppa005493m
ppa016176m
ppa003466m
ppa009675m
ppa023509m
ppa011920m
ppa013379m
scaffold_2
21010k
21020k
21030k
21040k
21050k
% gc
53%
27%
Malus x domestica Assembly v1.0-primary (reverse)
chr8
chr1
chr10
18100k
18110k
6020k
6010k
6000k
5990k
5980k
5970k
MDP0000259631_mRNA
MDP0000259632_mRNA
MDP0000259633_mRNA
MDP0000313744_mRNA
MDP0000119447_mRNA
MDP0000264634_mRNA
MDP0000264636_mRNA
MDP0000222938_mRNA
MDP0000264635_mRNA
MDP0000347610_mRNA

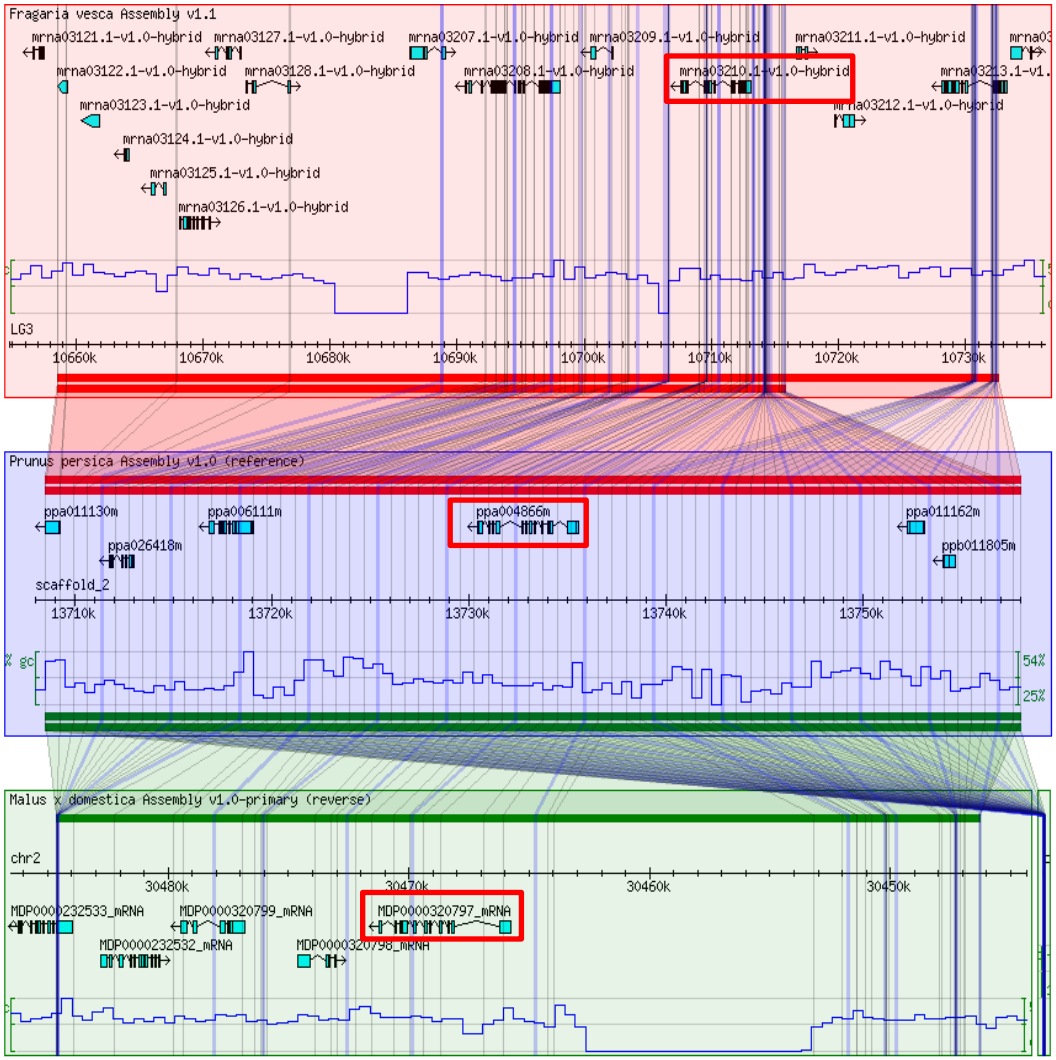

Fragaria vesca Assembly v1.1
mrna31253.1-v1.0-hybrid
mrna31256.1-v1.0-hybrid
mrna31261.1-v1.0-hybrid
mrna31264.1-v1.0-hybrid
mrna31265.1-v
mrna31254.1-v1.0-hybrid
mrna31258.1-v1.0-hybrid
mrna31263.1-v1.0-hybrid
mrna31255.1-v1.0-hybrid
mrna31262.1-v1.0-hybrid
mrna31257.1-v1.0-hybrid
mrna31259.1-v1.0-hybrid
mrna31260.1-v1.0-hybrid
LG3
9470k
9480k
9490k
9500k
9510k
9520k
9530k
9540k
Prunus persica Assembly v1.0 (reference)
ppa026600m
ppa014986m
ppa003706m
ppa004228m
ppa003338m
scaffold_4
10900k
10910k
10920k
10930k
10940k
% gc
53%
19%
Malus x domestica Assembly v1.0-primary (reverse)
chr11
32150k
32140k
32130k
32120k
32110k
32100k
MDP0000145094_mRNA
MDP0000872932_mRNA
MDP0000657507_mRNA
MDP0000158730_mRNA
MDP0000255131_mRNA
MDP0000145097_mRNA
Malus x domestica Assembly v1.0-prime
chr3
31760k
31750k
MDP0000196373_mRNA


Fragaria vesca Assembly v1.1
mrna10163.1-v1.0-hybrid
mrna10164.1-v1.0-hybrid
mrna10165.1-v1.0-hybrid
mrna10166.1-v1.0-hybrid
mrna10167.1-v1.0-hybrid
mrna10168.1-v1.0-hybrid
LG1
690k
700k
710k
720k
Prunus persica Assembly v1.0 (reference)
ppa013582m
ppa002867m
ppa005772m
ppa004012m
ppa003184m
ppa007425m
scaffold_7
22620k
22630k
22640k
22650k
22660k
% gc
51%
23%
Malus x domestica Assembly v1.0-primary (reverse)
chr2
1050k
1040k
1030k
1020k
MDP0000255770_mRNA
MDP0000634790_mRNA
MDP0000017709
Malus x domestica Assembly v
chr15
7360k
MDP0000744273_mRNA
chr15
chr14
21610k
Malus x domest


Fragaria vesca Assembly v1.1
mrna10549.1-v1.0-hybrid
mrna10552.1-v1.0-hybrid
mrna10557.1-v1.0-hybrid
mrna10561.1-v1.0-hybrid
mrna10565.1-v1.0-hybrid
mrna10568
mrna10550.1-v1.0-hybrid
mrna10553.1-v1.0-hybrid
mrna10558.1-v1.0-hybrid
mrna10562.1-v1.0-hybrid
mrna10567.1-v1.0-
mrna10551.1-v1.0-hybrid
mrna10556.1-v1.0-hybrid
mrna10559.1-v1.0-hybrid
mrna10564.1-v1.0-hybrid
mrna10554.1-v1.0-hybrid
mrna10560.1-v1.0-hybrid
mrna10566.1-v1.0-hybrid
mrna10555.1-v1.0-hybrid
mrna10563.1-v1.0-hybrid
LG2
5690k
15700k
15710k
15720k
15730k
15740k
15750k
15760k
15770k
Prunus persica Assembly v1.0 (reference)
ppa017534m
ppa011383m
ppa020872m
ppa023123m
ppa016740m
ppa003141m
ppa005776m
ppa020353m
ppa018221m
ppa004508m
ppa026610m
ppa014057
ppa010823m
ppa009269m
scaffold_8
21150k
21160k
21170k
21180k
21190k
% gc
59%
20%
Malus x domestica Asse
Malus x domestica Assembly v1.0-primary
Malus x domestica Assembly v1.0-primary
chr3
chr11
chr3
38010k
39030k
39040k
39050k
39060k
39070k
38200k
38210k
38220k
MDP0000119532_mRNA
MDP0000204381_mRNA
MDP0000235736_mRNA
MDP0000156350_mRNA
MDP0000735994_mRNA
MDP0000568498_mRNA
MDP0000303602_mRNA
MDP0000303601_mRNA

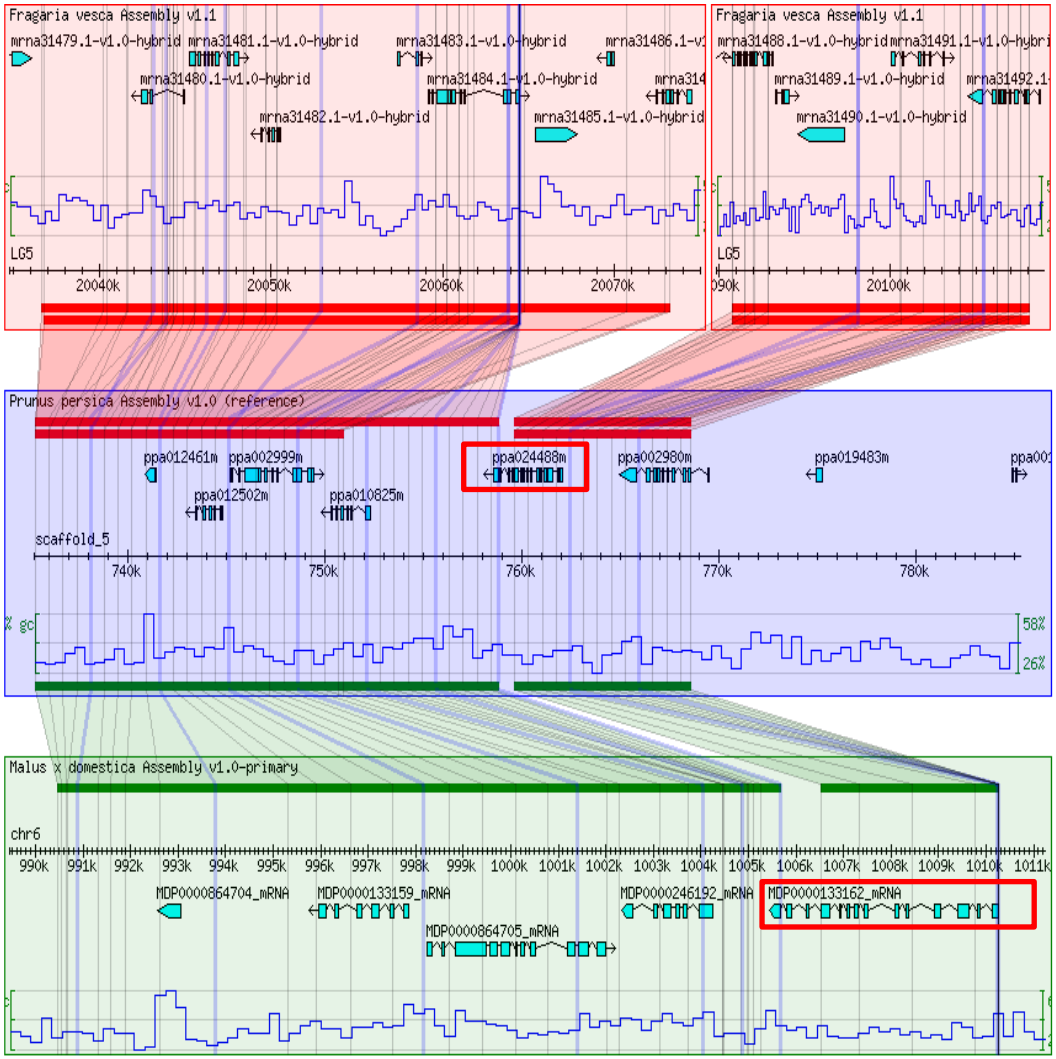

Fragaria vesca Assembly v1.1 (reverse)
mrna09658.1-v1.0-hybrid
mrna09654.1-v1.0-hybrid
mrna09652.1-v1.0-hybrid
mrna09648.1-v1.0-hybrid
mrna09657.1-v1.0-hybrid
mrna09653.1-v1.0-hybrid
mrna09651.1-v1.0-hybrid
mrna09656.1-v1.0-hybrid
mrna09650.1-v1.0-hybrid
mrna09655.1-v1.0-hybrid
mrna09649.1-v1.0-hybrid
mrna09635.1
LG6
8580k
8570k
8560k
8550k
8540k
8530k
8520k
8450k
Prunus persica Assembly v1.0 (reference)
ppa017561m
ppa026565m
ppa004621m
ppa009620m
ppa022386m
ppa002125m
ppa014157m
ppa003437m
ppa005071m
scaffold_6
21990k
22000k
22010k
22020k
22030k
% gc
60%
25%
Malus x domestica Assembly v1.0-primary
chr12
19400k
19500k
MDP0000720039_mRNA
MDP0000520536_mRNA
MDP0000464579_mRNA
MDP0000168714_mRNA
chr4
10600k

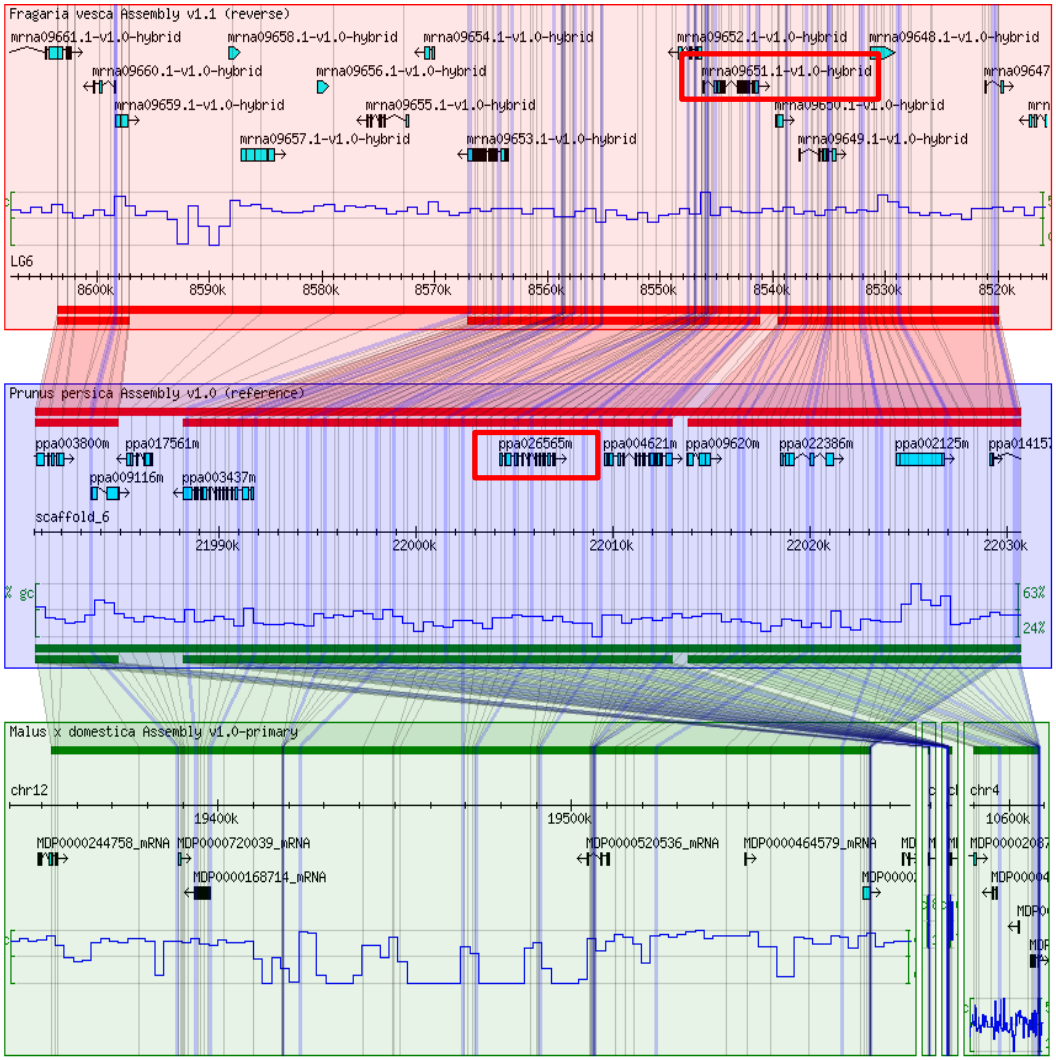

Fragaria vesca Assembly v1.1
mrna29765.1-v1.0-hybrid
mrna29767.1-v1.0-hybrid
mrna29770.1-v1.0-hybrid
mrna29772.1-v1.0-hybrid
mrna29766.1-v1.0-hybrid
mrna29768.1-v1.0-hybrid
mrna29771.1-v1.0-hybrid
mrna29773.1-v1.0-hybrid
mrna29769.1-v1.0-hybrid
mrna29774.1-v1.0-h
mrna29775.1-v1
LG3
4640k
4650k
4660k
4670k
4680k
4690k
Prunus persica Assembly v1.0 (reference)
ppa008828m
ppa000807m
ppa014139m
ppa024307m
ppa002065m
ppa015462m
ppa004687m
ppb020639m
ppa026709m
scaffold_4
2580k
2590k
2600k
2610k
2620k
% gc
52%
19%
Malus x domestica Assembly v1.0-primary (reverse)
chr10
32340k
32330k
32320k
32310k
32300k
32290k
32280k
32240k
32230k
MDP0000781875_mRNA
MDP0000179656_mRNA
MDP0000928368_mRNA
MDP0000311497_mRNA
MDP0000309351_mRNA
MDP0000179654_mRNA
MDP0000120040_mRNA
MDP0000219984_
Malus x domestica Assembly v1.0-primary
chr5
5320k
5330k
5340k
5350k
5360k
5370k
MDP0000276278_mRNA
MDP0000173151_mRNA
MDP0000173149_mRNA
MDP0000713715_mRNA
MDP0000344345_mRNA


Fragaria vesca Assembly v1.1
mrna03121.1-v1.0-hybrid
mrna03127.1-v1.0-hybrid
mrna03207.1-v1.0-hybrid
mrna03209.1-v1.0-hybrid
mrna03211.1-v1.0-hybrid
mrna03122.1-v1.0-hybrid
mrna03128.1-v1.0-hybrid
mrna03208.1-v1.0-hybrid
mrna03210.1-v1.0-hybrid
mrna03123.1-v1.0-hybrid
mrna03212.1-v1.0-hybrid
mrna03124.1-v1.0-hybrid
mrna03125.1-v1.0-hybrid
mrna03126.1-v1.0-hybrid
LG3
10660k
10670k
10680k
10690k
10700k
10710k
10720k
10730k
Prunus persica Assembly v1.0 (reference)
ppa011130m
ppa006111m
ppa004866m
ppa011162m
ppa026418m
ppb011805m
scaffold_2
13710k
13720k
13730k
13740k
13750k
% gc
54%
25%
Malus x domestica Assembly v1.0-primary (reverse)
chr2
30480k
30470k
30460k
30450k
MDP0000232533_mRNA
MDP0000320799_mRNA
MDP0000320797_mRNA
MDP0000232532_mRNA
MDP0000320798_mRNA

Fragaria vesca Assembly v1.1 (reverse)
mrna28551.1-v1.0-hybrid
mrna28549.1-v1.0-hybrid
mrna28544.1-v1.0-hybrid
mrna34542.1-v1.0-hybrid
mrna28541.1-v1
mrna28550.1-v1.0-hybrid
mrna28548.1-v1.0-hybrid
mrna28543.1-v1.0-hybrid
mrna34540.1-v1.0-hybrid
mrna28547.1-v1.0-hybrid
mrna28542.1-v1.0-hybrid
mrna28546.1-v1.0-hybrid
mrna34543.1-v1.0-hybrid
mrna28545.1-v1.0-hybrid
mrna34541.1-v1.0-hybrid
LG3
17210k
17200k
17190k
17180k
17170k
17160k
17150k
17140k
17130k
17120k
17110k
Prunus persica Assembly v1.0 (reference)
ppa006518m
ppa022847m
ppa003207m
ppa001557m
scaffold_6
6780k
6790k
6800k
6810k
6820k
% gc
49%
13%
Malus x domestica Assembly v1.0-primary (reverse)
chr15
400k
28390k
28380k
28370k
28360k
28350k
28340k
28330k
28320k
28310k
28300k
28290k
28280k
28270k
28260k
MDP0000139654_mRNA
MDP0000133760_mRNA
MDP0000225981_mRNA
MDP0000225980_mRNA
Malus x domestica Ass
chr11
12040k
12050k
MDP0000903417_mRNA

Fragaria vesca Assembly v1.1 (reverse)
mrna14601.1-v1.0-hybrid
mrna14597.1-v1.0-hybrid
mrna14592.1-v1.0-hybrid
mrna14588.1-v1.0-hybrid
mrna14600.1-v1.0-hybrid
mrna14595.1-v1.0-hybrid
mrna14591.1-v1.0-hybrid
mrna14587.1-v1.0-hybrid
mrna14599.1-v1.0-hybrid
mrna14594.1-v1.0-hybrid
mrna14590.1-v1.0-hybrid
mrna14586.1-v1.0-hybr
mrna14598.1-v1.0-hybrid
mrna14593.1-v1.0-hybrid
mrna14589.1-v1.0-hybrid
mrna14585.1-v1.0-h
mrna14596.1-v1.0-hybrid
LG1
7870k
7860k
7850k
7840k
7830k
7820k
7810k
7800k
7790k
7780k
7770k
Prunus persica Assembly v1.0 (reference)
ppa009119m
ppa022013m
ppa006927m
ppa011976m
ppa005827m
ppa021513m
ppa009202m
ppa002122m
ppa015768m
ppa024476m
ppa018677m
scaffold_7
17610k
17620k
17630k
17640k
17650k
% gc
57%
27%
Malus x domestica Assembly v1.0-primary (reverse)
chr2
11700k
11690k
11680k
11670k
11660k
11650k
11640k
MDP0000861535_mRNA
MDP0000273966_mRNA
MDP0000267909_mRNA
MDP0000273963_mRNA
MDP0000273964_mRNA


Fragaria vesca Assembly v1.1
mrna31479.1-v1.0-hybrid
mrna31481.1-v1.0-hybrid
mrna31483.1-v1.0-hybrid
mrna31486.1-v1
mrna31480.1-v1.0-hybrid
mrna31484.1-v1.0-hybrid
mrna314
mrna31482.1-v1.0-hybrid
mrna31485.1-v1.0-hybrid
LG5
20040k
20050k
20060k
20070k
Fragaria vesca Assembly v1.1
mrna31488.1-v1.0-hybrid
mrna31491.1-v1.0-hybri
mrna31489.1-v1.0-hybrid
mrna31492.1
mrna31490.1-v1.0-hybrid
LG5
090k
20100k
Prunus persica Assembly v1.0 (reference)
ppa012461m
ppa002999m
ppa024488m
ppa002980m
ppa019483m
ppa001
ppa012502m
ppa010825m
scaffold_5
740k
750k
760k
770k
780k
% gc
58%
26%
Malus x domestica Assembly v1.0-primary
chr6
990k
991k
992k
993k
994k
995k
996k
997k
998k
999k
1000k
1001k
1002k
1003k
1004k
1005k
1006k
1007k
1008k
1009k
1010k
1011k
MDP0000864704_mRNA
MDP0000133159_mRNA
MDP0000246192_mRNA
MDP0000133162_mRNA
MDP0000864705_mRNA


Fragaria vesca Assembly v1.1 (reverse)
mrna09661.1-v1.0-hybrid
mrna09658.1-v1.0-hybrid
mrna09654.1-v1.0-hybrid
mrna09652.1-v1.0-hybrid
mrna09648.1-v1.0-hybrid
mrna09660.1-v1.0-hybrid
mrna09656.1-v1.0-hybrid
mrna09651.1-v1.0-hybrid
mrna09659.1-v1.0-hybrid
mrna09655.1-v1.0-hybrid
mrna09650.1-v1.0-hybrid
mrna09657.1-v1.0-hybrid
mrna09653.1-v1.0-hybrid
mrna09649.1-v1.0-hybrid
LG6
8600k
8590k
8580k
8570k
8560k
8550k
8540k
8530k
8520k
Prunus persica Assembly v1.0 (reference)
ppa003800m
ppa017561m
ppa026565m
ppa004621m
ppa009620m
ppa022386m
ppa002125m
ppa009116m
ppa003437m
scaffold_6
21990k
22000k
22010k
22020k
22030k
% gc
63%
24%
Malus x domestica Assembly v1.0-primary
chr12
19400k
19500k
chr4
10500k
MDP0000244758_mRNA
MDP0000720039_mRNA
MDP0000520536_mRNA
MDP0000464579_mRNA
MDP0000168714_mRNA