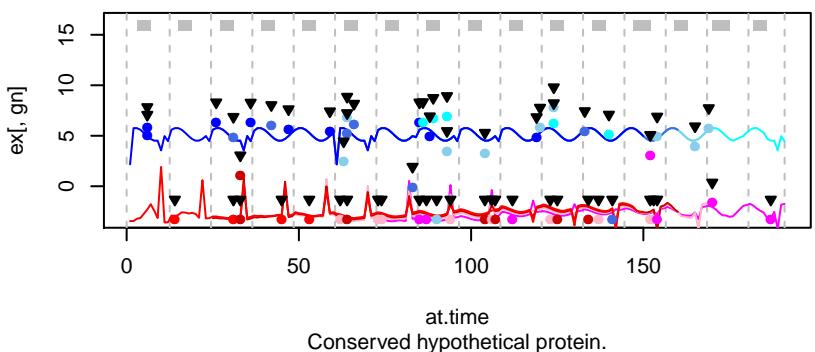
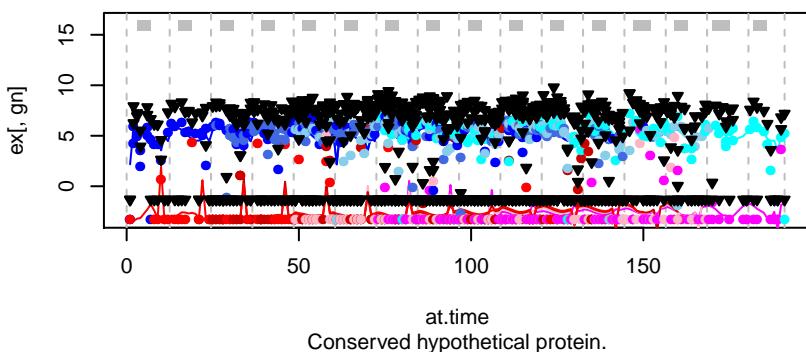


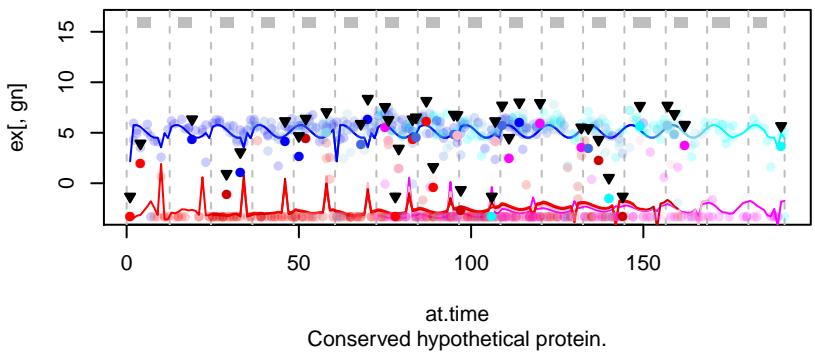
Os10g0489650 : AT2G32120.1 : HSP70T-2
kos_tak



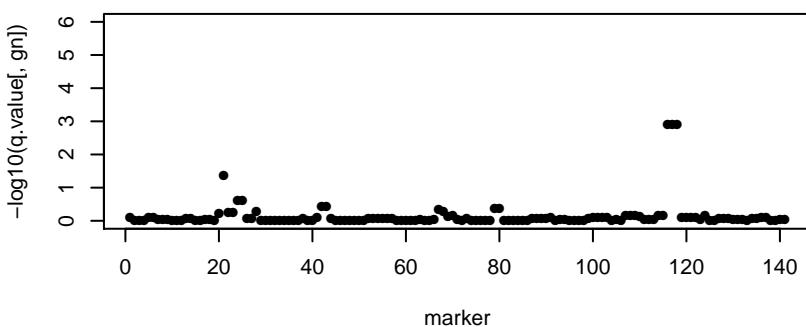
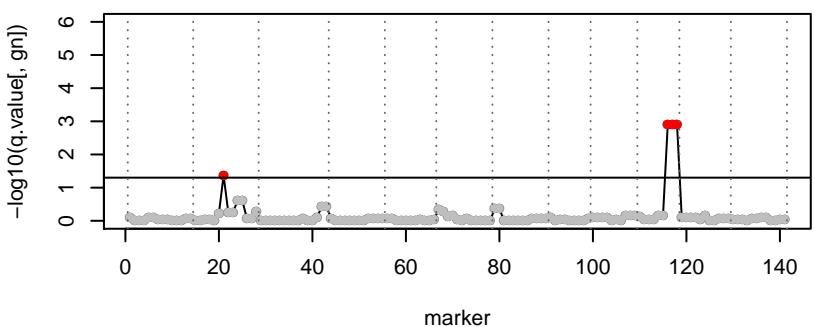
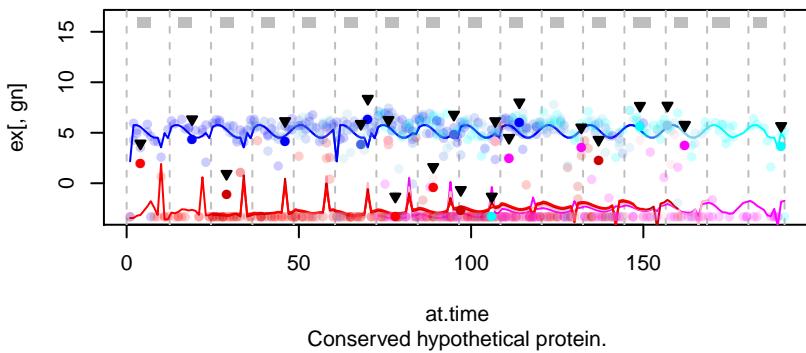
Os10g0489650 : AT2G32120.1 : HSP70T-2
Background



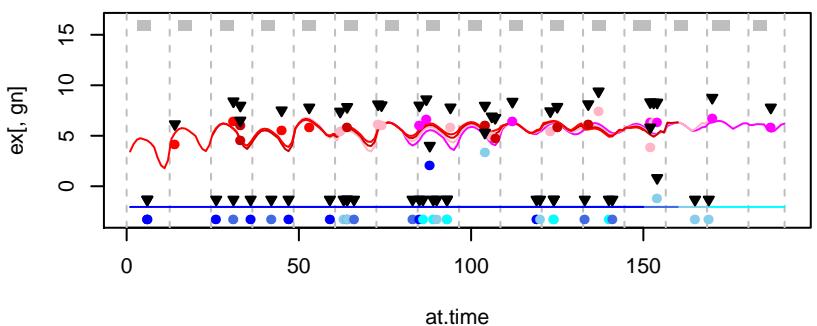
Os10g0489650 : AT2G32120.1 : HSP70T-2
919.651550678297_Most significant each peak



Os10g0489650 : AT2G32120.1 : HSP70T-2
913.539949967805_Most significant peak

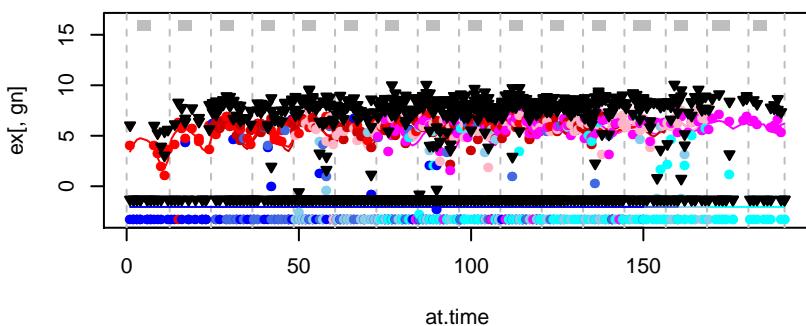


Os03g0247300 : AT1G58440.1 : SQE1,XF1
kos_tak



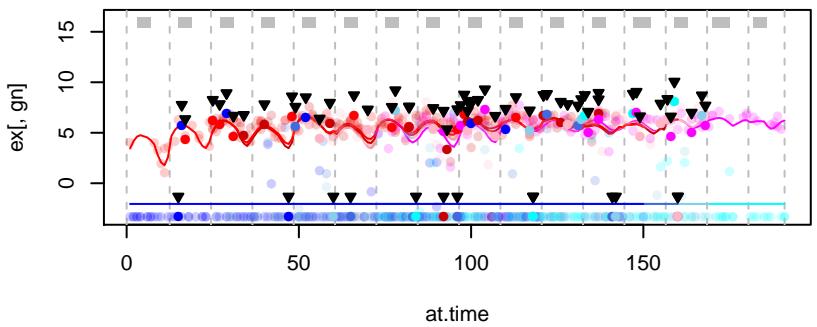
Protein of unknown function DUF1618 domain containing protein.

Os03g0247300 : AT1G58440.1 : SQE1,XF1
Background



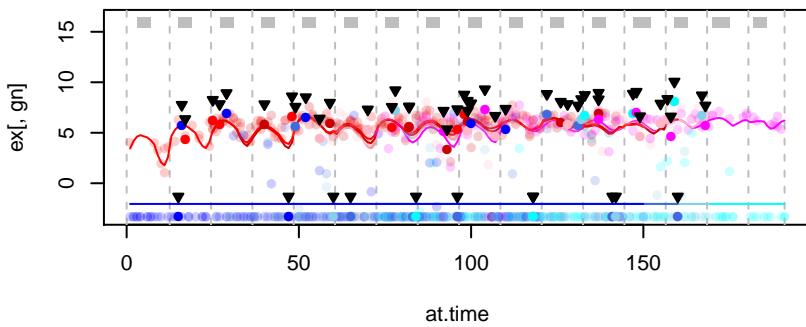
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1152.08402859275_Most significant each peak

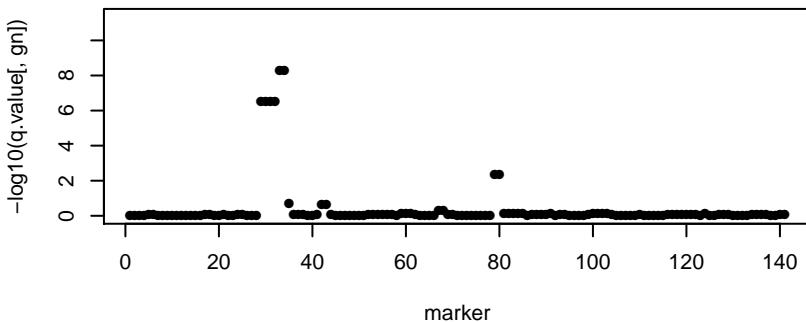
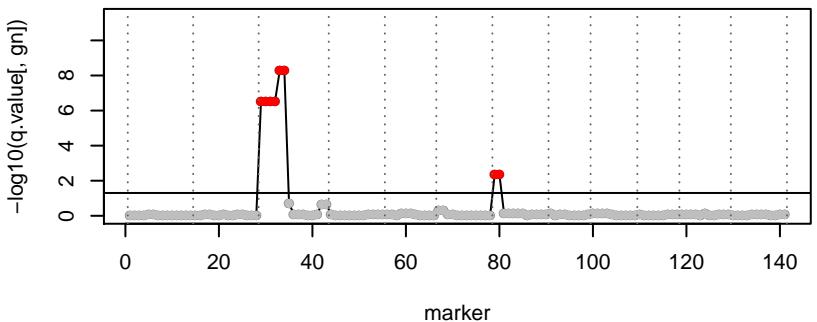


Protein of unknown function DUF1618 domain containing protein.

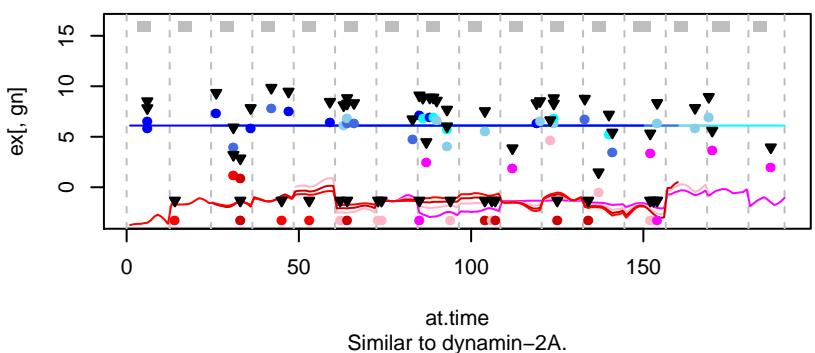
Os03g0247300 : AT1G58440.1 : SQE1,XF1
1200.77223692043_Most significant peak



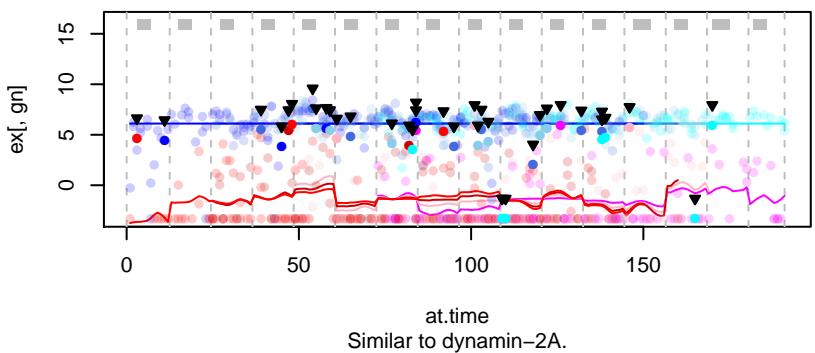
Protein of unknown function DUF1618 domain containing protein.



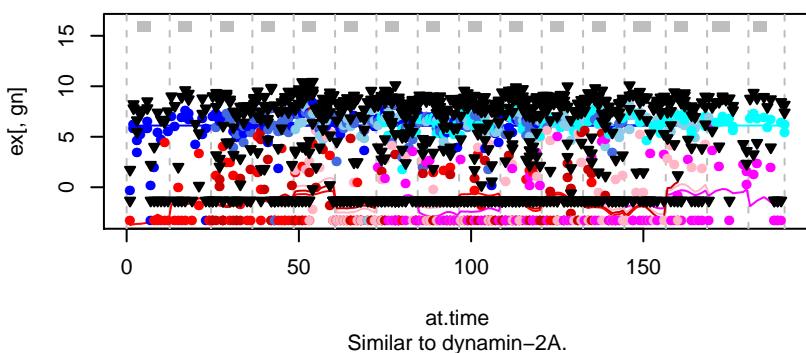
Os03g0732200 : AT1G10290.1 : ADL6,DRP2A
kos_tak



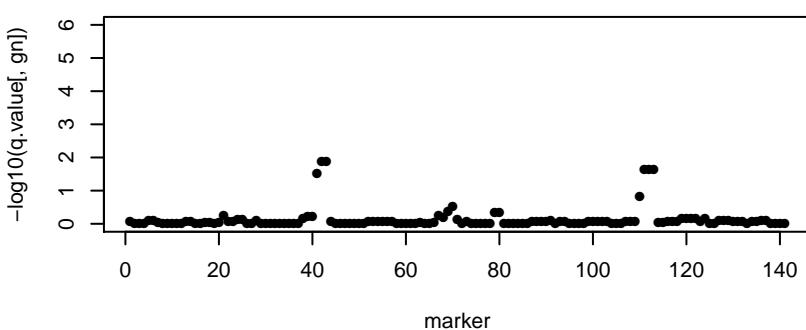
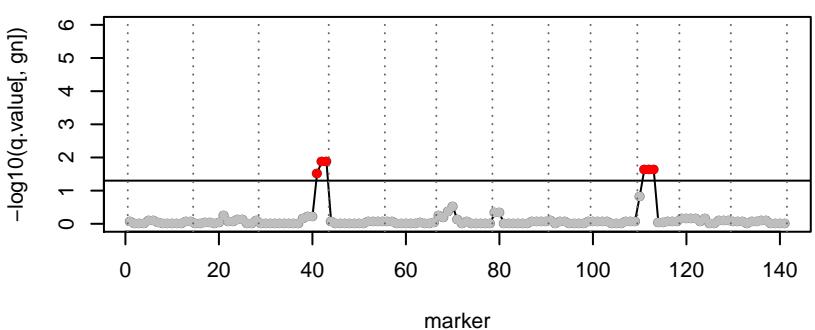
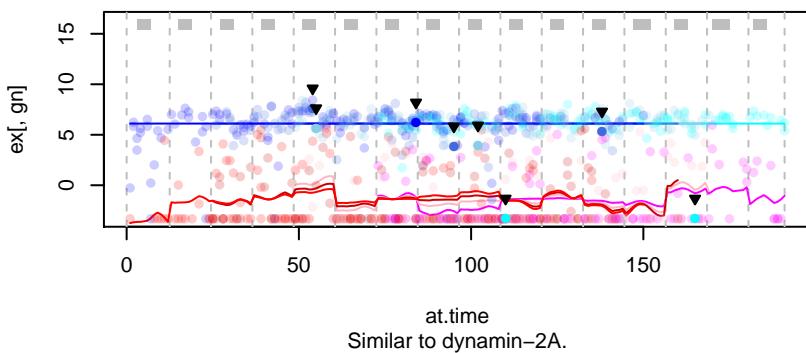
Os03g0732200 : AT1G10290.1 : ADL6,DRP2A
1384.66429896254_Most significant each peak



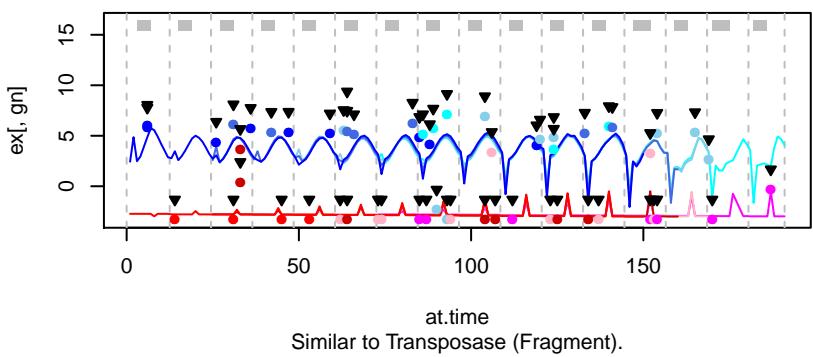
Os03g0732200 : AT1G10290.1 : ADL6,DRP2A
Background



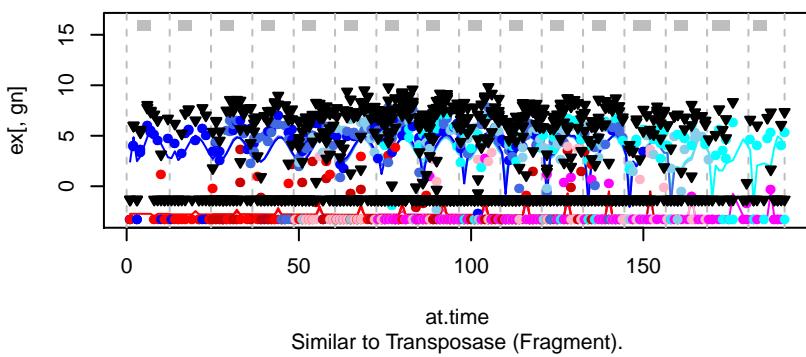
Os03g0732200 : AT1G10290.1 : ADL6,DRP2A
1394.91521900489_Most significant peak



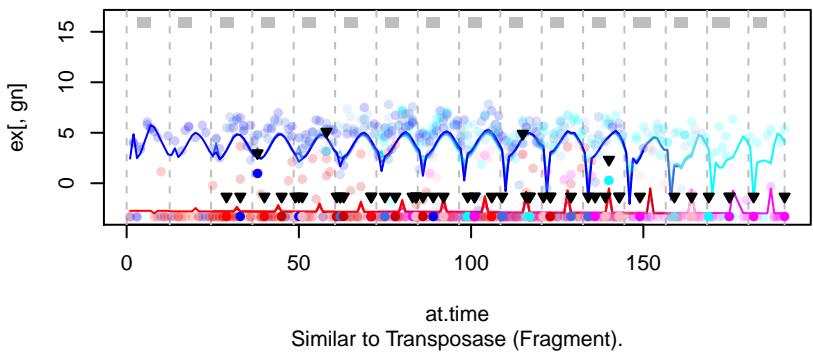
Os06g0226000 : AT5G41980.1 :
kos_tak



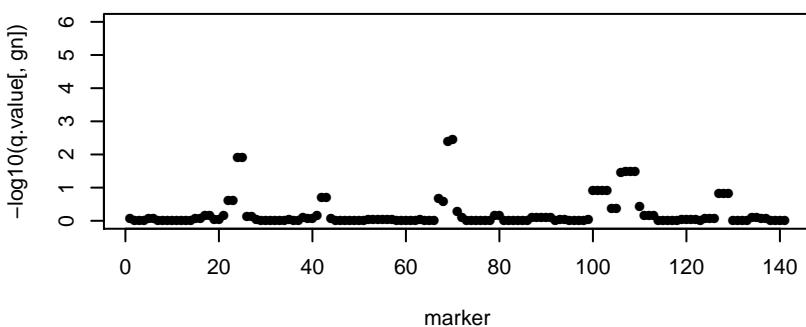
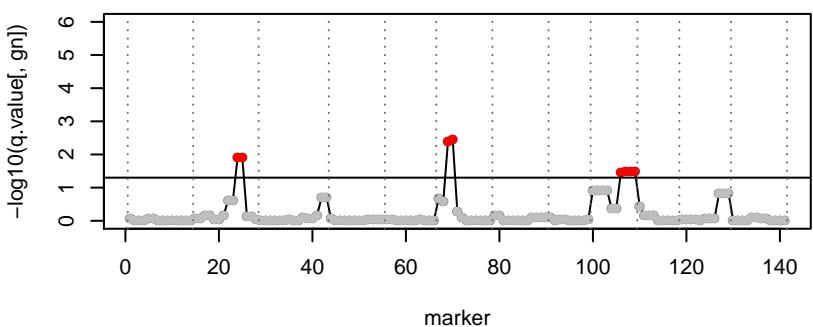
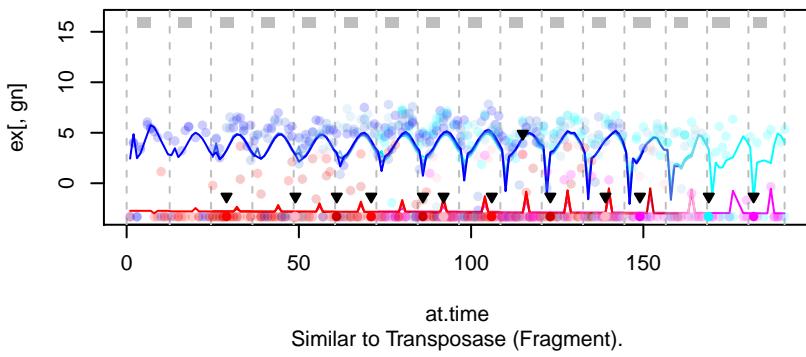
Os06g0226000 : AT5G41980.1 :
Background



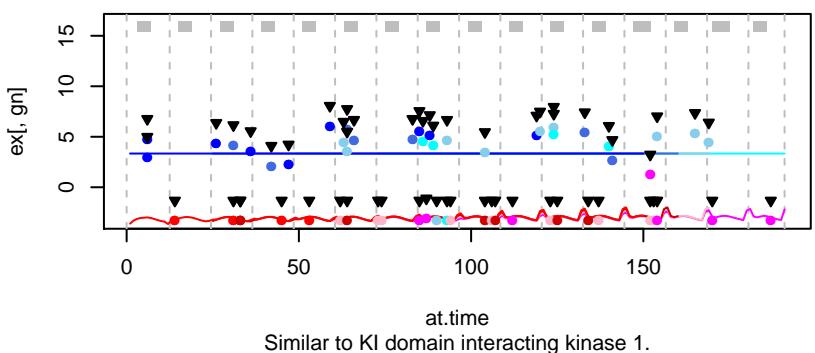
Os06g0226000 : AT5G41980.1 :
1339.11356097655_Most significant each peak



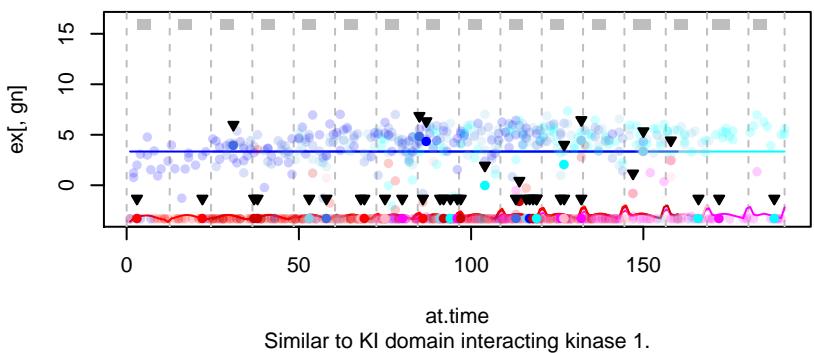
Os06g0226000 : AT5G41980.1 :
1375.64283220927_Most significant peak



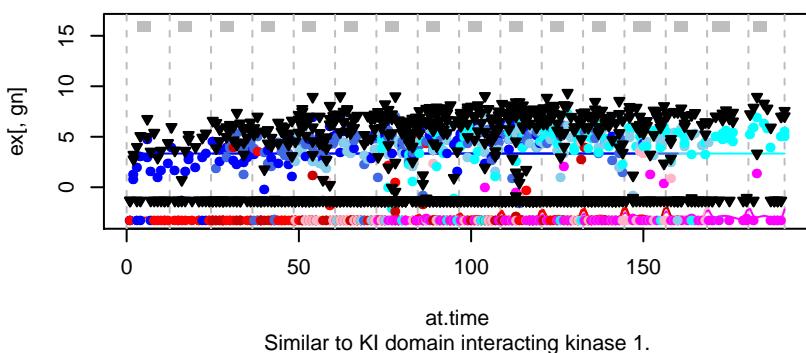
Os07g0541500 : AT4G23190.1 : AT-RLK3,CRK11
kos_tak



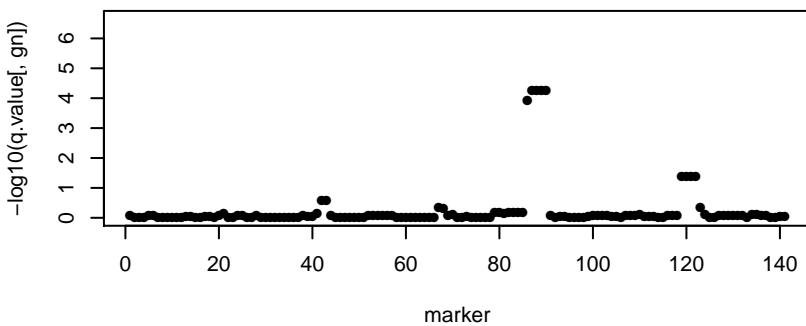
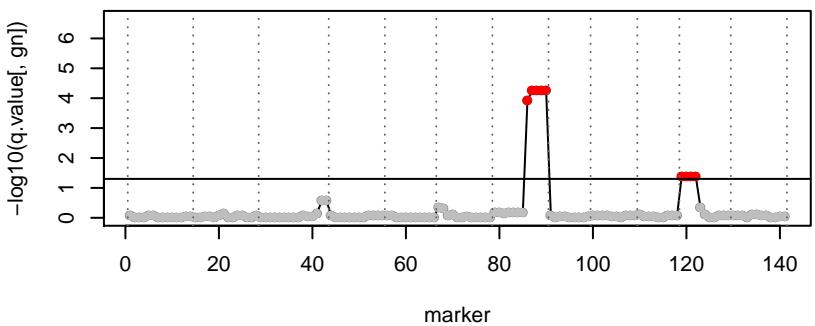
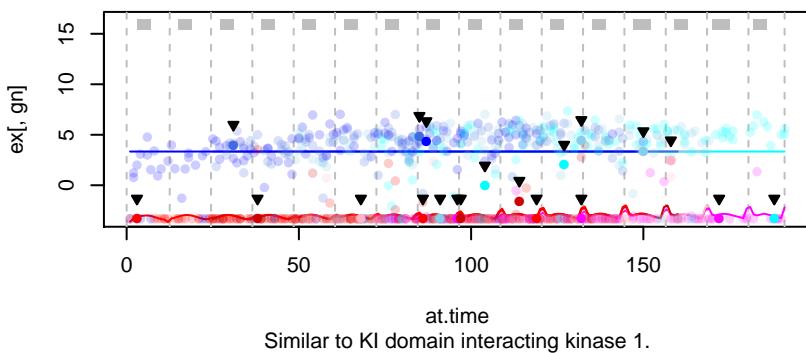
Os07g0541500 : AT4G23190.1 : AT-RLK3,CRK11
1062.68300442822_Most significant each peak



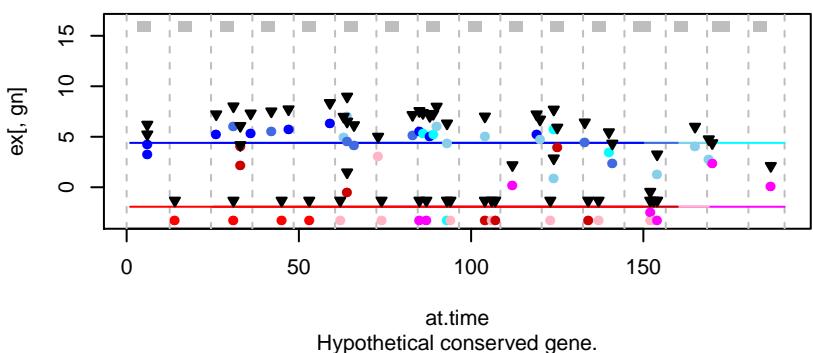
Os07g0541500 : AT4G23190.1 : AT-RLK3,CRK11
Background



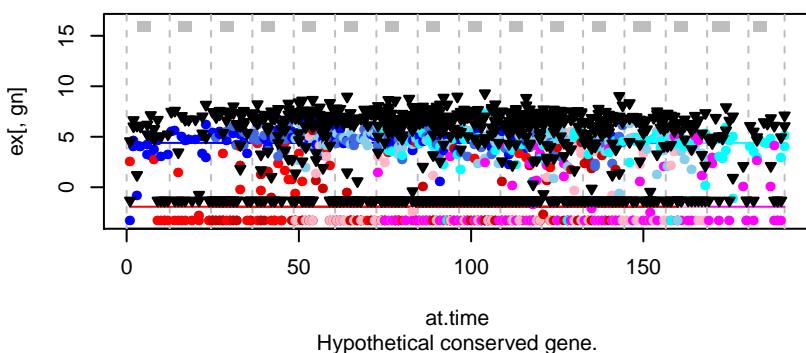
Os07g0541500 : AT4G23190.1 : AT-RLK3,CRK11
1070.70225912062_Most significant peak



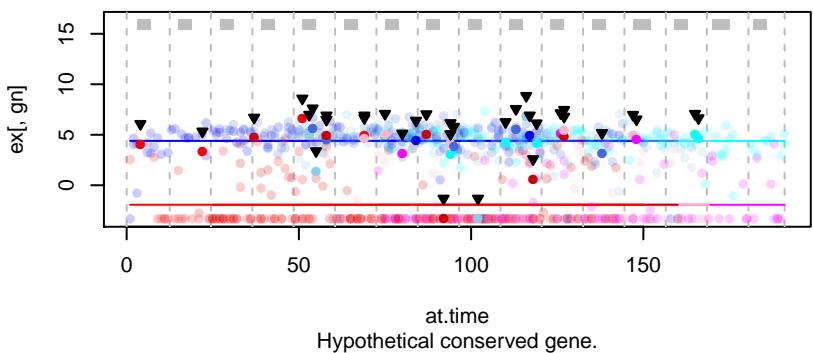
Os11g0146950 : AT4G15890.1 :
kos_tak



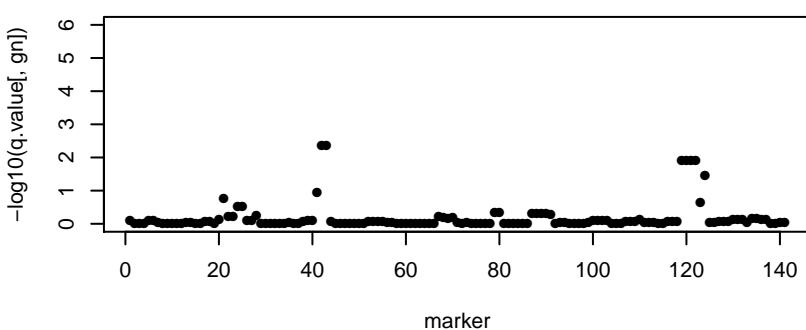
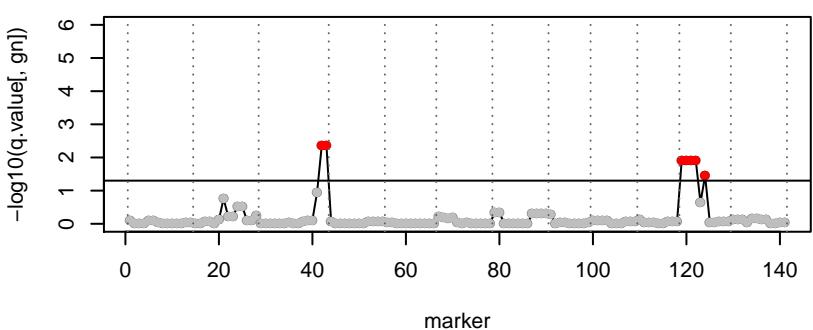
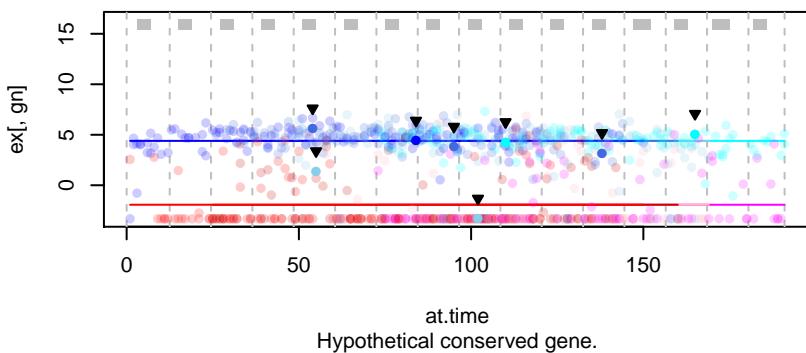
Os11g0146950 : AT4G15890.1 :
Background



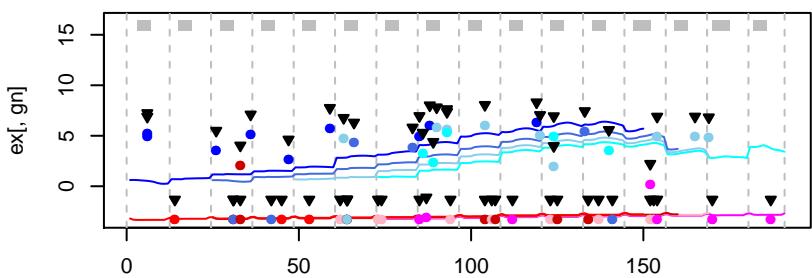
Os11g0146950 : AT4G15890.1 :
1326.40416238342_Most significant each peak



Os11g0146950 : AT4G15890.1 :
1286.05164710181_Most significant peak

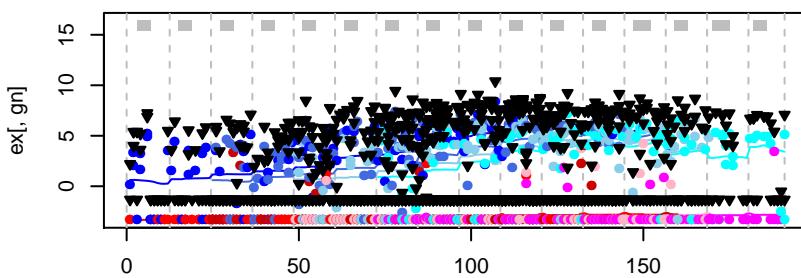


Os07g0543500 : AT5G18980.1 :
kos_tak



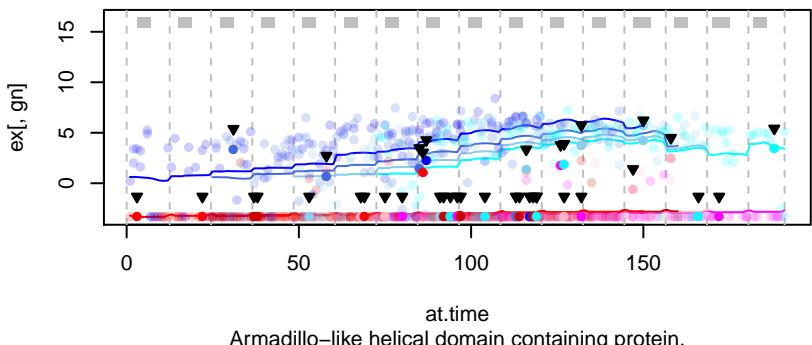
at.time
Armadillo-like helical domain containing protein.

Os07g0543500 : AT5G18980.1 :
Background



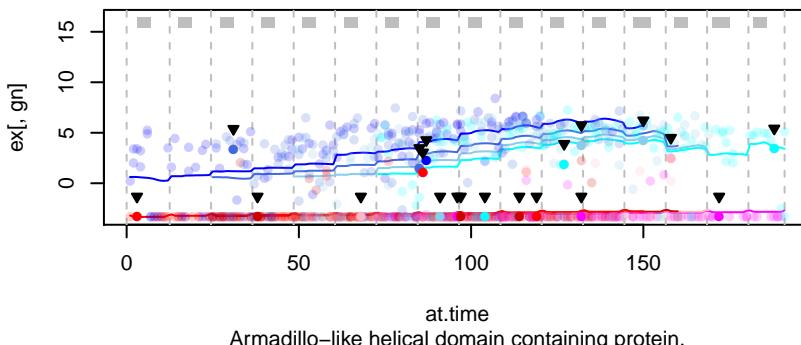
at.time
Armadillo-like helical domain containing protein.

Os07g0543500 : AT5G18980.1 :
1198.23108803674_Most significant each peak

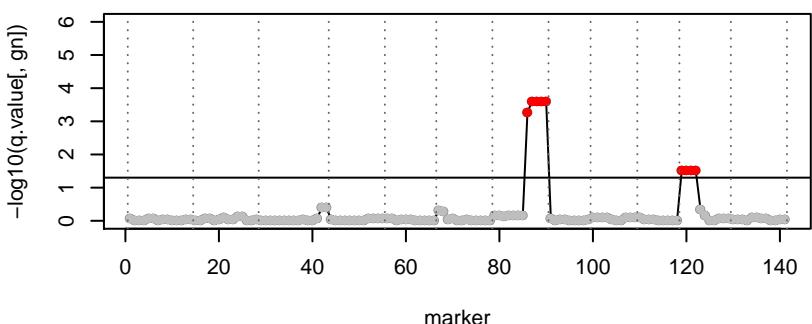


at.time
Armadillo-like helical domain containing protein.

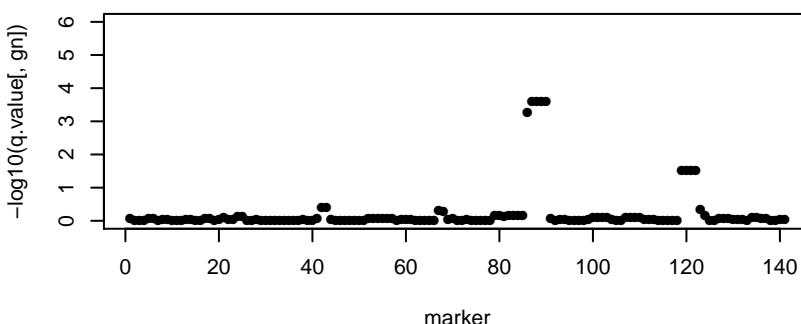
Os07g0543500 : AT5G18980.1 :
1217.71252942667_Most significant peak



at.time
Armadillo-like helical domain containing protein.

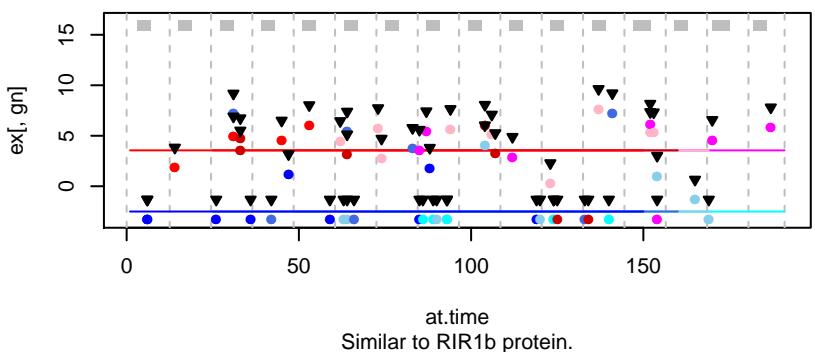


marker

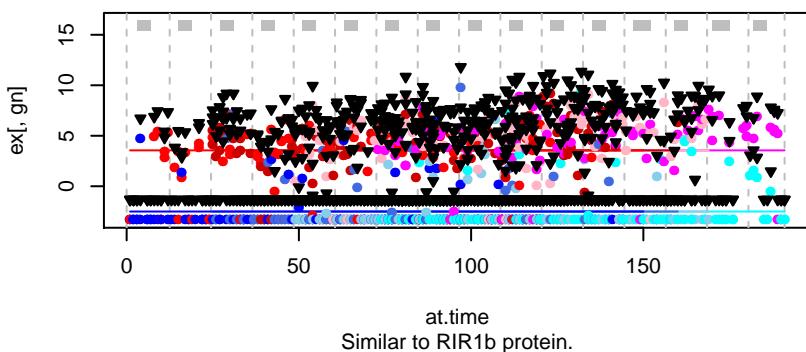


marker

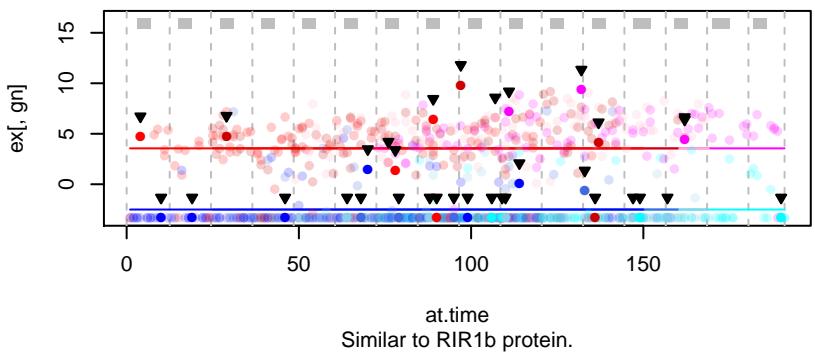
Os10g0569800 : AT5G15940.1 :
kos_tak



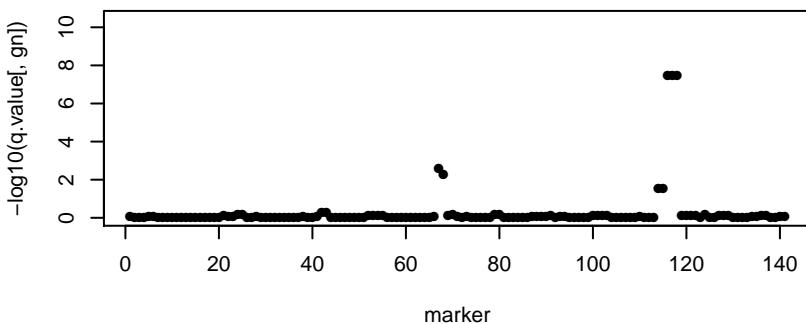
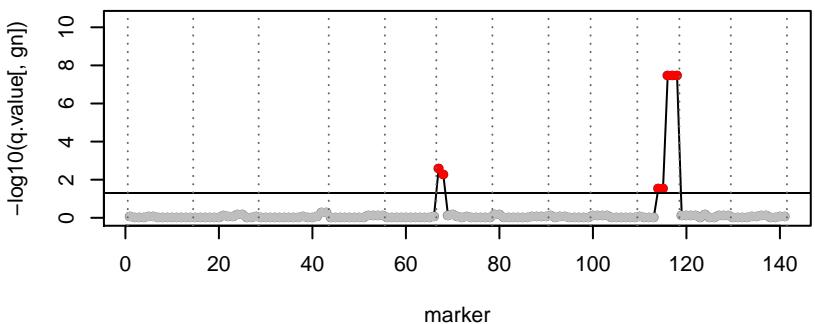
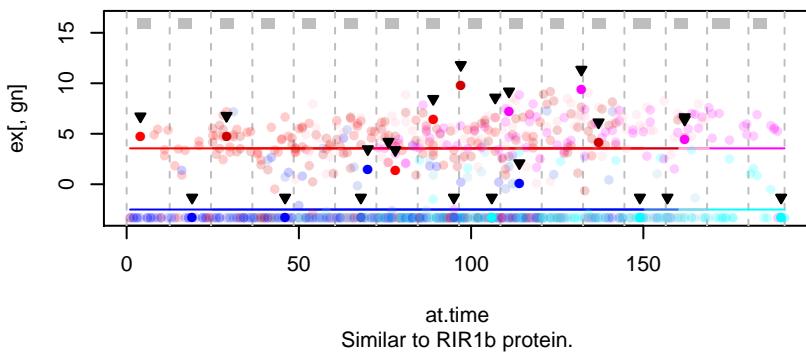
Os10g0569800 : AT5G15940.1 :
Background



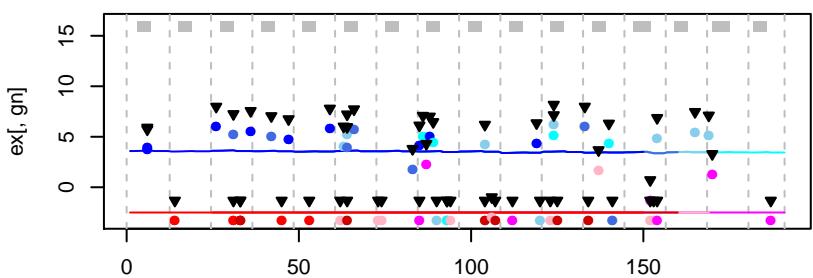
Os10g0569800 : AT5G15940.1 :
1519.3881426869_Most significant each peak



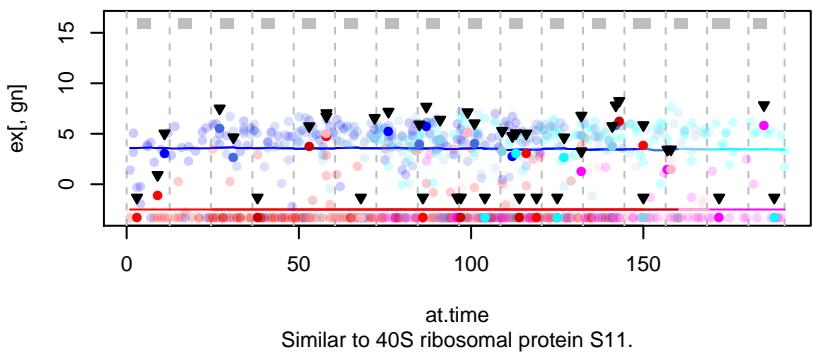
Os10g0569800 : AT5G15940.1 :
1558.07962016979_Most significant peak



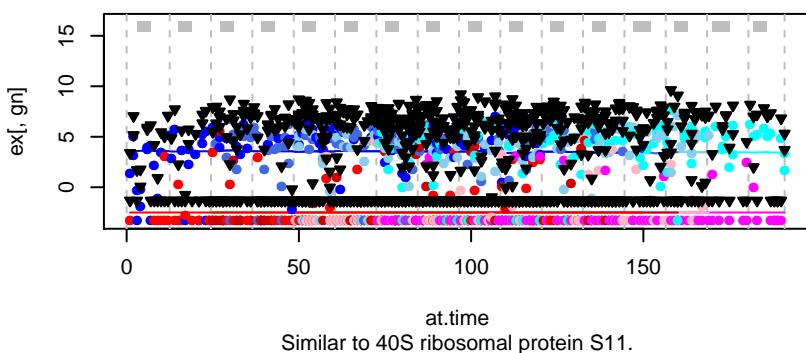
Os07g0565000 : :
kos_tak



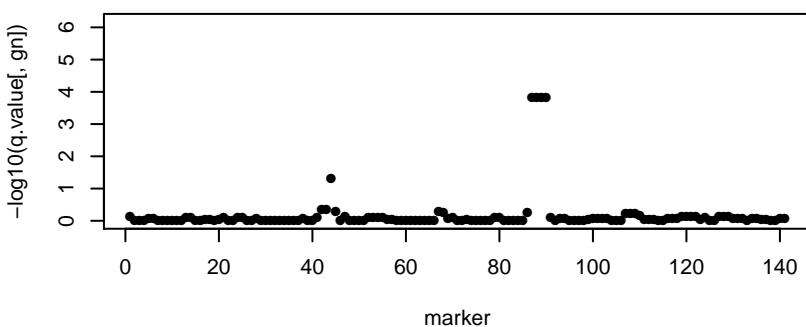
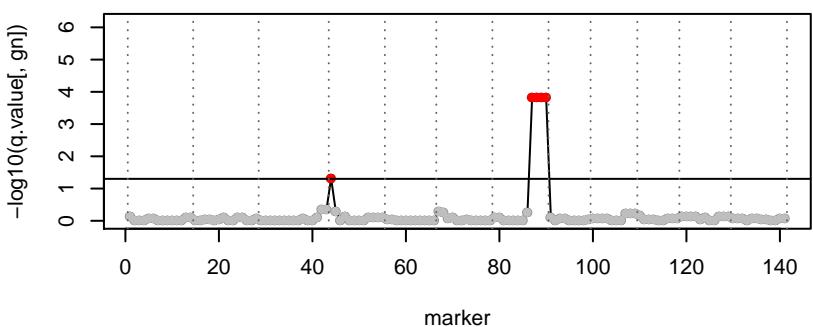
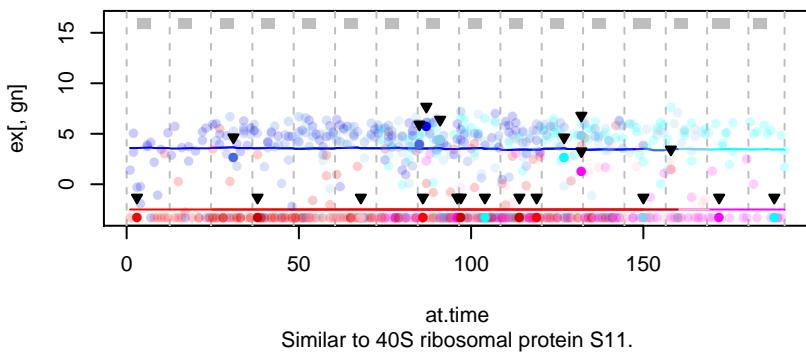
Os07g0565000 : :
1506.05869916771_Most significant each peak



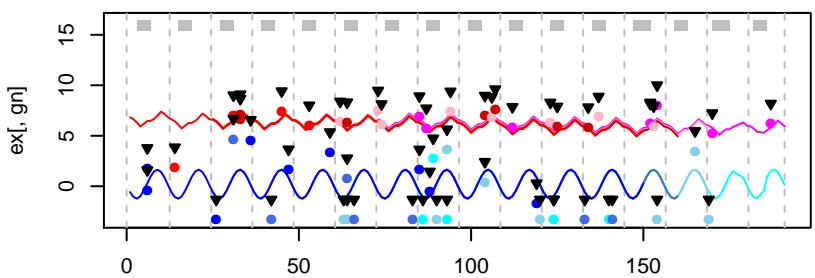
Os07g0565000 : :
Background



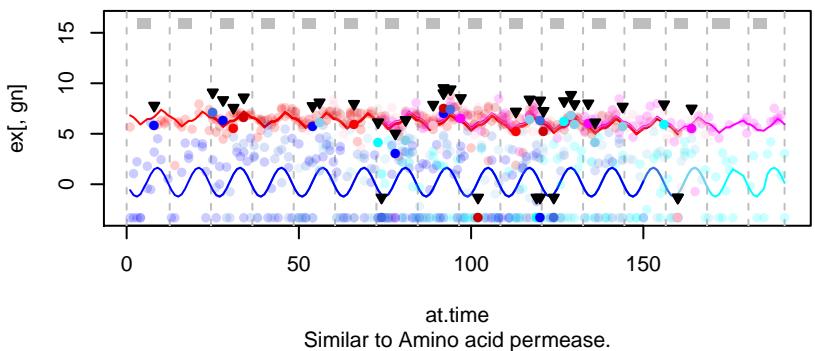
Os07g0565000 : :
1510.90643991244_Most significant peak



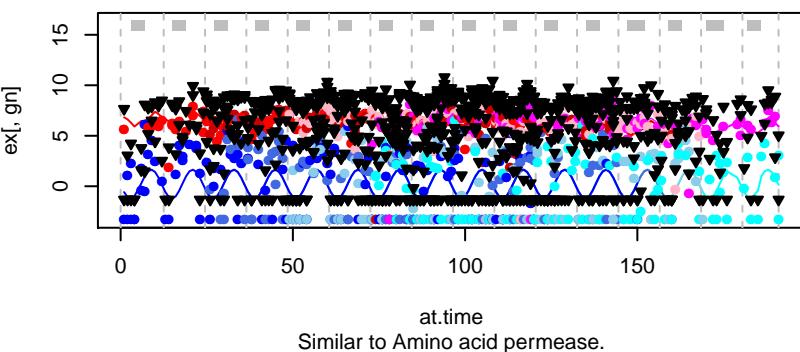
Os07g0100800 : AT2G39890.1 : ATPROT1,PROT1
kos_tak



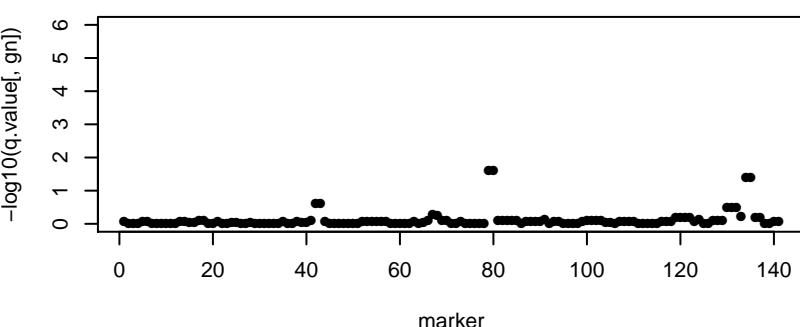
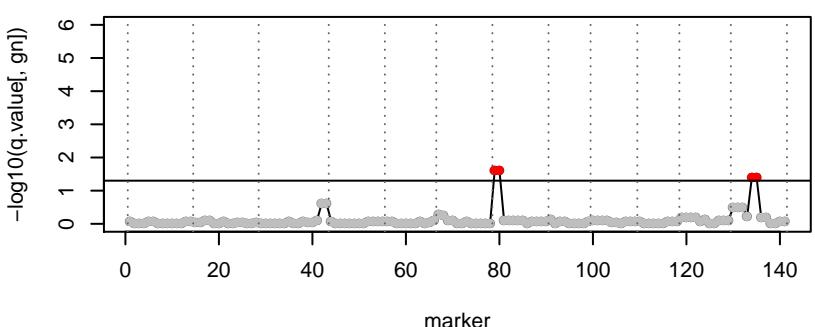
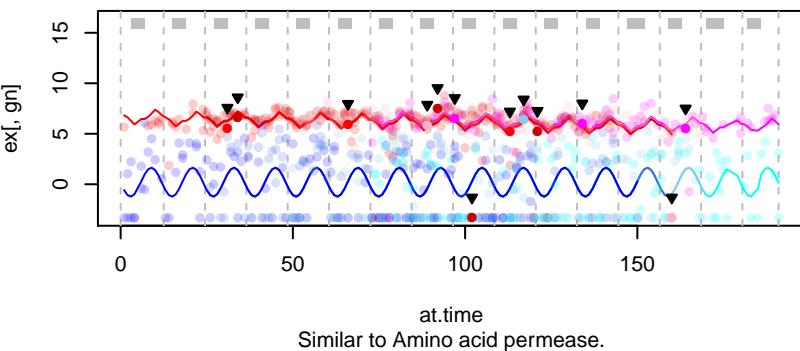
Os07g0100800 : AT2G39890.1 : ATPROT1,PROT1
1582.88852065723_Most significant each peak



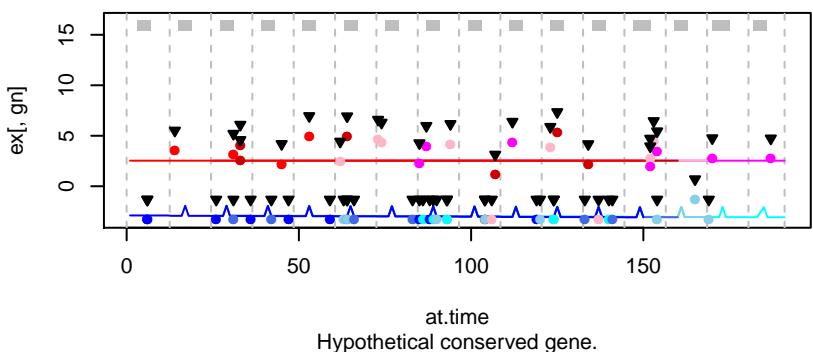
Os07g0100800 : AT2G39890.1 : ATPROT1,PROT1
Background



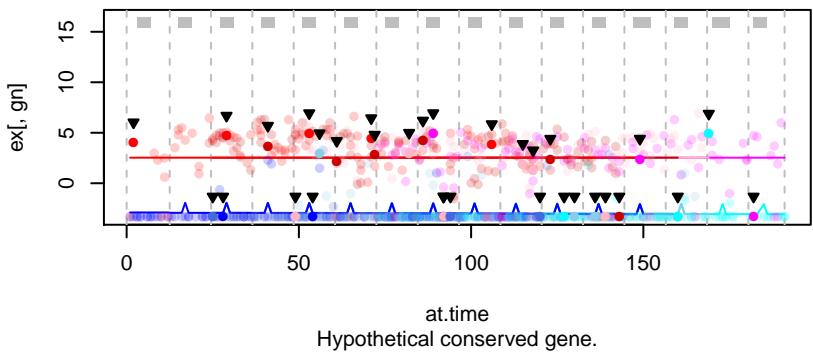
Os07g0100800 : AT2G39890.1 : ATPROT1,PROT1
1540.60502491533_Most significant peak



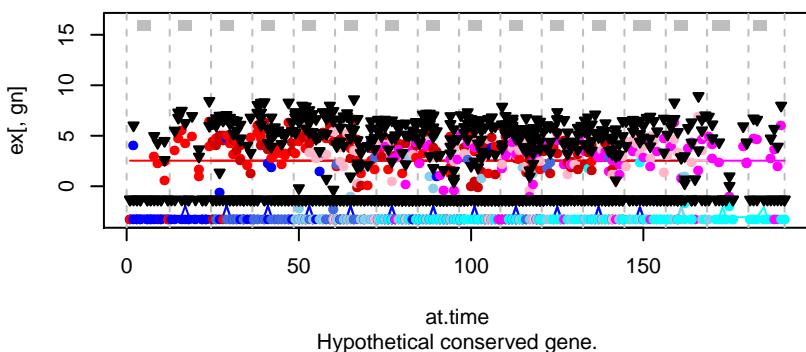
Os12g0487832 : :
kos_tak



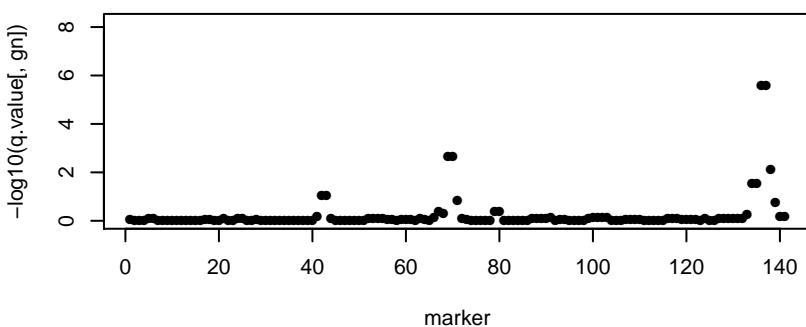
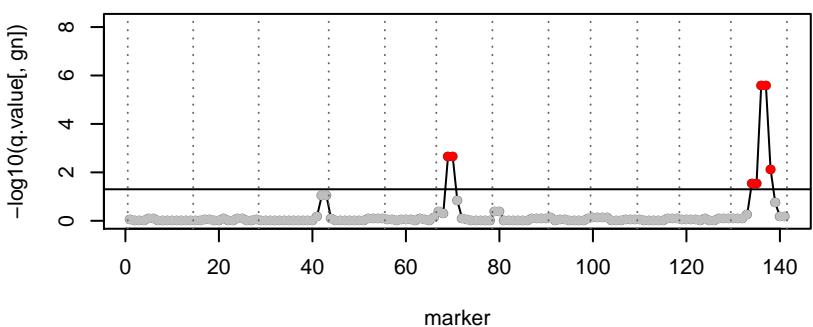
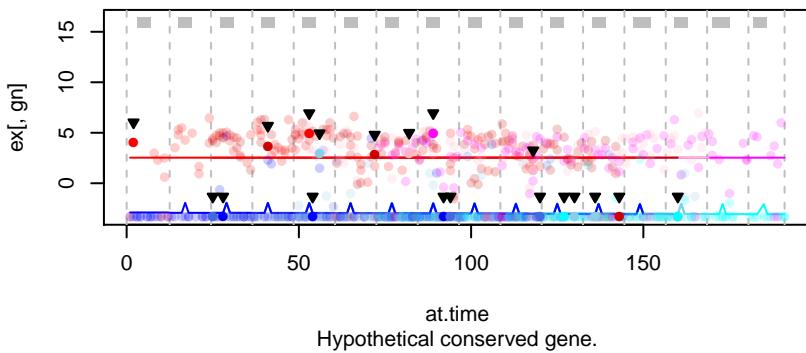
Os12g0487832 : :
1105.43574358173_Most significant each peak



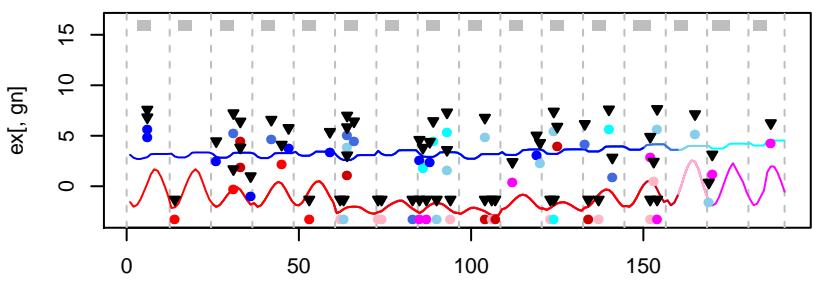
Os12g0487832 : :
Background



Os12g0487832 : :
1117.59540556571_Most significant peak

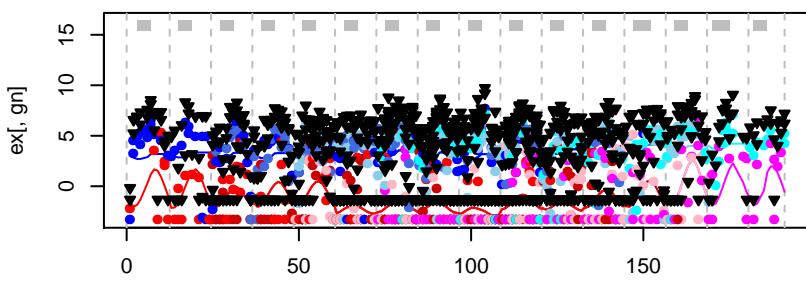


Os06g0154400 : AT1G29120.5 :
kos_tak



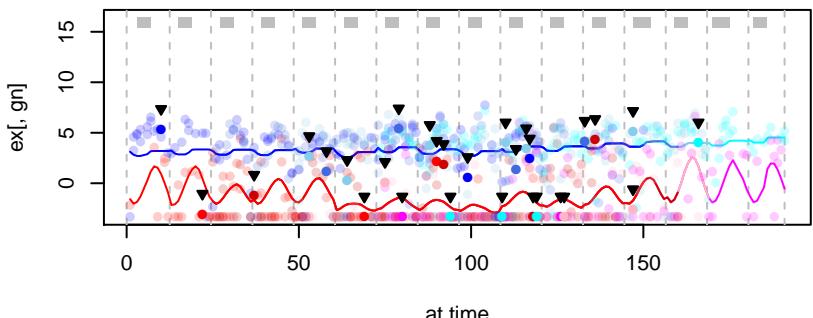
Domain of unknown function DUF676, lipase-like domain containing protein.

Os06g0154400 : AT1G29120.5 :
Background



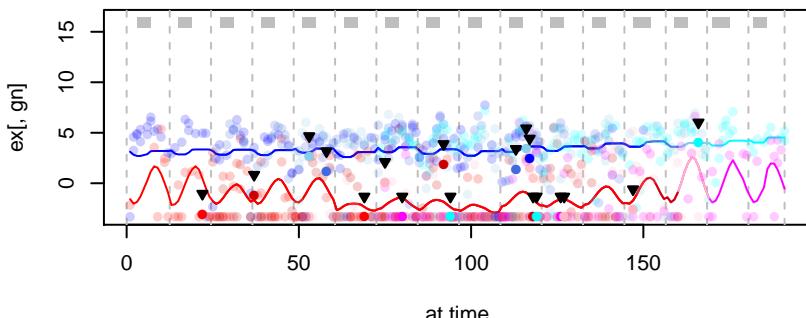
Domain of unknown function DUF676, lipase-like domain containing protein.

Os06g0154400 : AT1G29120.5 :
1679.74333846774_Most significant each peak

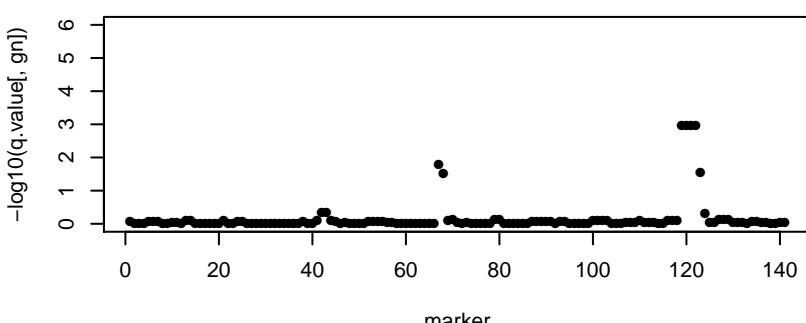
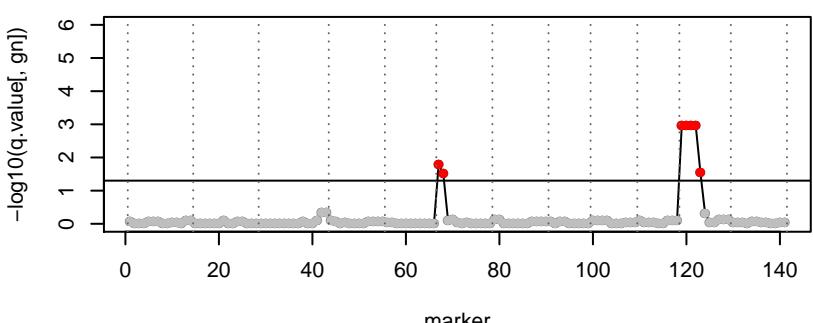


Domain of unknown function DUF676, lipase-like domain containing protein.

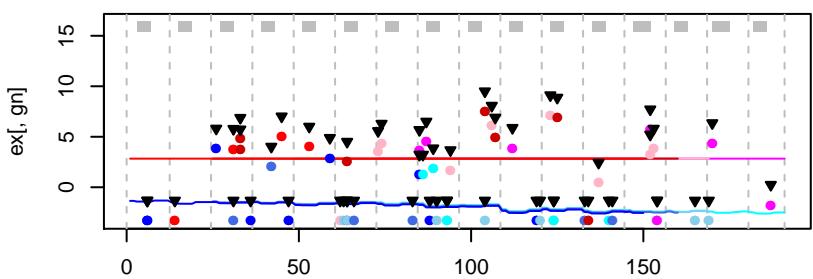
Os06g0154400 : AT1G29120.5 :
1695.6917220062_Most significant peak



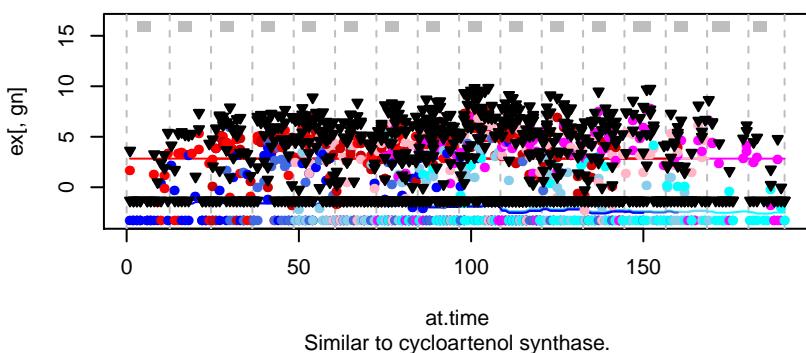
Domain of unknown function DUF676, lipase-like domain containing protein.



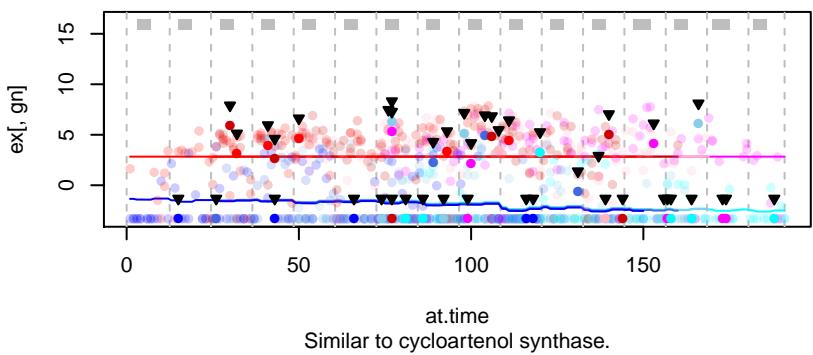
Os02g0140500 : AT5G49760.1 :
kos_tak



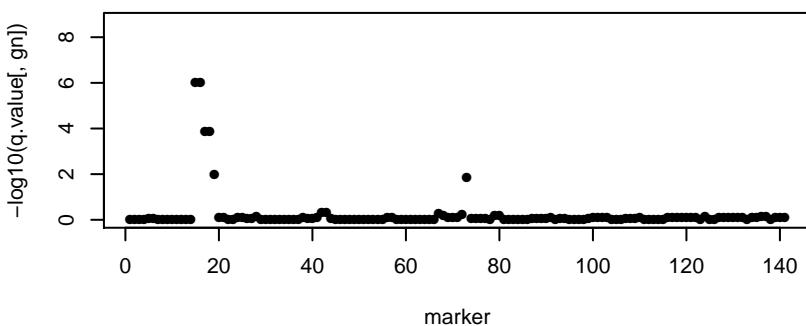
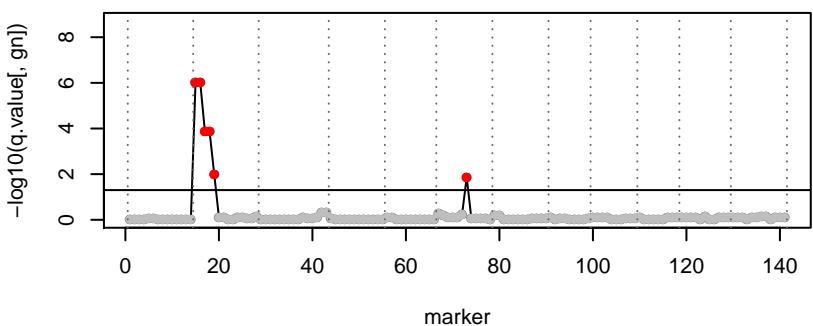
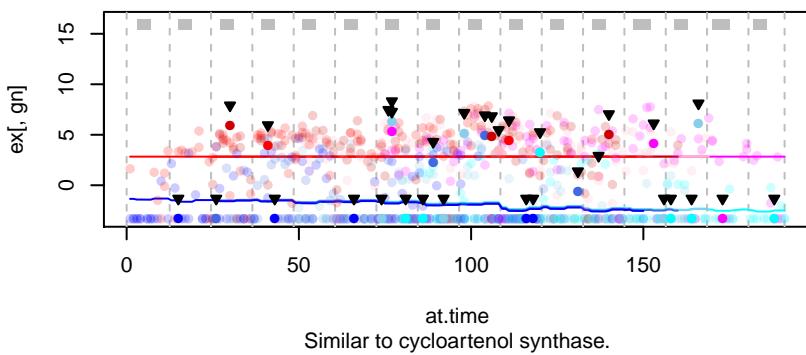
Os02g0140500 : AT5G49760.1 :
Background



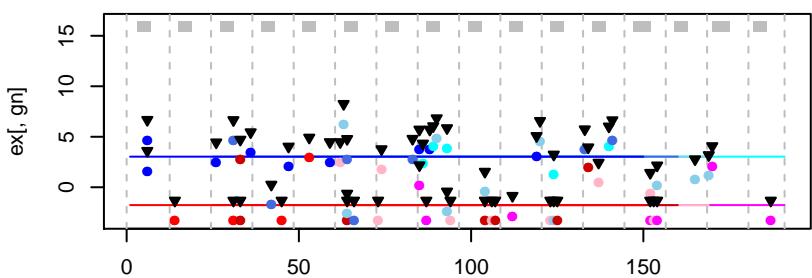
Os02g0140500 : AT5G49760.1 :
1847.33269685115_Most significant each peak



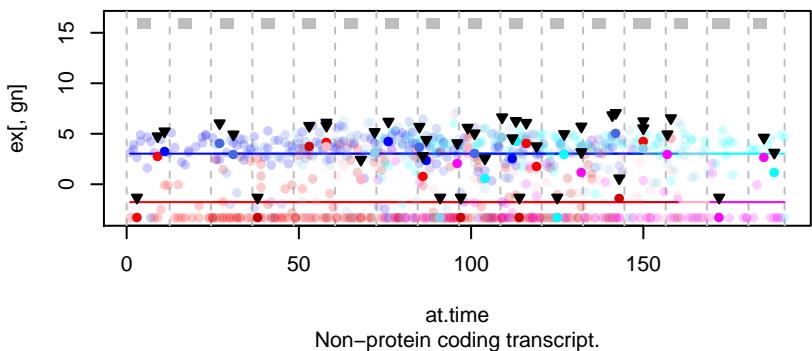
Os02g0140500 : AT5G49760.1 :
1838.09766406407_Most significant peak



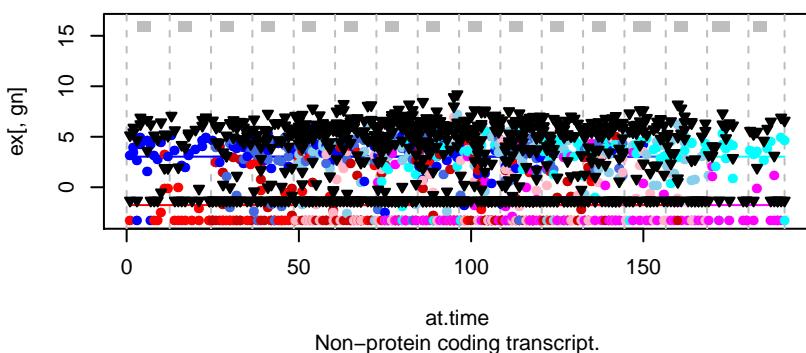
Os07g0579050 : :
kos_tak



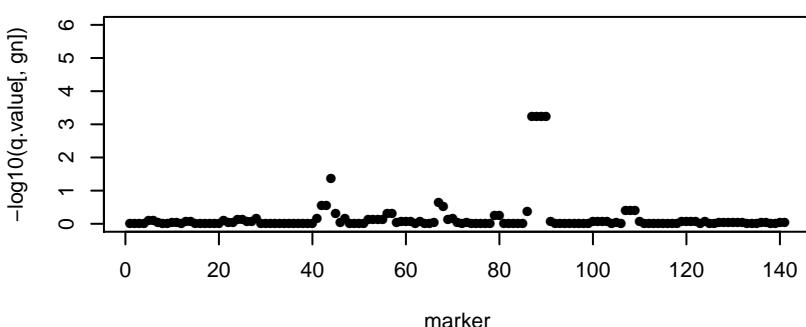
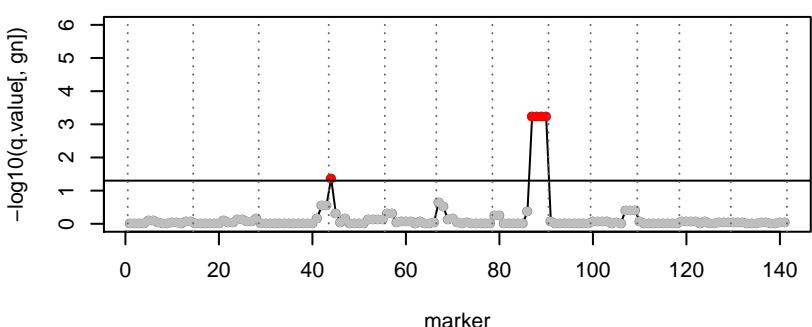
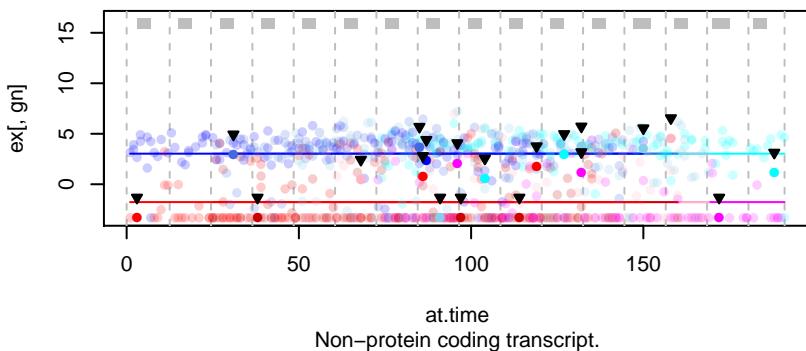
Os07g0579050 : :
1550.85485500743_Most significant each peak



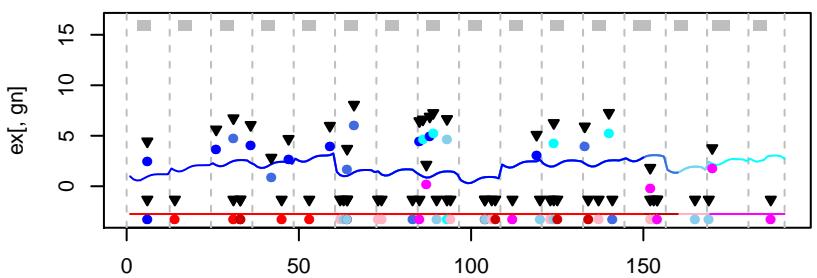
Os07g0579050 : :
Background



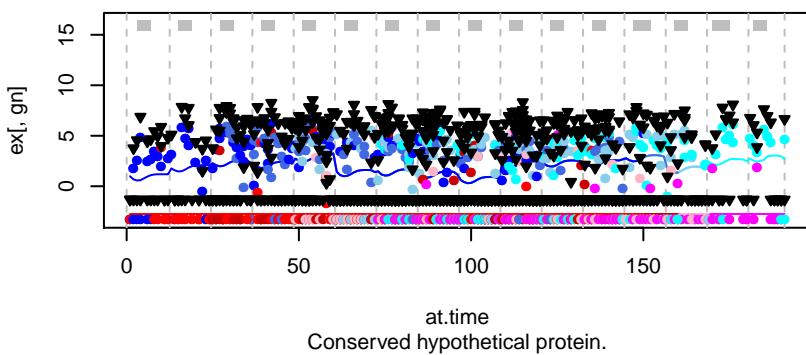
Os07g0579050 : :
1555.54203872301_Most significant peak



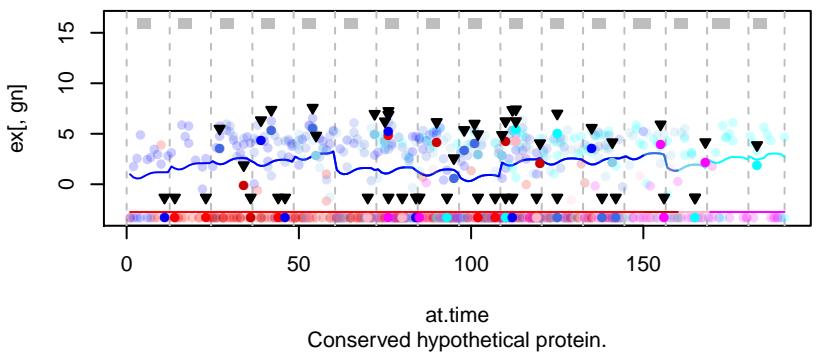
Os04g0290400 : AT2G22330.1 : CYP79B3
kos_tak



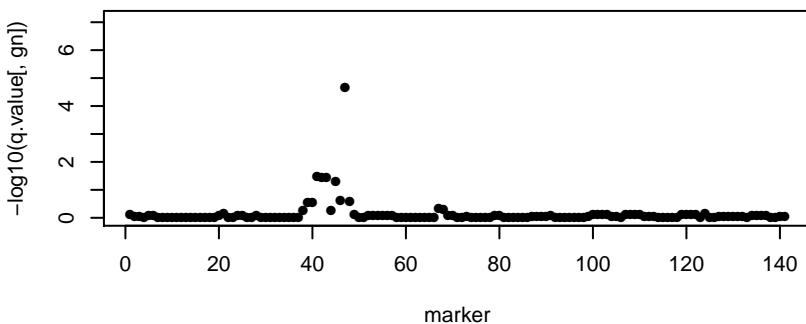
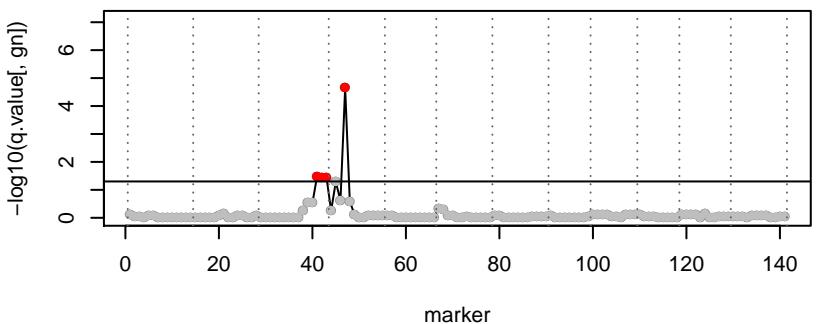
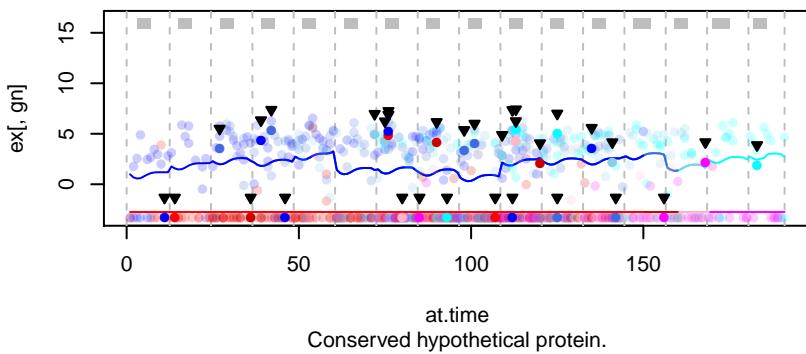
Os04g0290400 : AT2G22330.1 : CYP79B3
Background



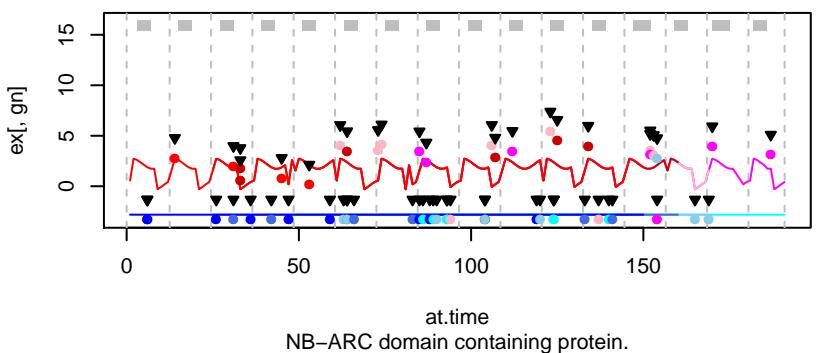
Os04g0290400 : AT2G22330.1 : CYP79B3
1663.39591288672_Most significant each peak



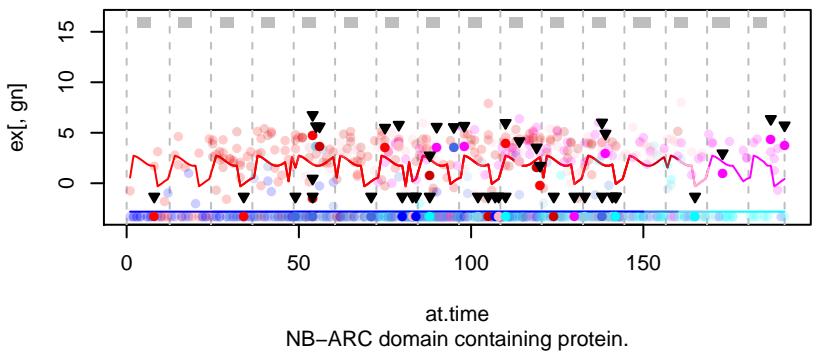
Os04g0290400 : AT2G22330.1 : CYP79B3
1682.81283877412_Most significant peak



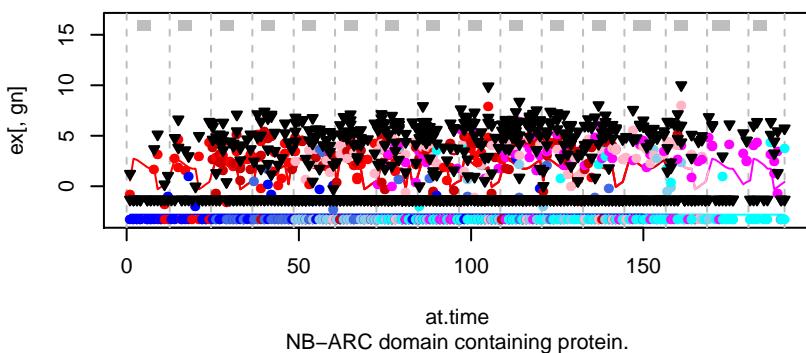
Os08g0539900 : AT3G07040.1 : RPM1,RPS3
kos_tak



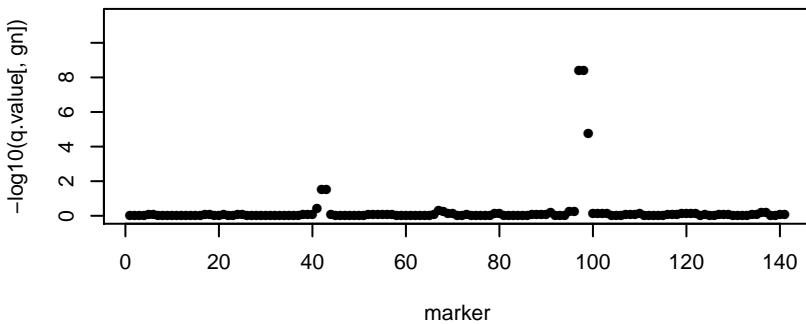
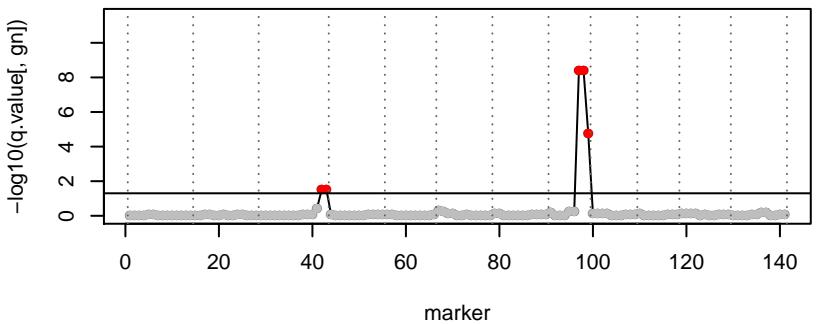
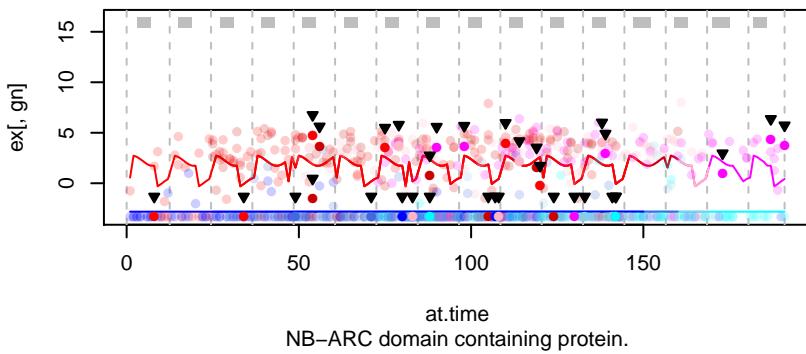
Os08g0539900 : AT3G07040.1 : RPM1,RPS3
1296.86021838894_Most significant each peak



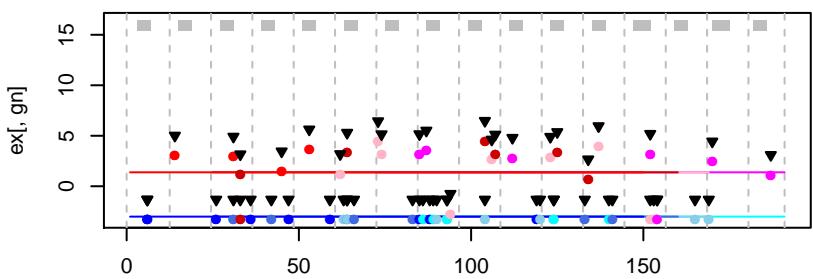
Os08g0539900 : AT3G07040.1 : RPM1,RPS3
Background



Os08g0539900 : AT3G07040.1 : RPM1,RPS3
1314.68514091922_Most significant peak

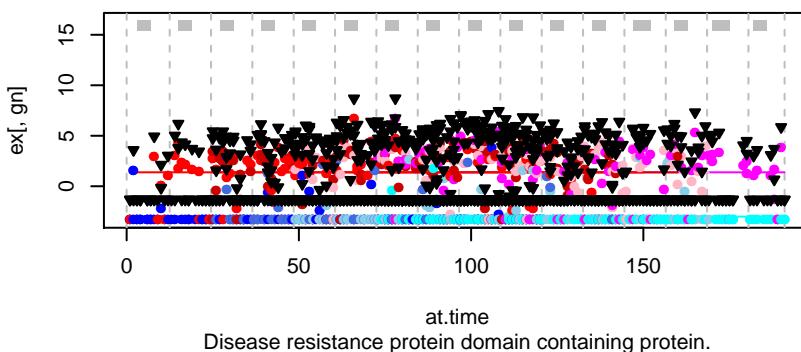


Os12g0481700 : AT3G14470.1 :
kos_tak



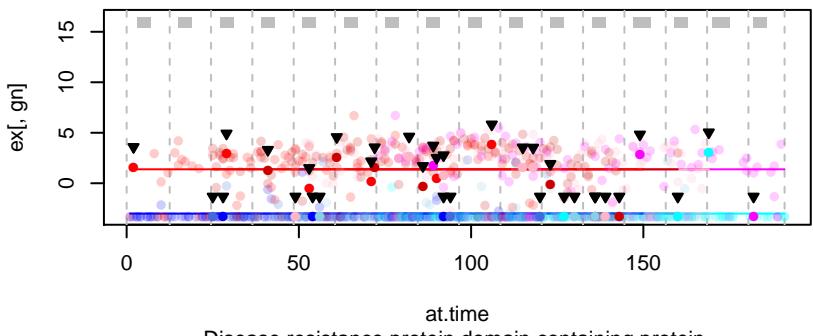
Disease resistance protein domain containing protein.

Os12g0481700 : AT3G14470.1 :
Background



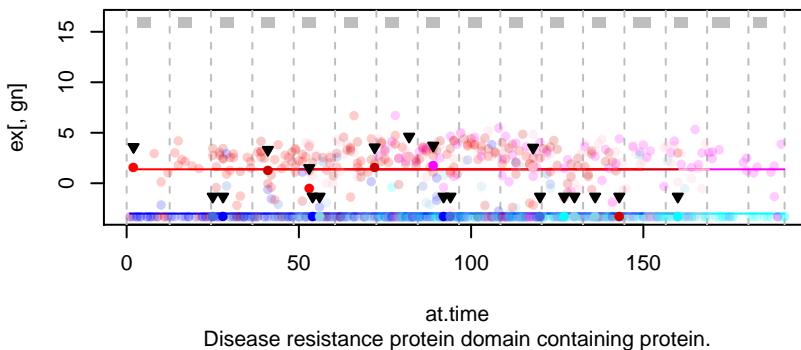
Disease resistance protein domain containing protein.

Os12g0481700 : AT3G14470.1 :
1045.66936201458_Most significant each peak

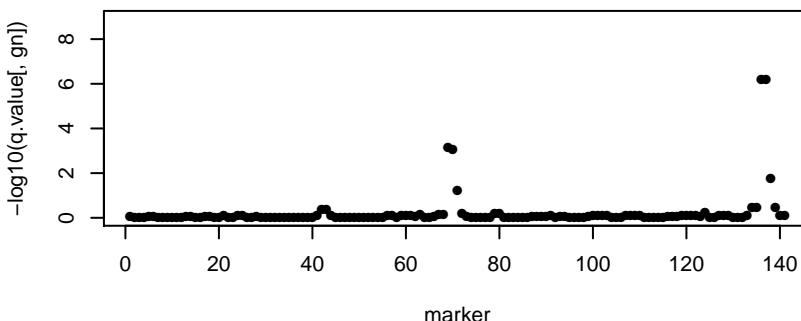
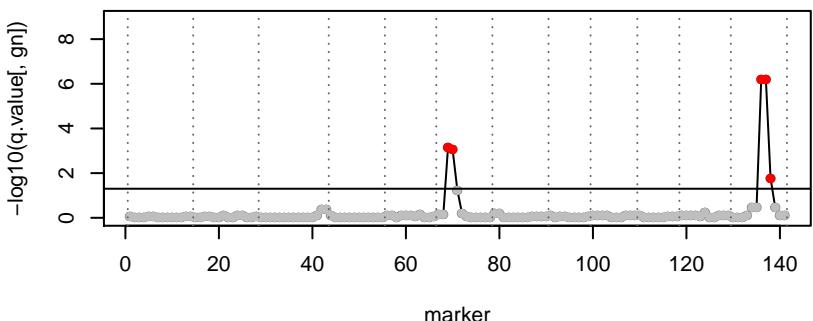


Disease resistance protein domain containing protein.

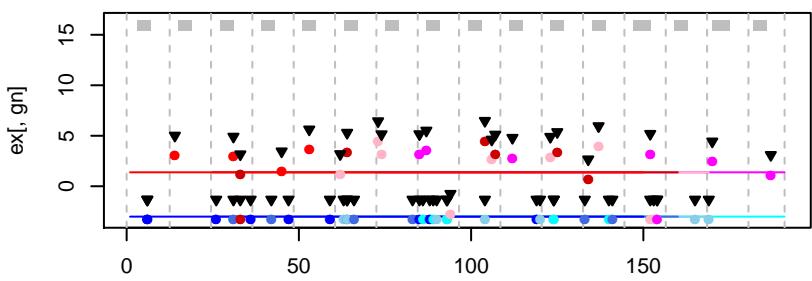
Os12g0481700 : AT3G14470.1 :
1055.96543415675_Most significant peak



Disease resistance protein domain containing protein.

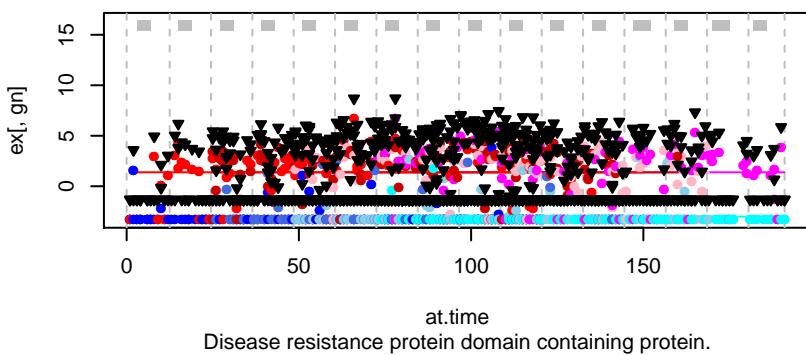


Os12g0482100 : AT1G61180.2 :
kos_tak



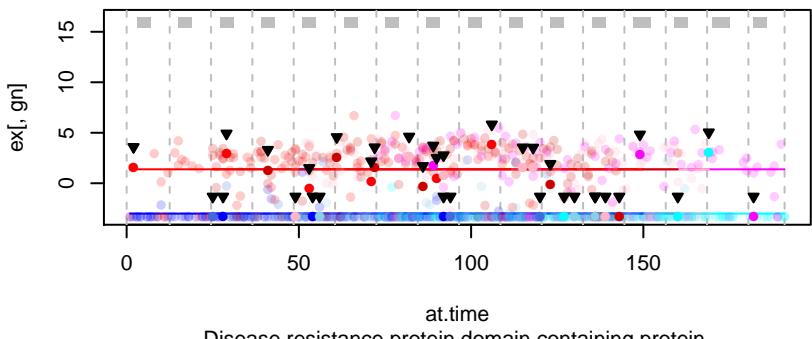
Disease resistance protein domain containing protein.

Os12g0482100 : AT1G61180.2 :
Background



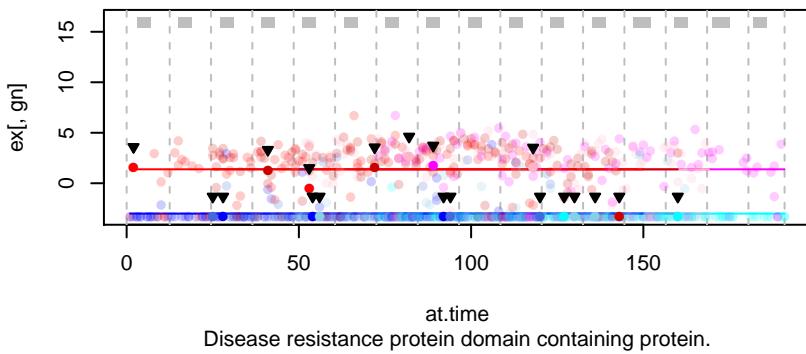
Disease resistance protein domain containing protein.

Os12g0482100 : AT1G61180.2 :
1045.66936201458_Most significant each peak

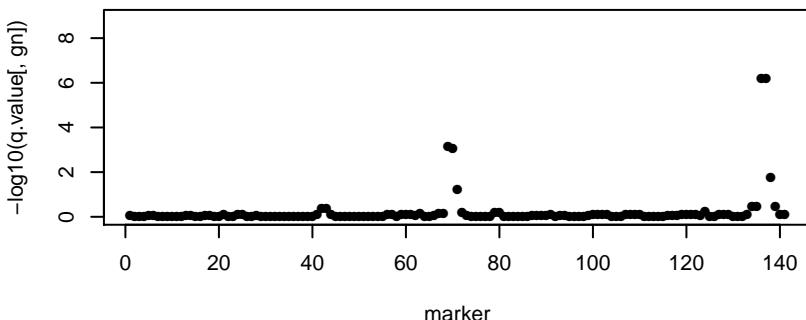
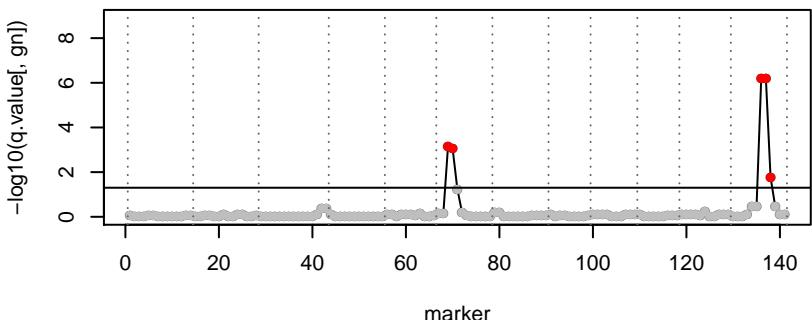


Disease resistance protein domain containing protein.

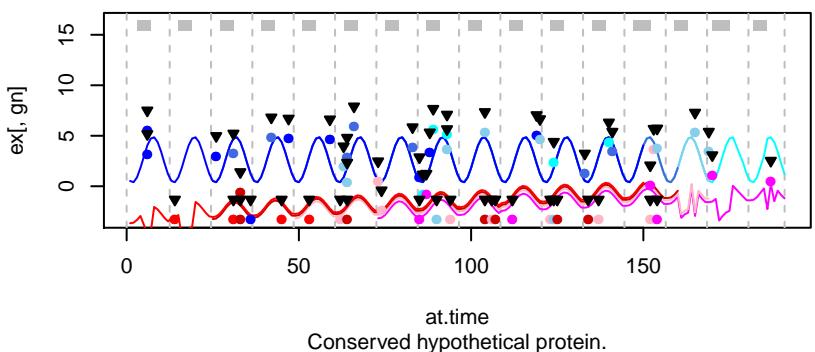
Os12g0482100 : AT1G61180.2 :
1055.96543415675_Most significant peak



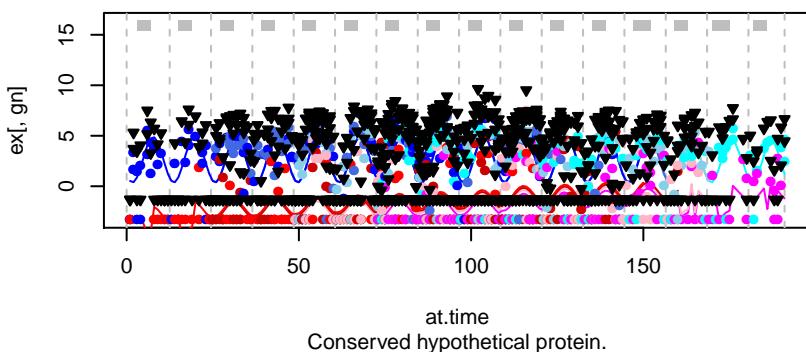
Disease resistance protein domain containing protein.



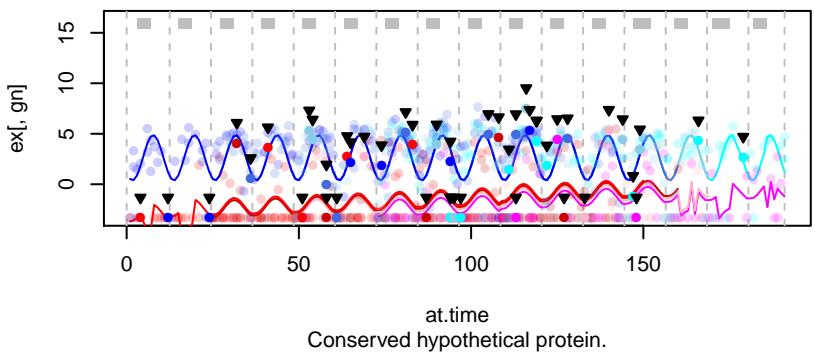
Os11g0444400 : AT5G43670.1 :
kos_tak



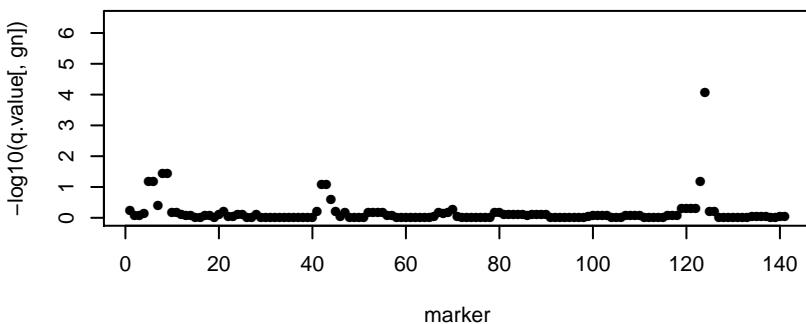
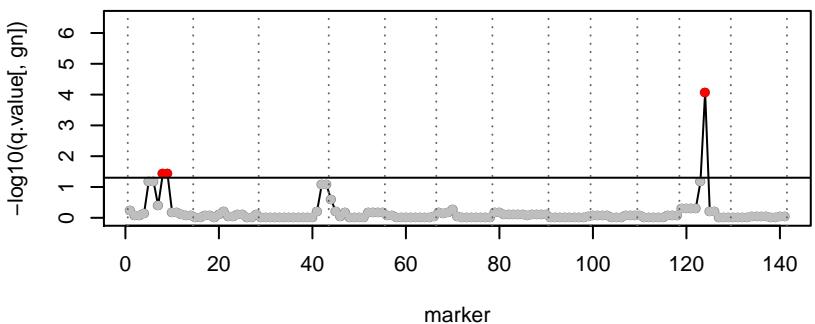
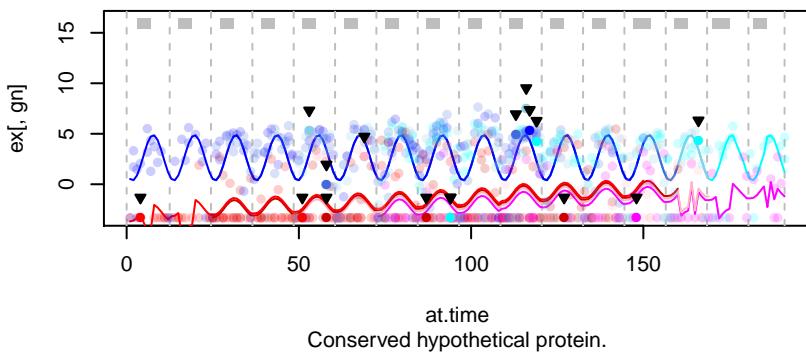
Os11g0444400 : AT5G43670.1 :
Background



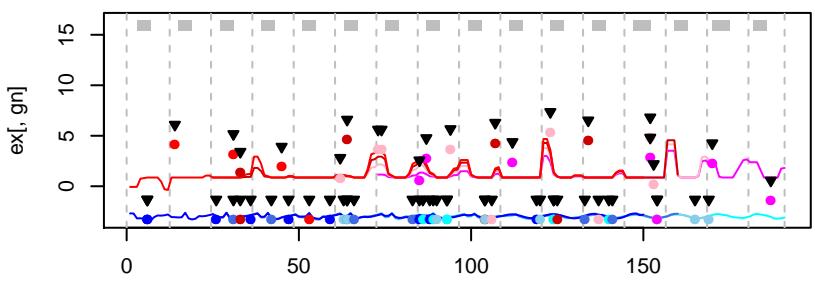
Os11g0444400 : AT5G43670.1 :
1676.39704035124_Most significant each peak



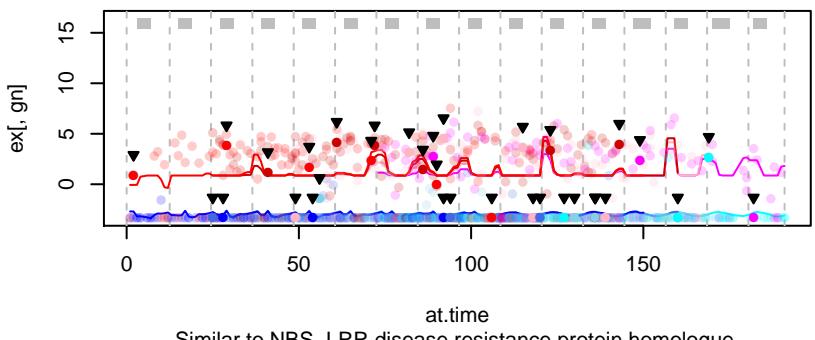
Os11g0444400 : AT5G43670.1 :
1687.88844910474_Most significant peak



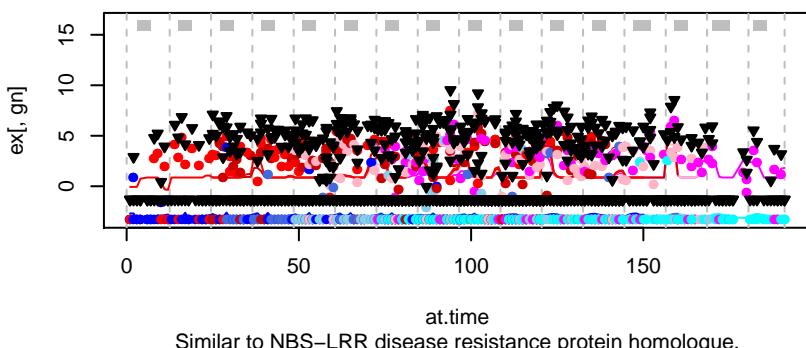
Os12g0481400 : AT3G50950.2 : ZAR1
kos_tak



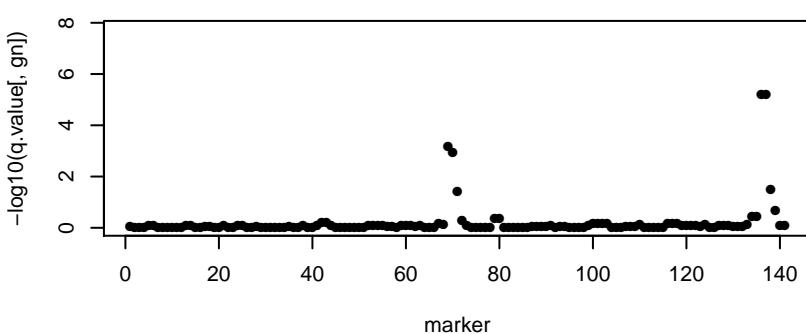
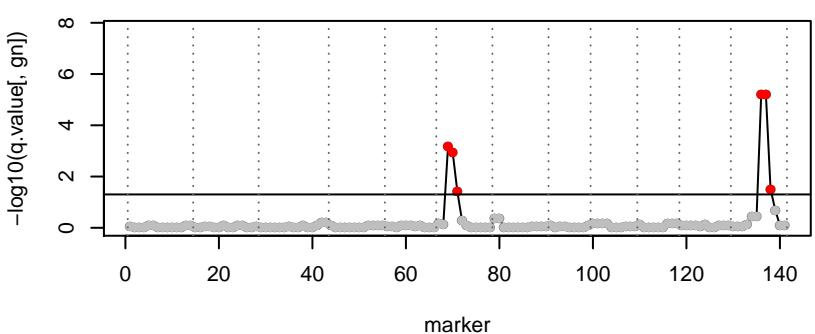
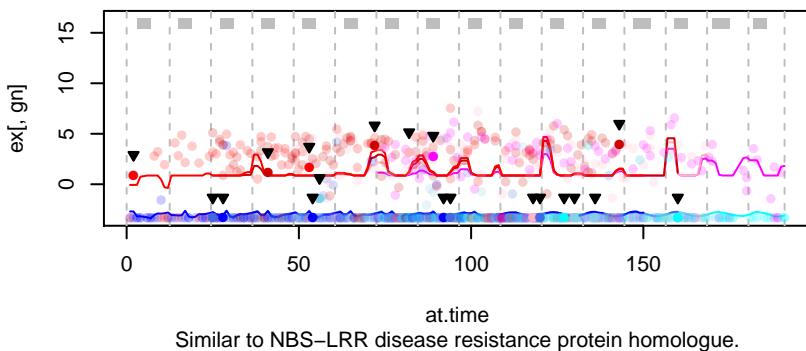
Os12g0481400 : AT3G50950.2 : ZAR1
1310.02498770097_Most significant each peak



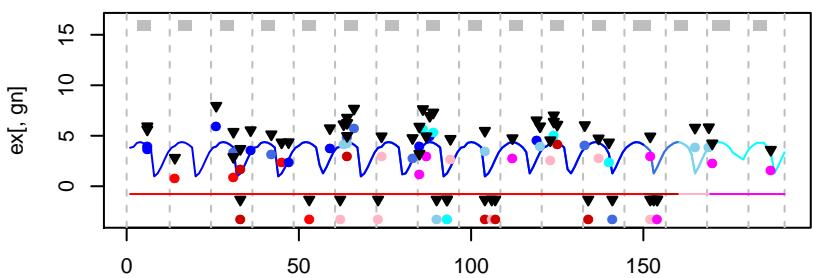
Os12g0481400 : AT3G50950.2 : ZAR1
Background



Os12g0481400 : AT3G50950.2 : ZAR1
1317.4996595753_Most significant peak

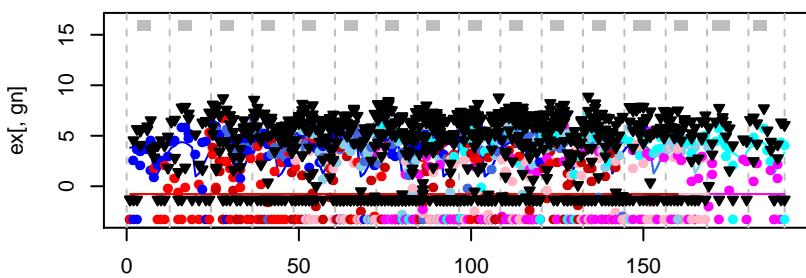


Os12g0483700 : AT4G04885.1 : PCFS4
kos_tak



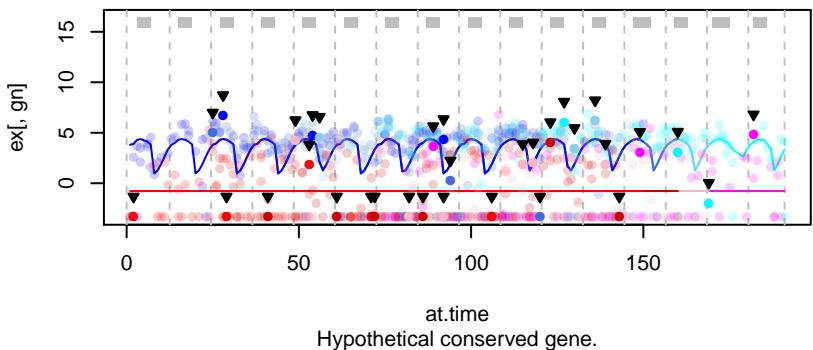
at.time
Hypothetical conserved gene.

Os12g0483700 : AT4G04885.1 : PCFS4
Background



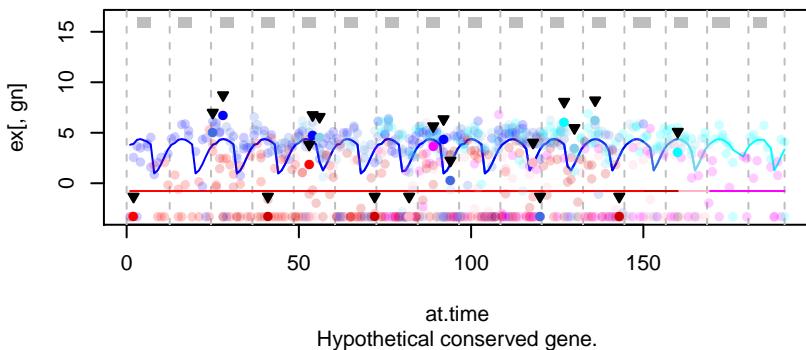
at.time
Hypothetical conserved gene.

Os12g0483700 : AT4G04885.1 : PCFS4
1861.87050634835_Most significant each peak

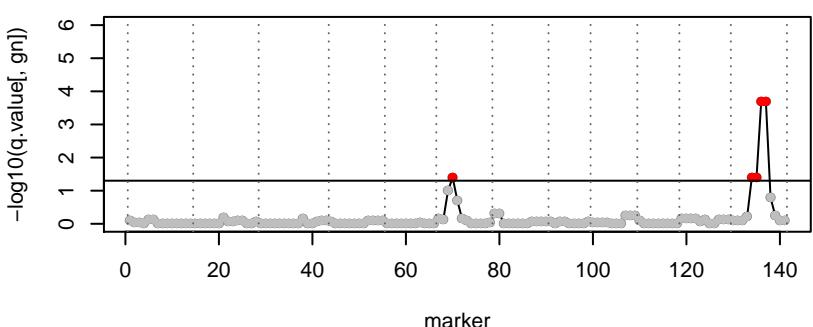


at.time
Hypothetical conserved gene.

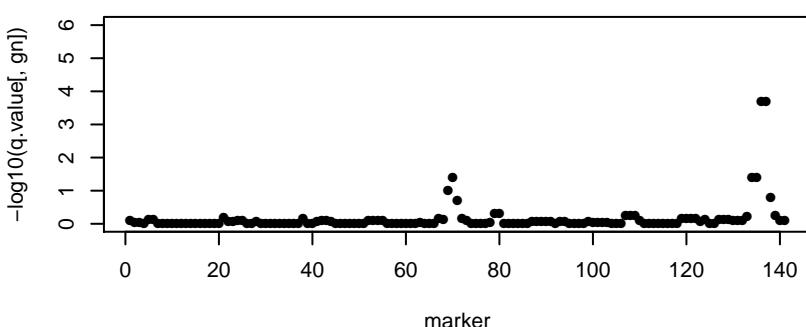
Os12g0483700 : AT4G04885.1 : PCFS4
1861.50125249628_Most significant peak



at.time
Hypothetical conserved gene.

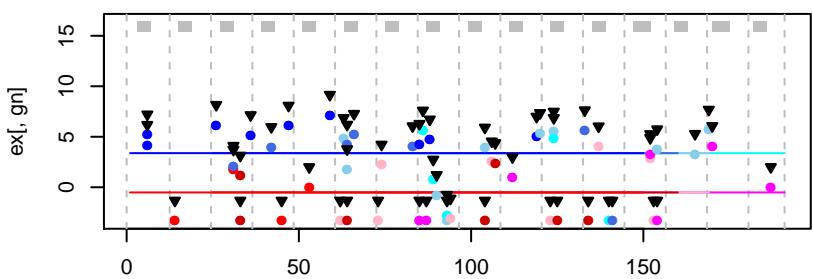


marker



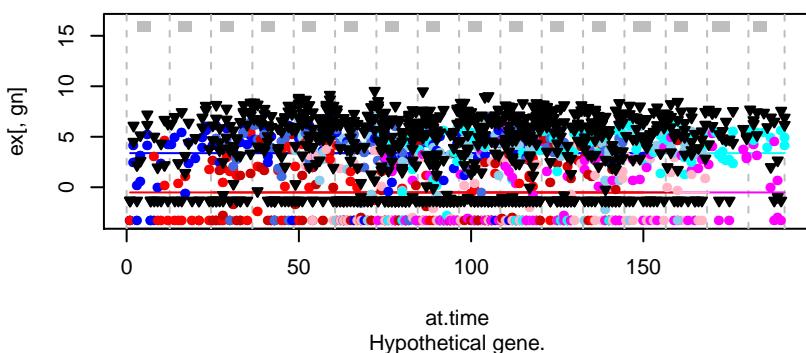
marker

Os10g0195175 : AT1G58050.1 :
kos_tak



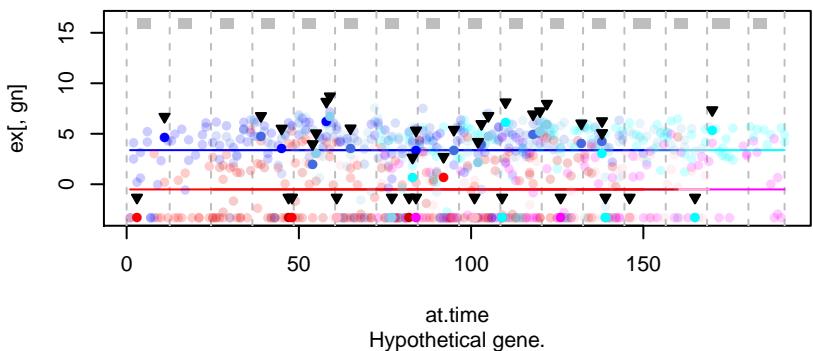
at.time
Hypothetical gene.

Os10g0195175 : AT1G58050.1 :
Background



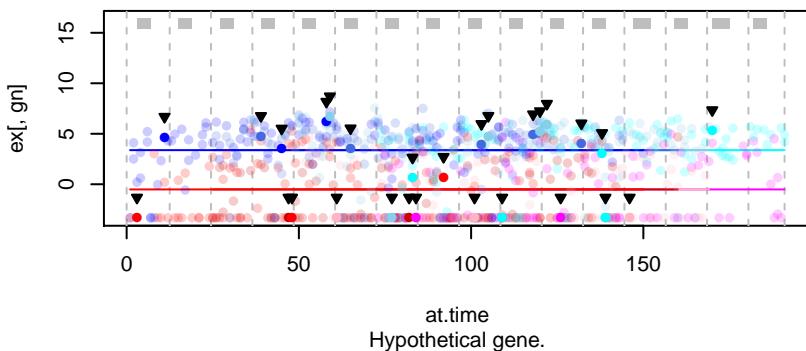
at.time
Hypothetical gene.

Os10g0195175 : AT1G58050.1 :
2052.77170506072_Most significant each peak

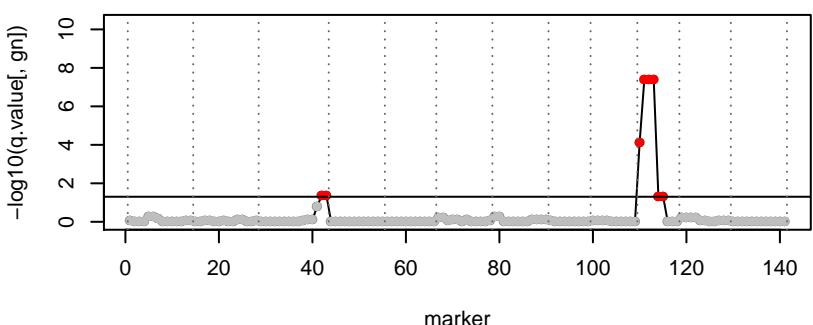


at.time
Hypothetical gene.

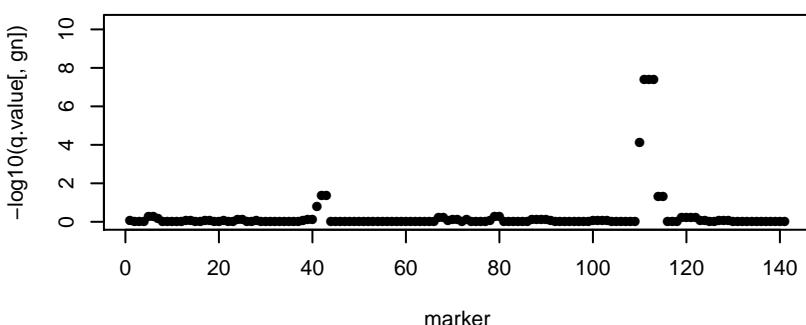
Os10g0195175 : AT1G58050.1 :
2070.20243911444_Most significant peak



at.time
Hypothetical gene.

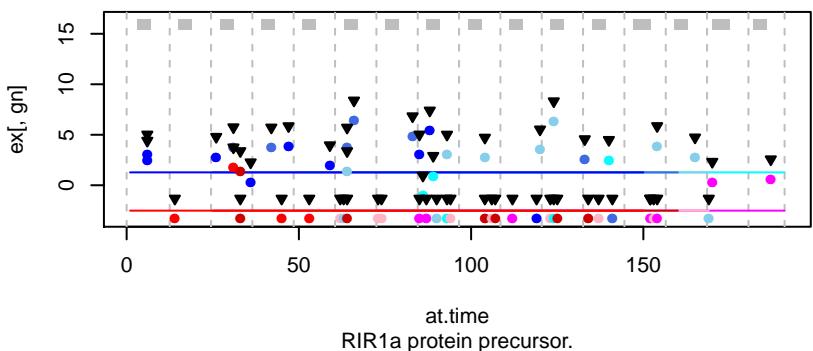


marker

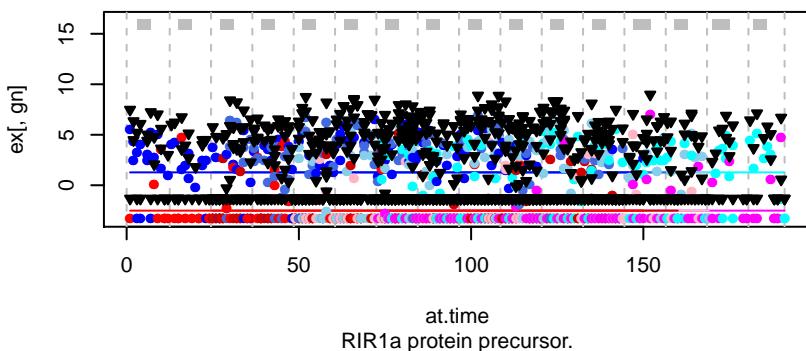


marker

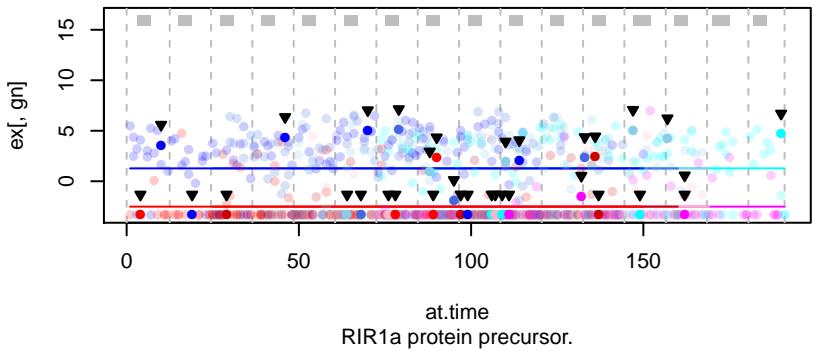
Os10g0569400 : AT4G32820.2 :
kos_tak



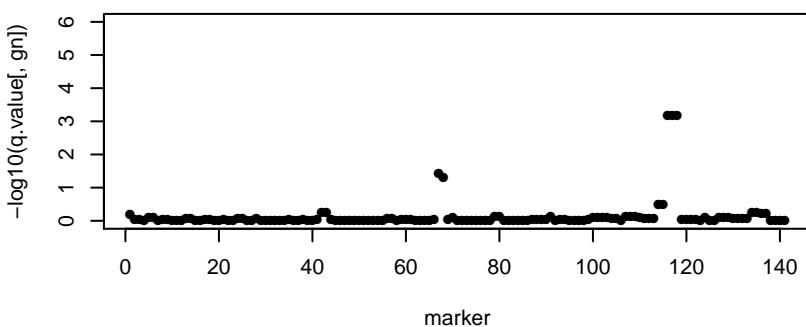
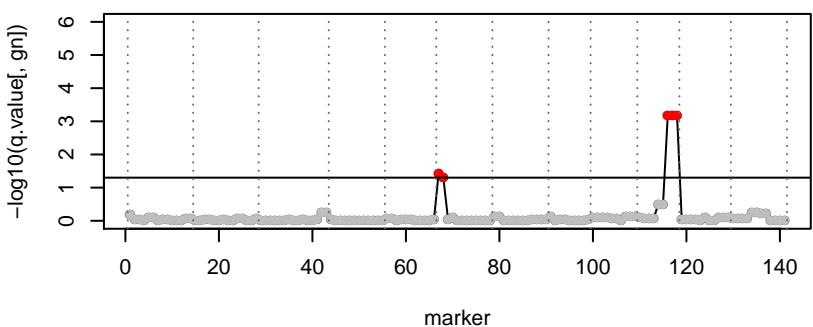
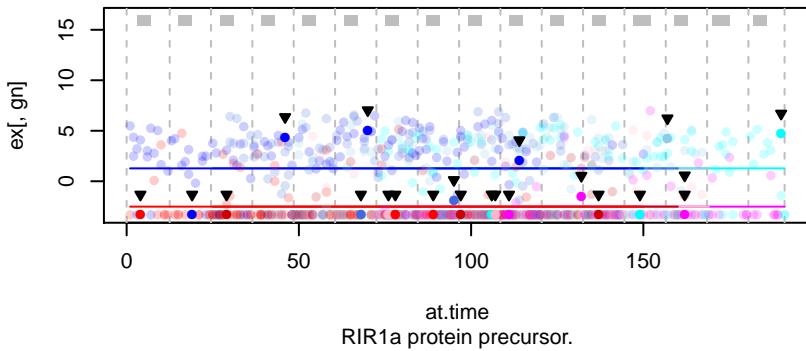
Os10g0569400 : AT4G32820.2 :
Background



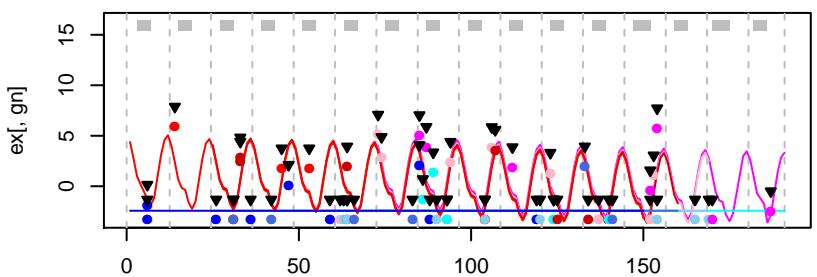
Os10g0569400 : AT4G32820.2 :
1834.83707343395_Most significant each peak



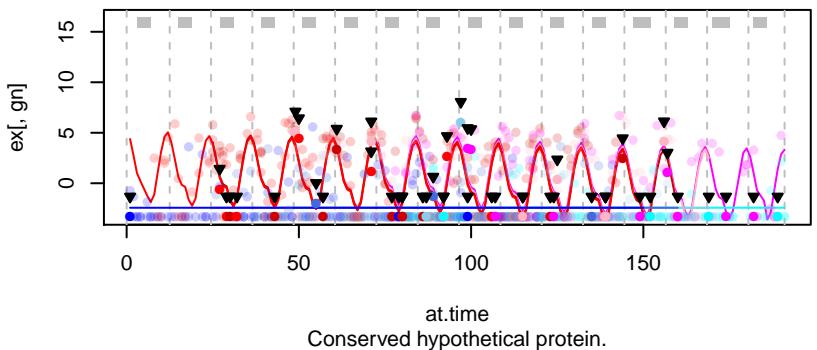
Os10g0569400 : AT4G32820.2 :
1837.96318101105_Most significant peak



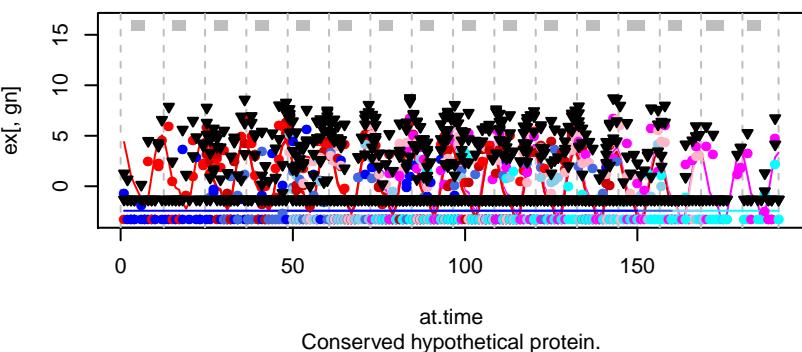
Os08g0255020 : AT1G66970.2 : GDPDL1,SVL2
kos_tak



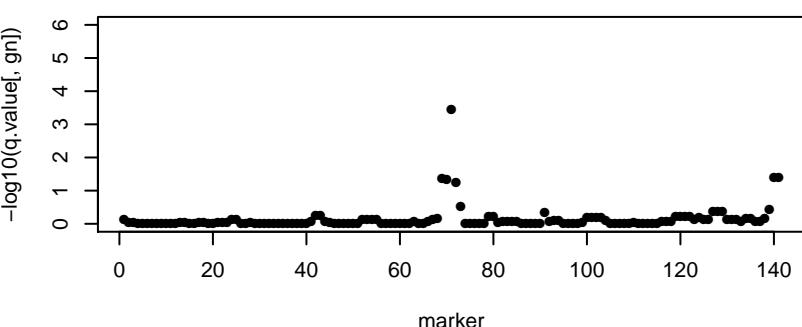
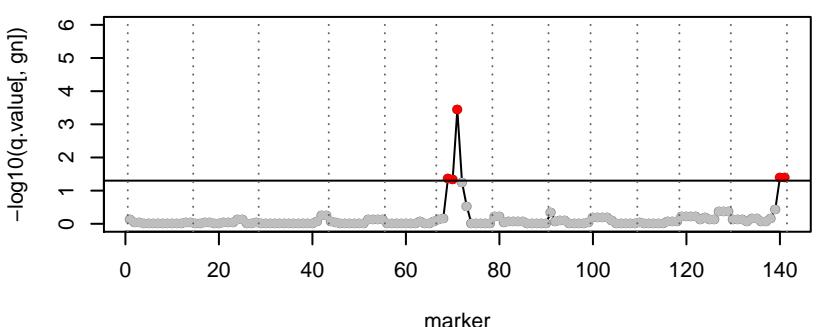
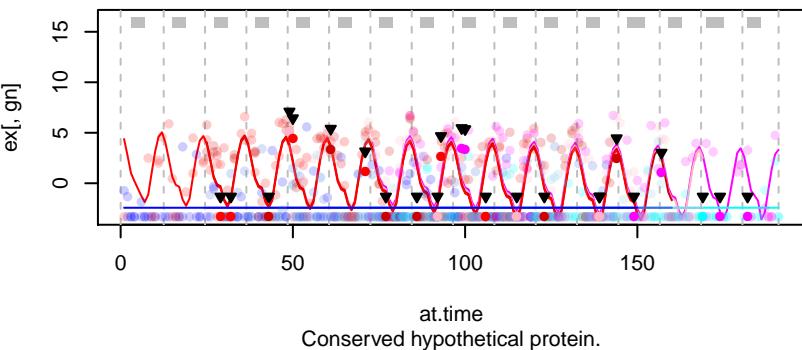
Os08g0255020 : AT1G66970.2 : GDPDL1,SVL2
1488.9747981574_Most significant each peak



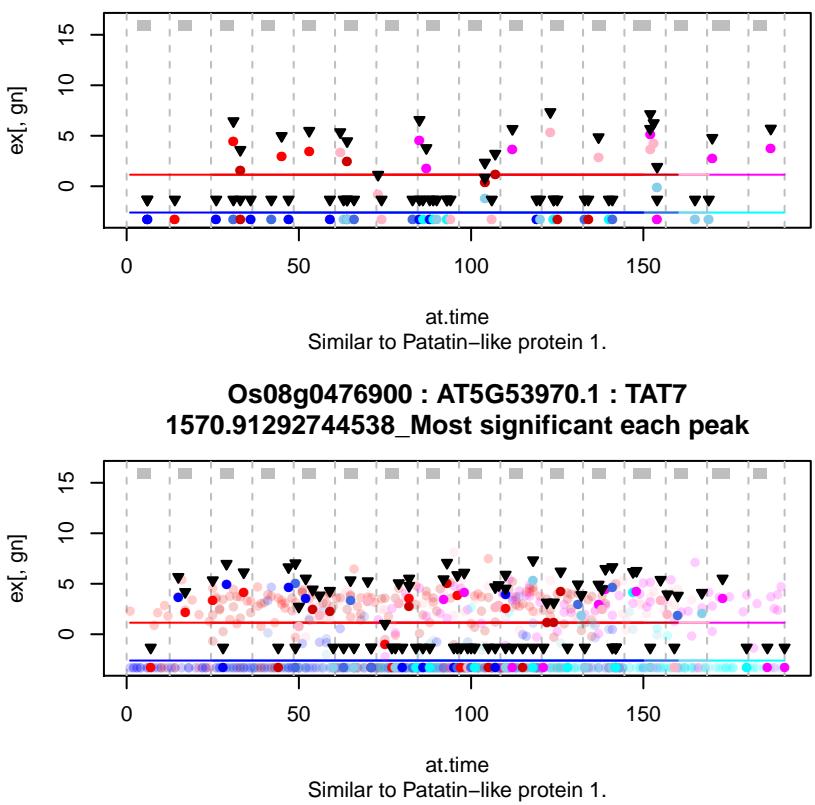
Os08g0255020 : AT1G66970.2 : GDPDL1,SVL2
Background



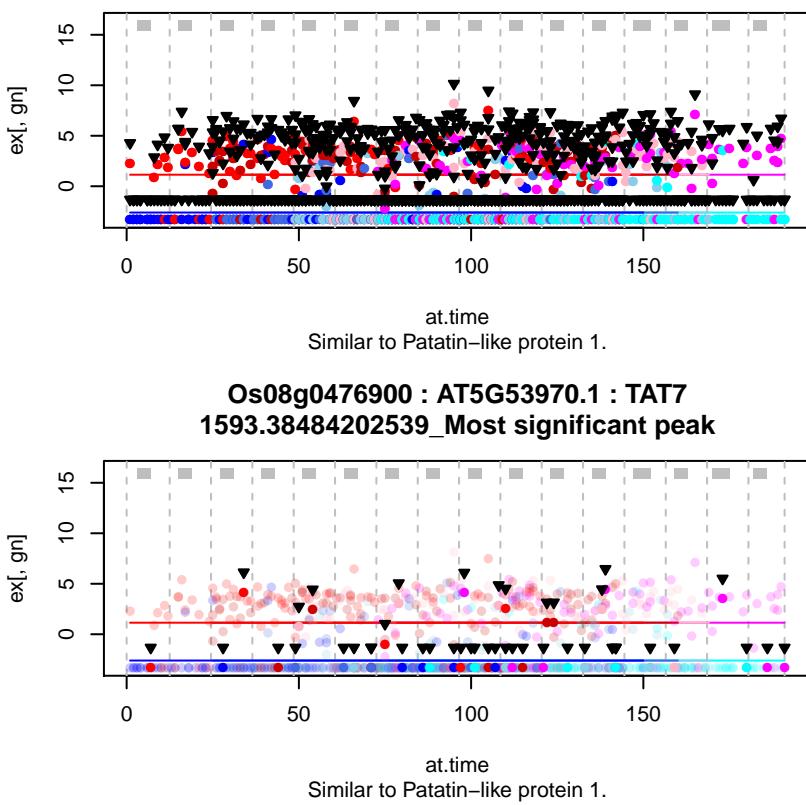
Os08g0255020 : AT1G66970.2 : GDPDL1,SVL2
1502.41373531651_Most significant peak



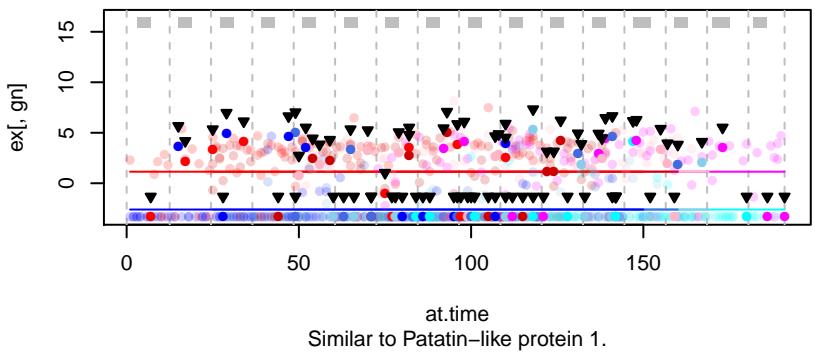
Os08g0476900 : AT5G53970.1 : TAT7
kos_tak



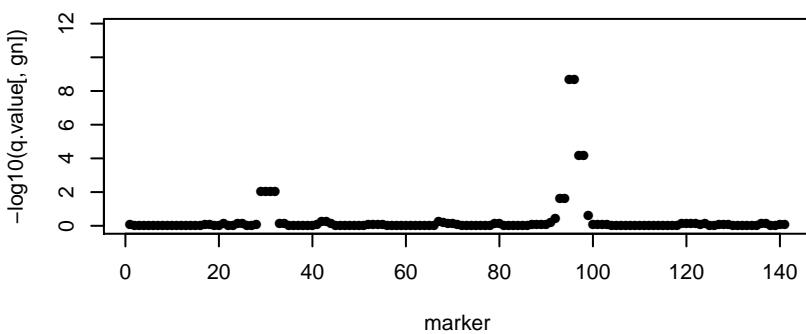
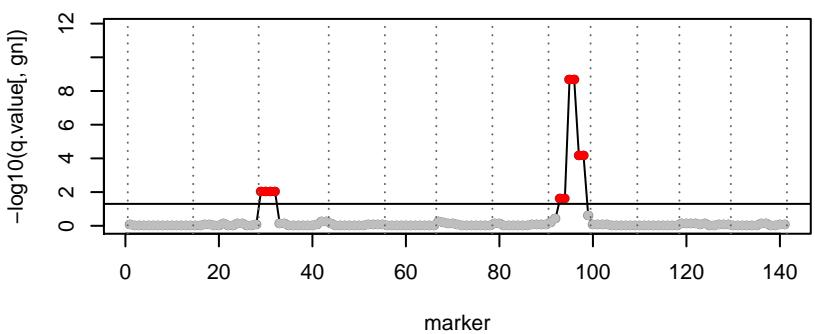
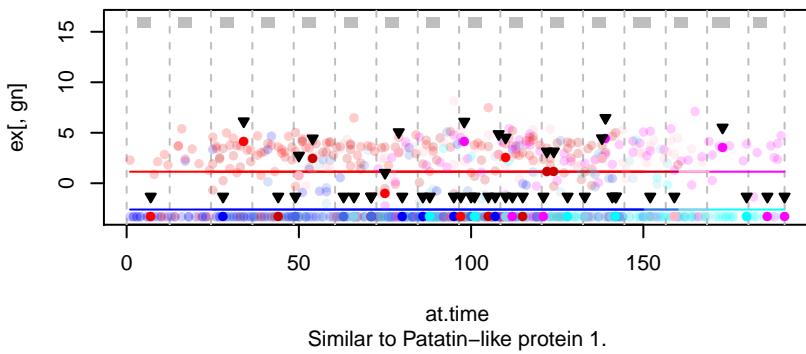
Os08g0476900 : AT5G53970.1 : TAT7
Background



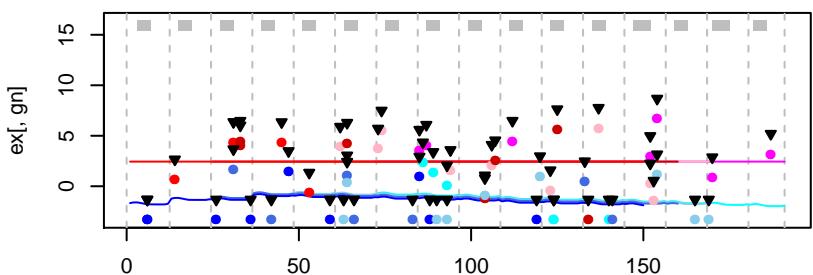
Os08g0476900 : AT5G53970.1 : TAT7
1570.91292744538_Most significant each peak



Os08g0476900 : AT5G53970.1 : TAT7
1593.38484202539_Most significant peak

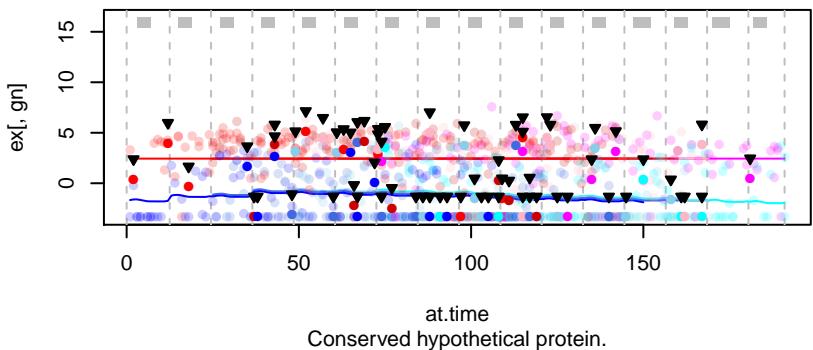


Os04g0446401 : :
kos_tak



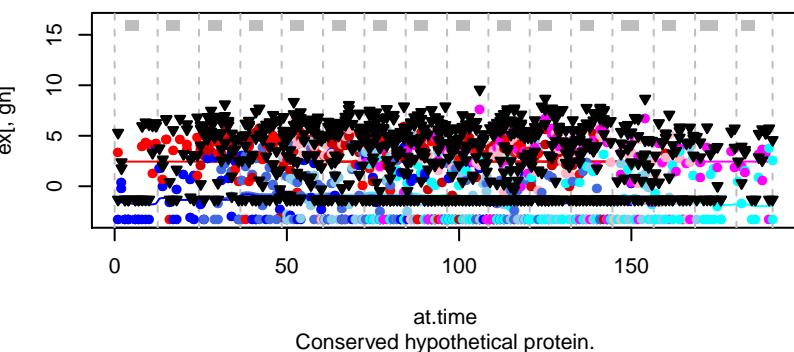
at.time
Conserved hypothetical protein.

Os04g0446401 : :
1870.04304521876_Most significant each peak



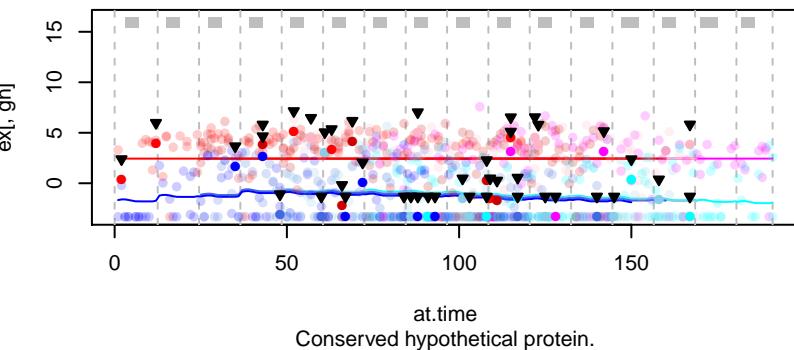
at.time
Conserved hypothetical protein.

Os04g0446401 : :
Background

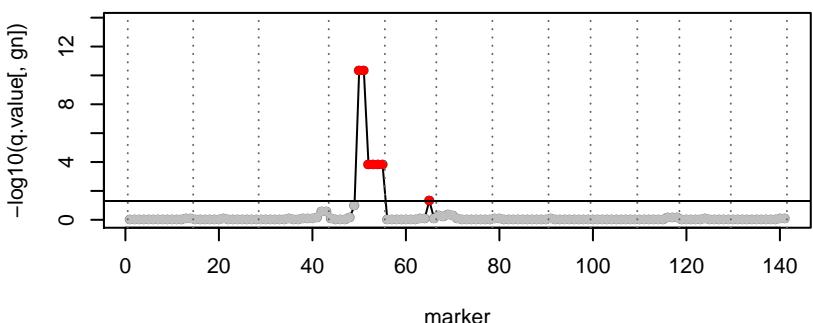


at.time
Conserved hypothetical protein.

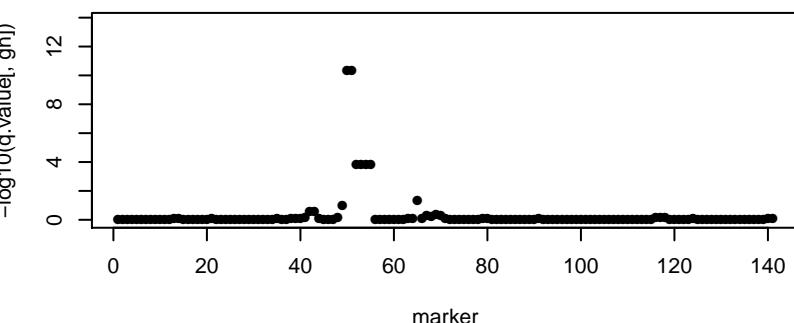
Os04g0446401 : :
1845.55064465502_Most significant peak



at.time
Conserved hypothetical protein.

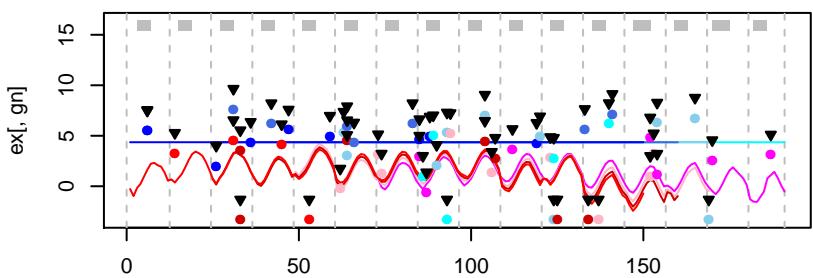


marker



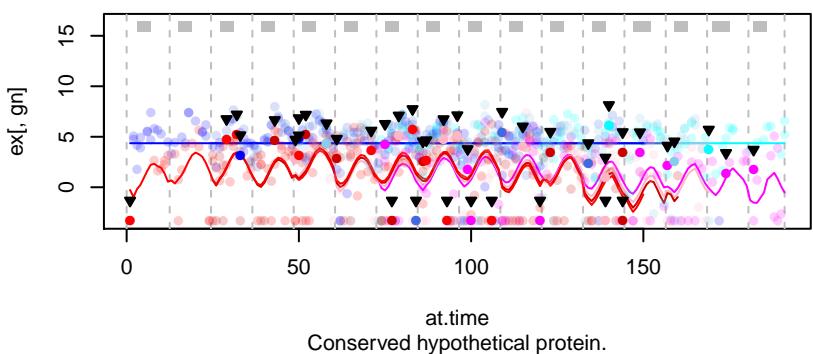
marker

Os06g0329200 : AT1G58230.1 :
kos_tak



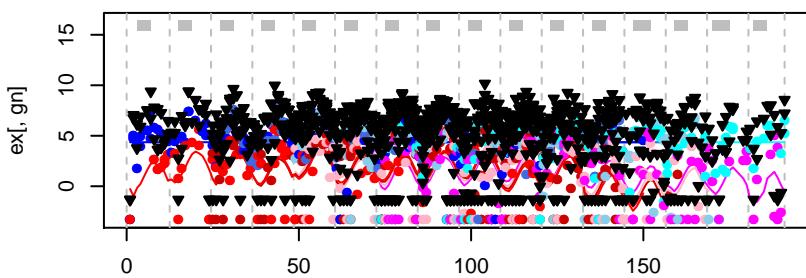
at.time
Conserved hypothetical protein.

Os06g0329200 : AT1G58230.1 :
1753.0967210382_Most significant each peak



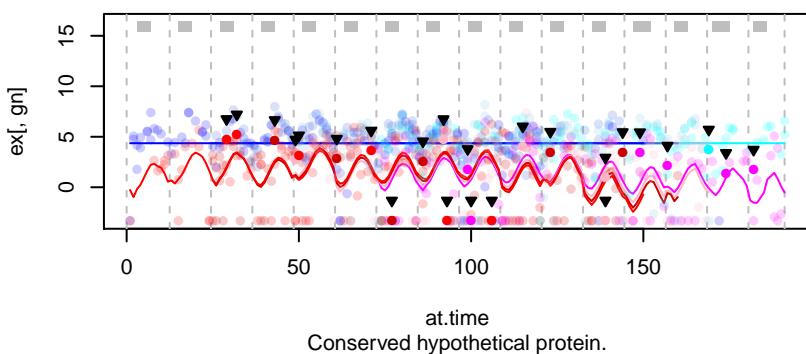
at.time
Conserved hypothetical protein.

Os06g0329200 : AT1G58230.1 :
Background

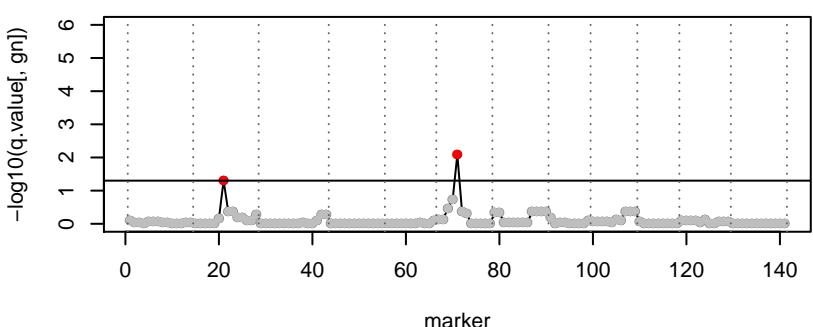


at.time
Conserved hypothetical protein.

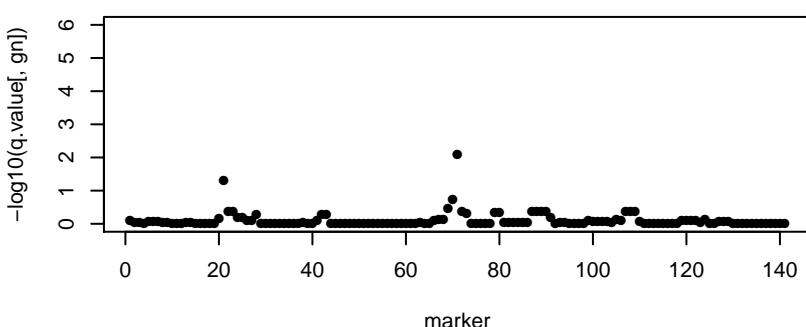
Os06g0329200 : AT1G58230.1 :
1762.77266104381_Most significant peak



at.time
Conserved hypothetical protein.

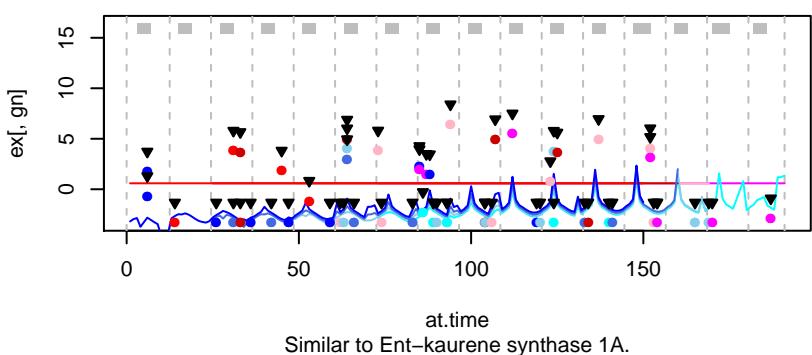


marker

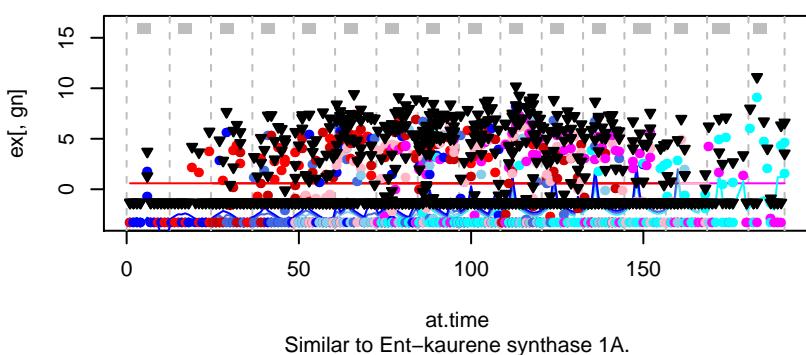


marker

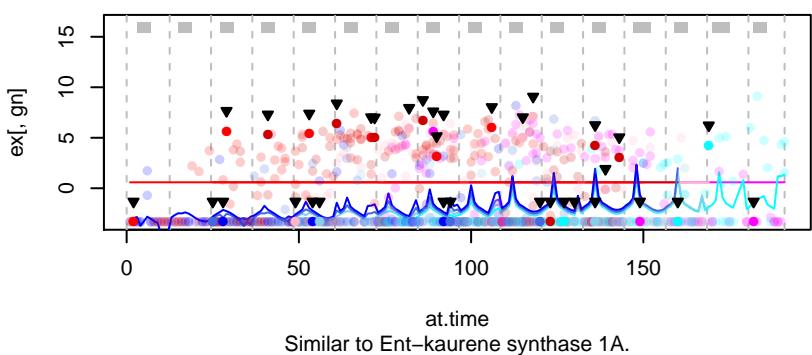
Os12g0491800 : AT1G79460.1 : ATKS,ATKS1,GA2,KS,KS1
kos_tak



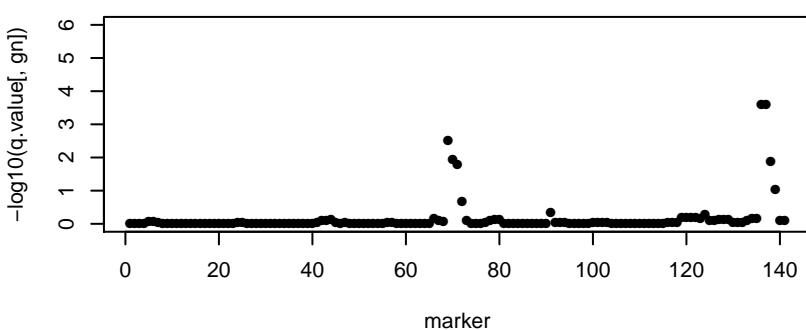
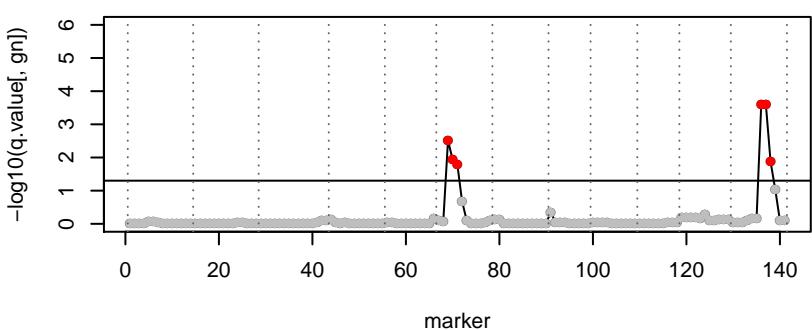
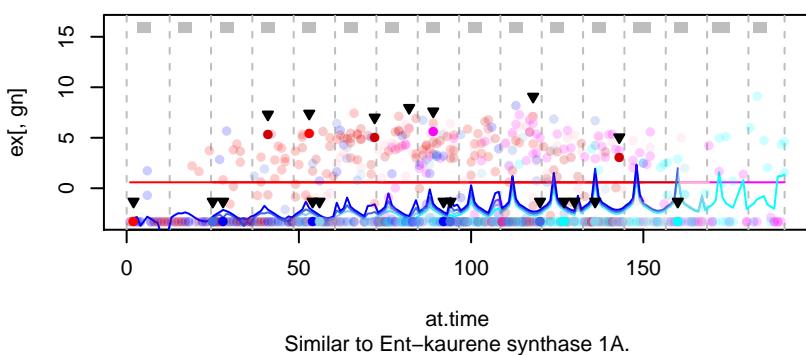
Os12g0491800 : AT1G79460.1 : ATKS,ATKS1,GA2,KS,KS1
Background

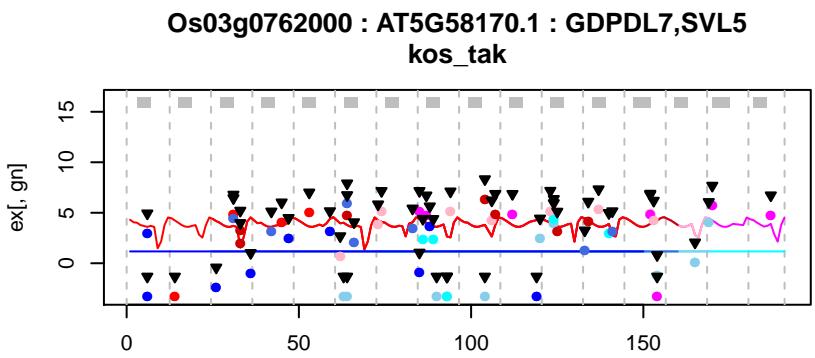


Os12g0491800 : AT1G79460.1 : ATKS,ATKS1,GA2,KS,KS1
2106.95223282039_Most significant each peak

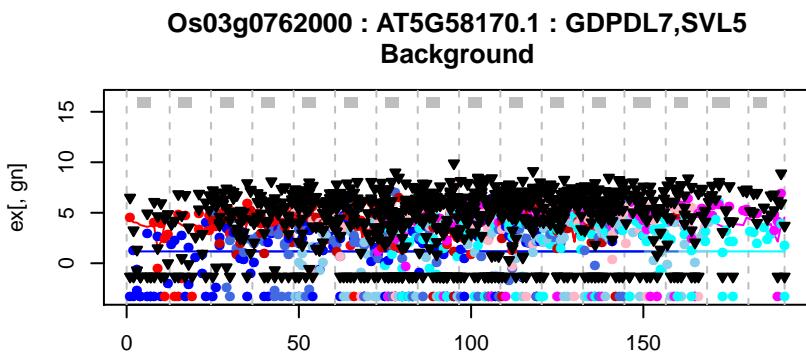
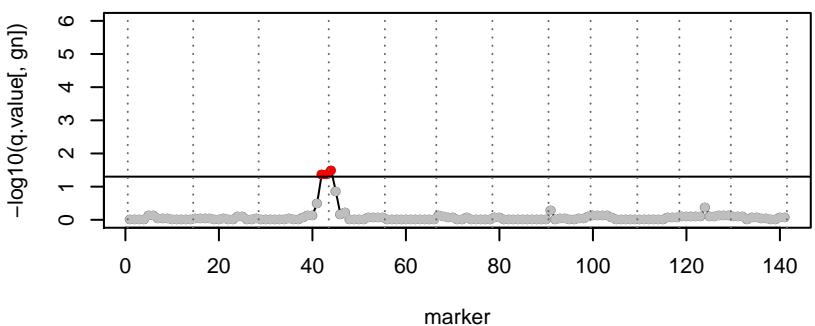
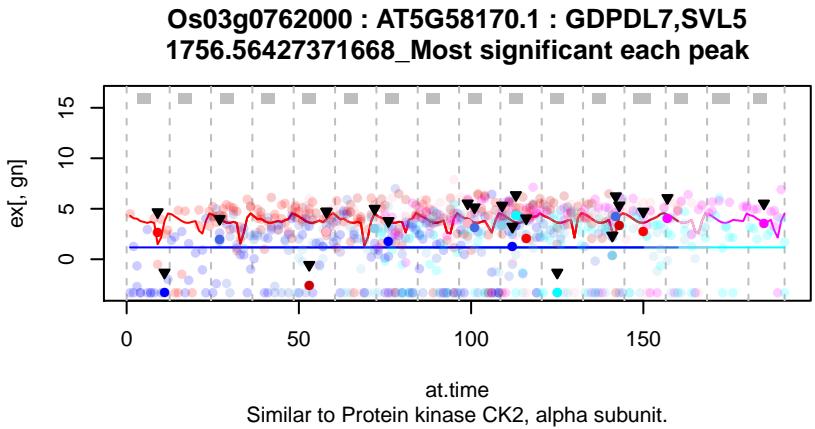


Os12g0491800 : AT1G79460.1 : ATKS,ATKS1,GA2,KS,KS1
2115.23998345504_Most significant peak

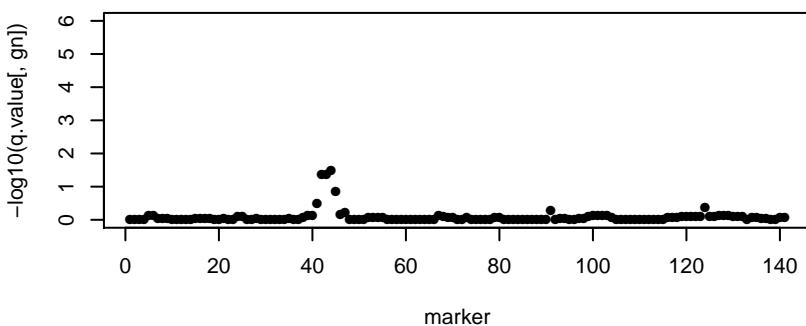
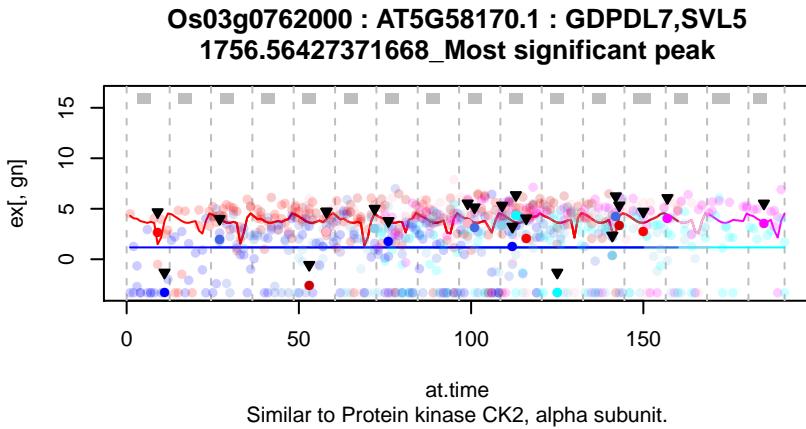




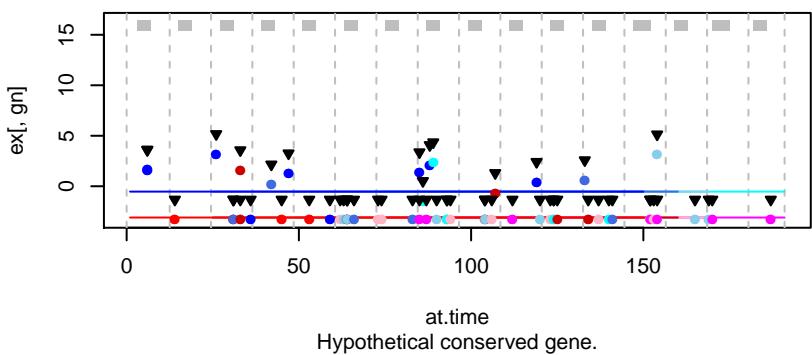
Similar to Protein kinase CK2, alpha subunit.



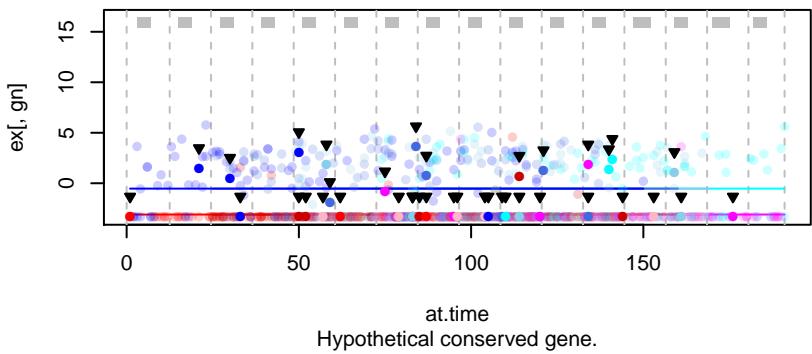
Similar to Protein kinase CK2, alpha subunit.



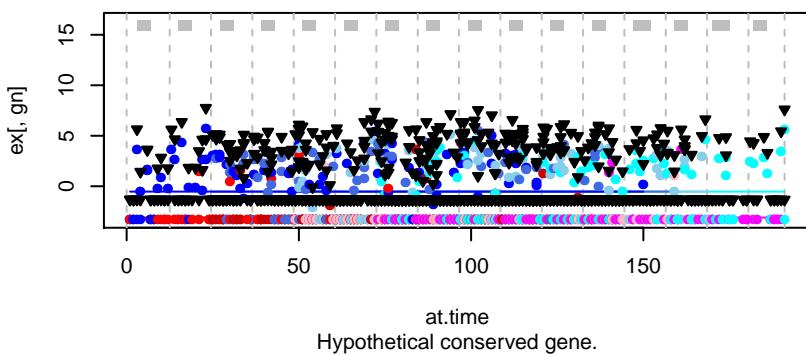
Os01g0258100 : :
kos_tak



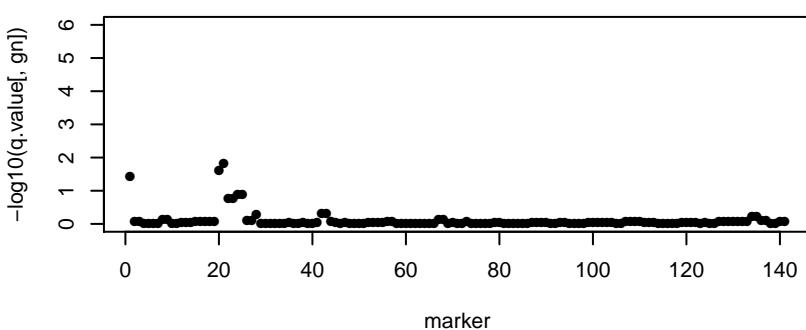
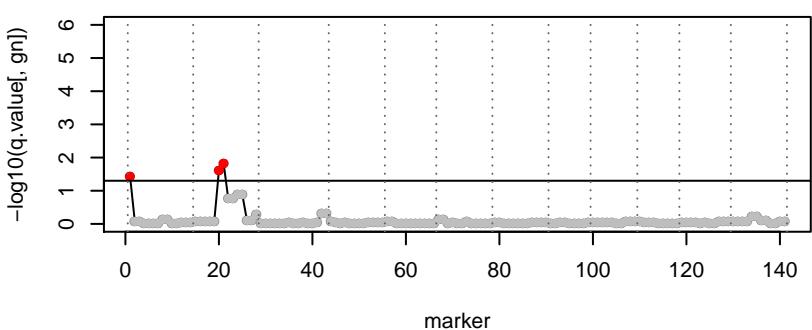
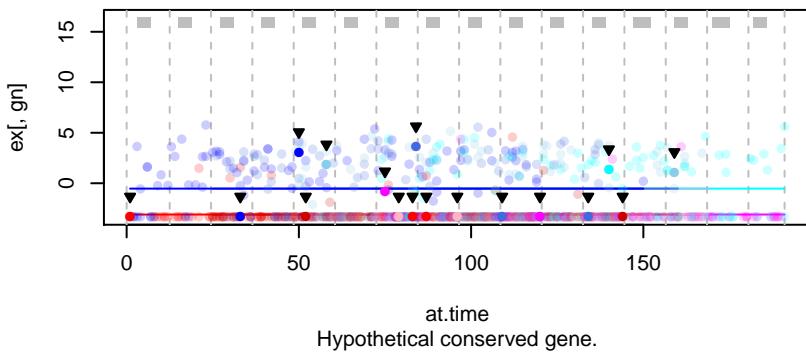
Os01g0258100 : :
1346.15535259798_Most significant each peak



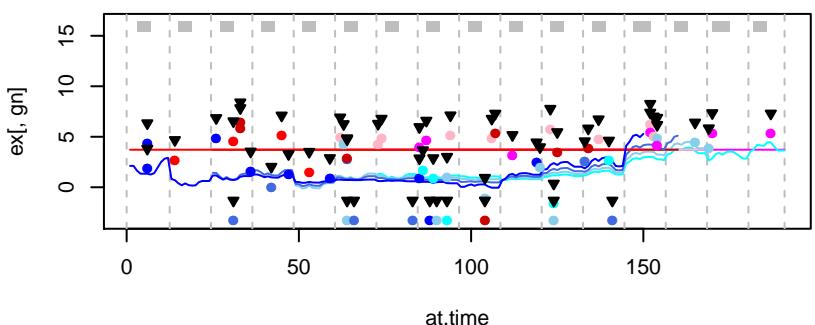
Os01g0258100 : :
Background



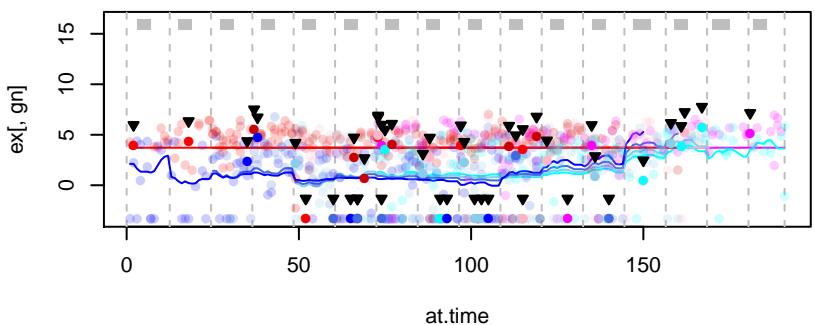
Os01g0258100 : :
1361.42718838425_Most significant peak



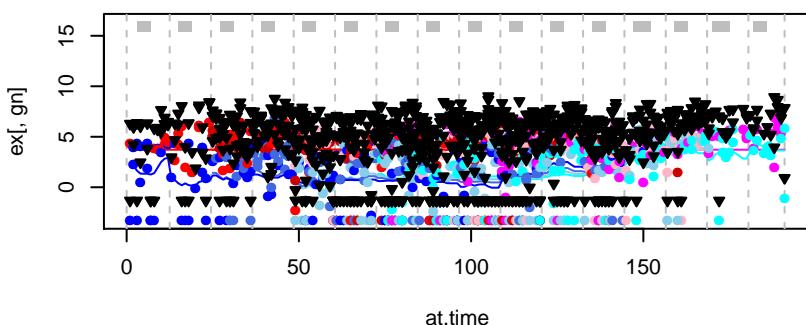
Os05g0568100 : AT4G22220.1 : ATISU1,ISU1
kos_tak



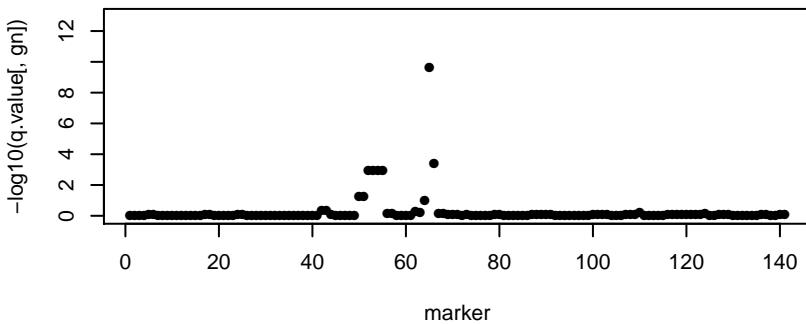
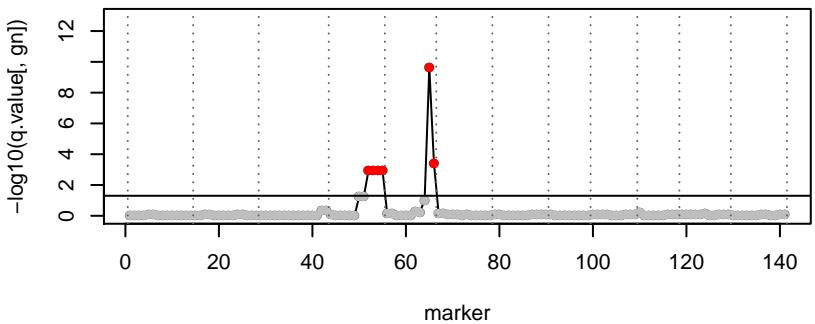
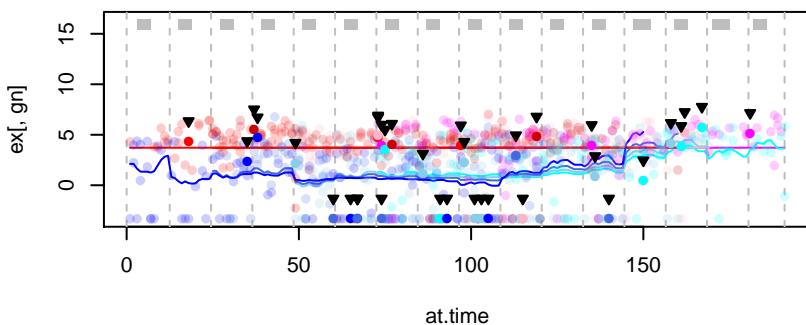
Os05g0568100 : AT4G22220.1 : ATISU1,ISU1
1738.62694127425_Most significant each peak



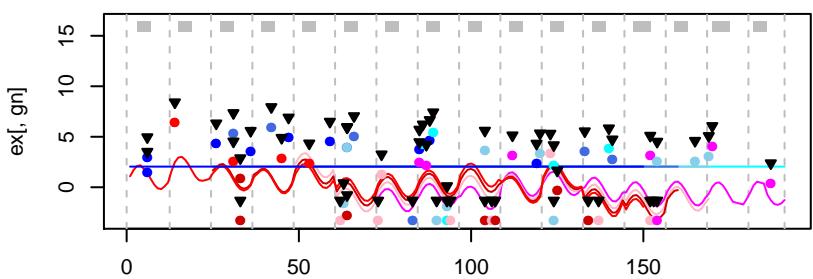
Os05g0568100 : AT4G22220.1 : ATISU1,ISU1
Background



Os05g0568100 : AT4G22220.1 : ATISU1,ISU1
1738.20169977804_Most significant peak

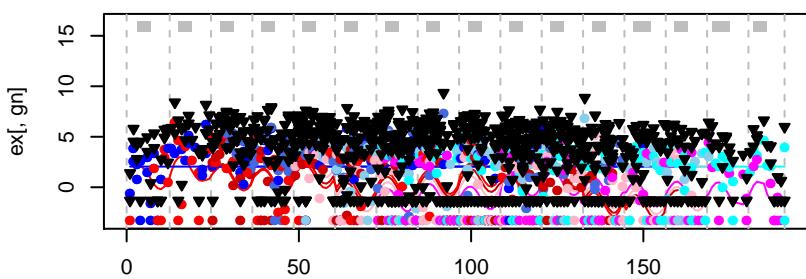


Os09g0519200 : AT4G38710.2 :
kos_tak



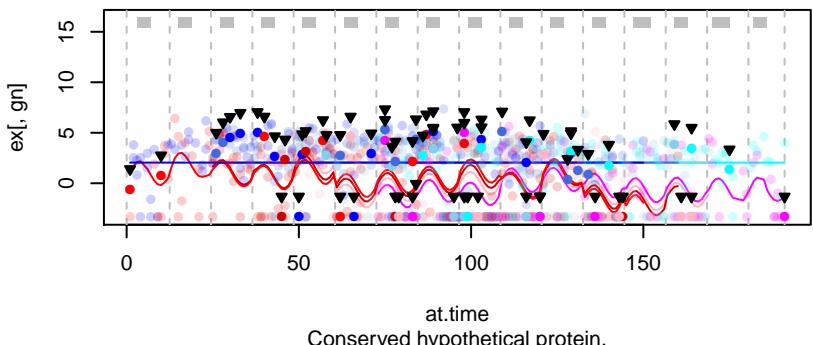
at.time
Conserved hypothetical protein.

Os09g0519200 : AT4G38710.2 :
Background



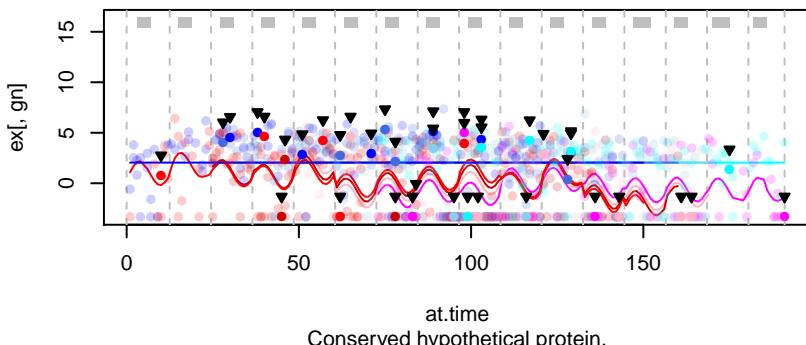
at.time
Conserved hypothetical protein.

Os09g0519200 : AT4G38710.2 :
2008.30434681739_Most significant each peak

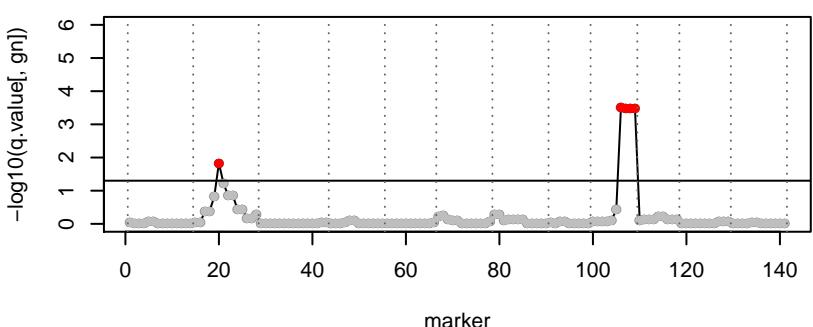


at.time
Conserved hypothetical protein.

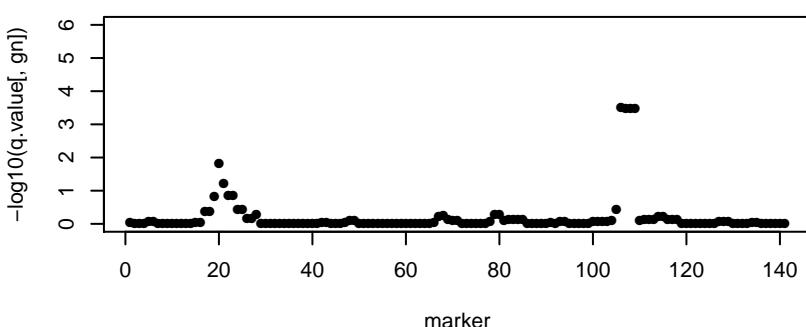
Os09g0519200 : AT4G38710.2 :
2032.09591671668_Most significant peak



at.time
Conserved hypothetical protein.

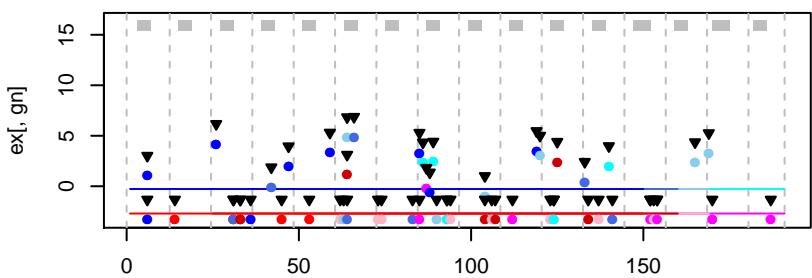


marker



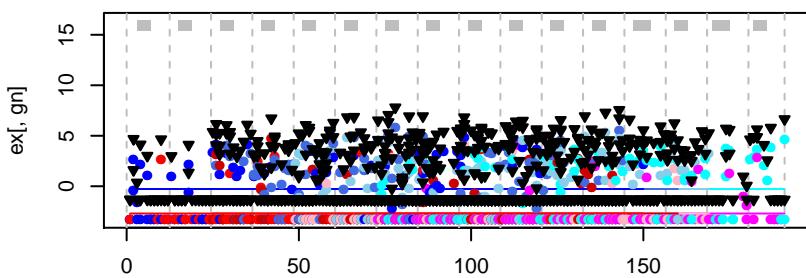
marker

Os04g0293700 : AT3G60310.1 :
kos_tak



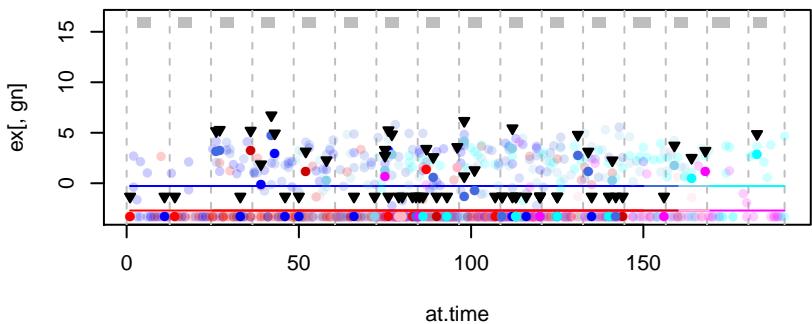
Transposase, Ptta/En/Spm, plant domain containing protein.

Os04g0293700 : AT3G60310.1 :
Background



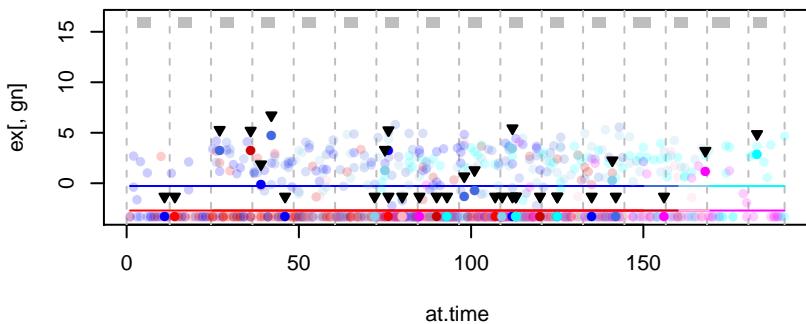
Transposase, Ptta/En/Spm, plant domain containing protein.

Os04g0293700 : AT3G60310.1 :
1623.62274479516_Most significant each peak

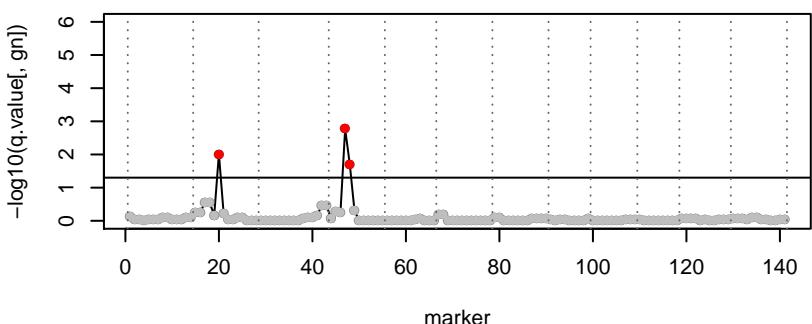


Transposase, Ptta/En/Spm, plant domain containing protein.

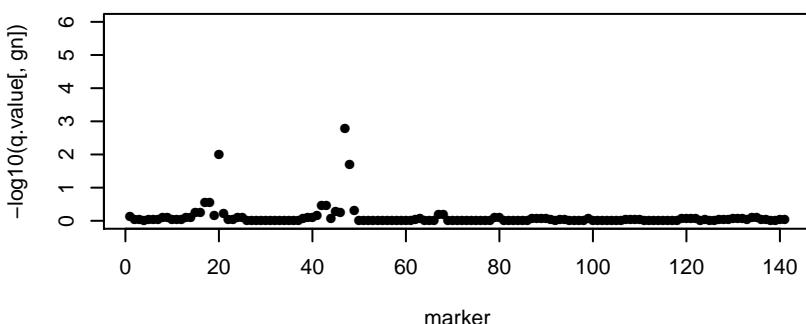
Os04g0293700 : AT3G60310.1 :
1628.48228864552_Most significant peak



Transposase, Ptta/En/Spm, plant domain containing protein.

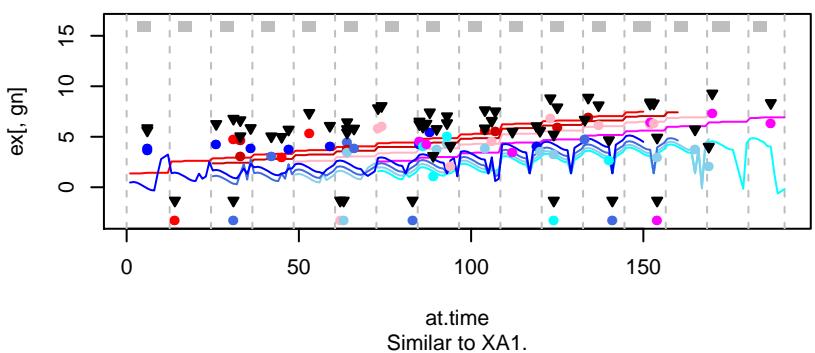


marker

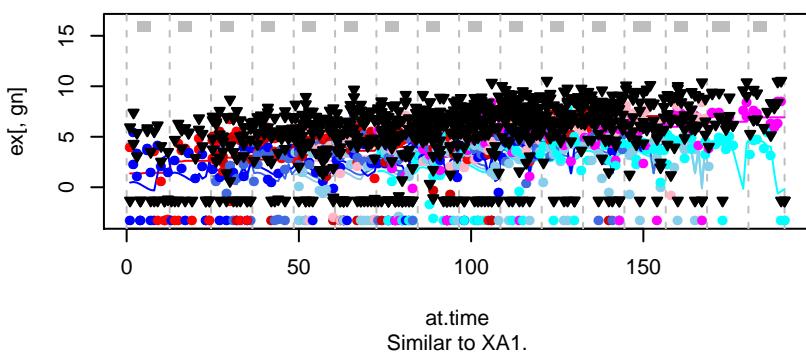


marker

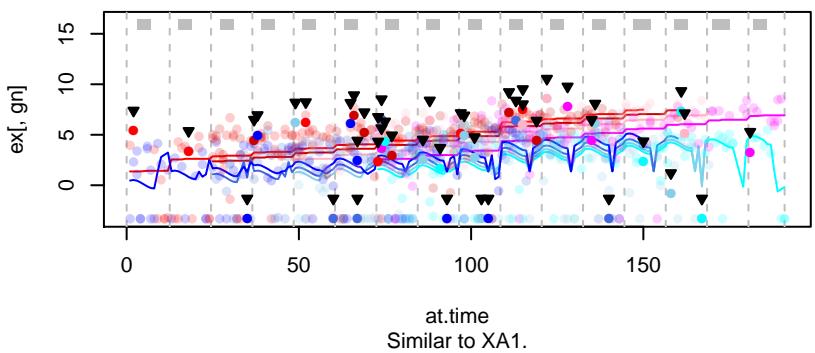
Os04g0623066 : AT1G69550.1 :
kos_tak



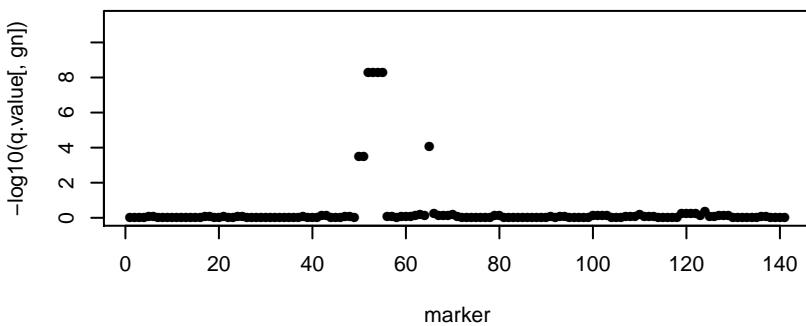
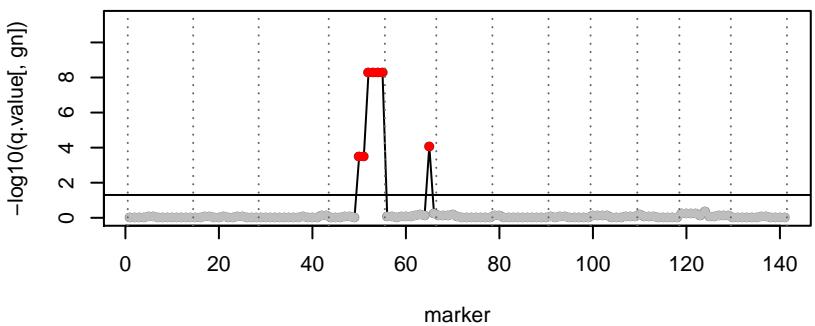
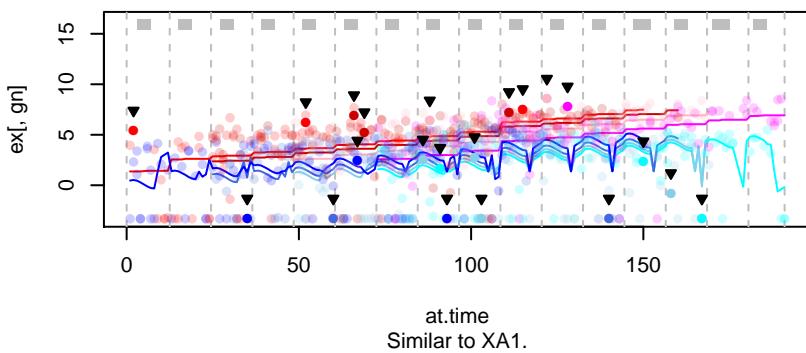
Os04g0623066 : AT1G69550.1 :
Background



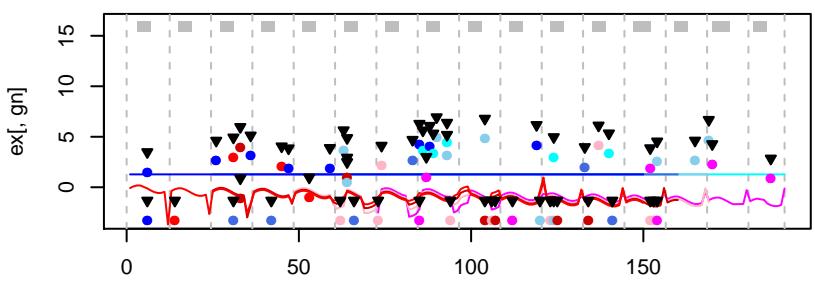
Os04g0623066 : AT1G69550.1 :
1625.46881036049_Most significant each peak



Os04g0623066 : AT1G69550.1 :
1616.10128548366_Most significant peak

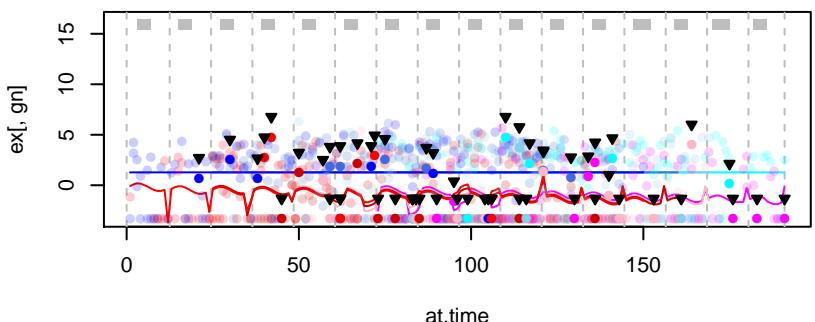


Os01g0175500 : AT1G56410.1 : ERD2,HSP70T-1
kos_tak



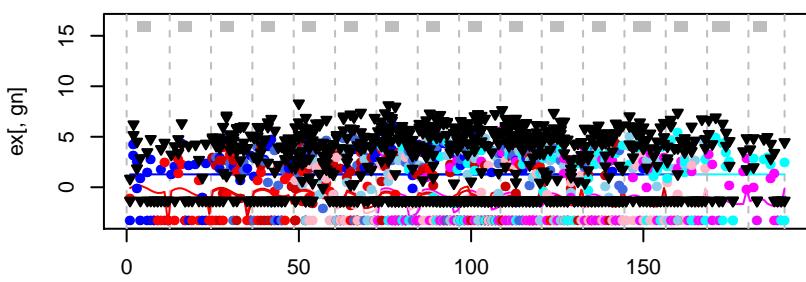
Protein of unknown function DUF1618 domain containing protein.

Os01g0175500 : AT1G56410.1 : ERD2,HSP70T-1
2122.89522479221_Most significant each peak



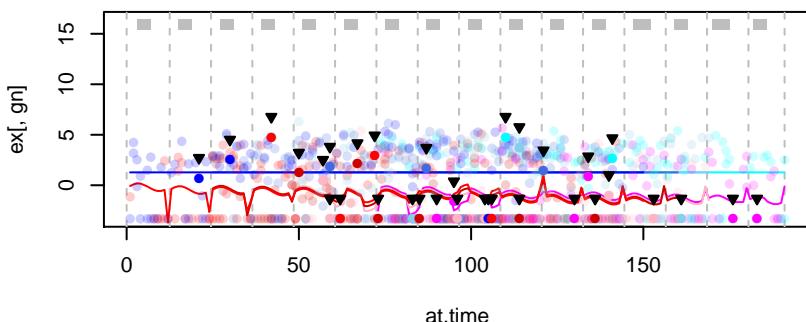
Protein of unknown function DUF1618 domain containing protein.

Os01g0175500 : AT1G56410.1 : ERD2,HSP70T-1
Background

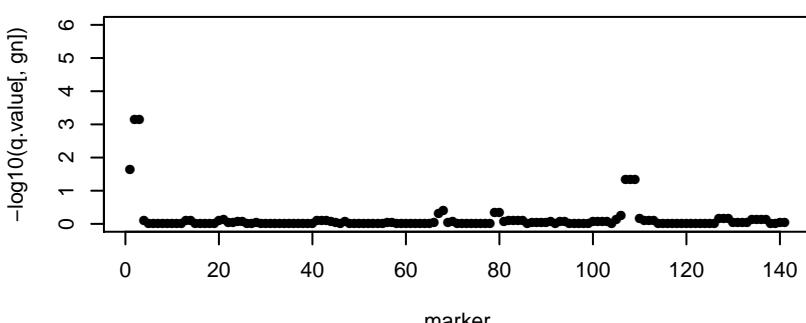
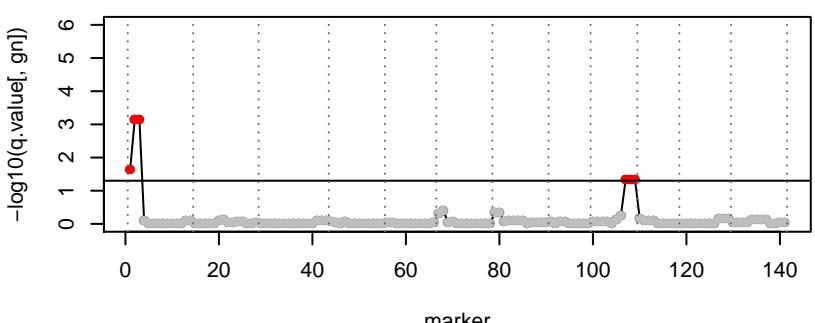


Protein of unknown function DUF1618 domain containing protein.

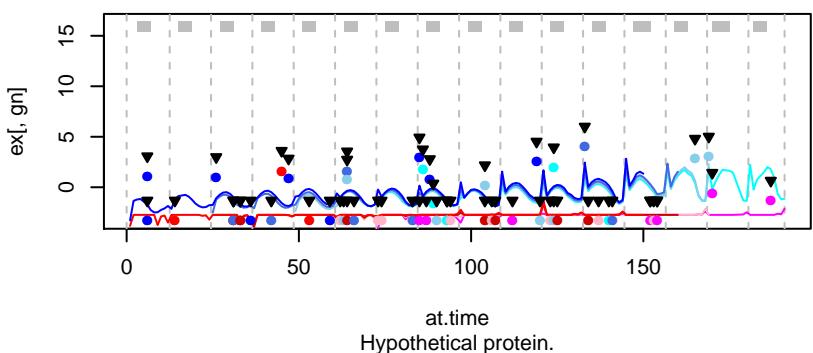
Os01g0175500 : AT1G56410.1 : ERD2,HSP70T-1
2140.88983743039_Most significant peak



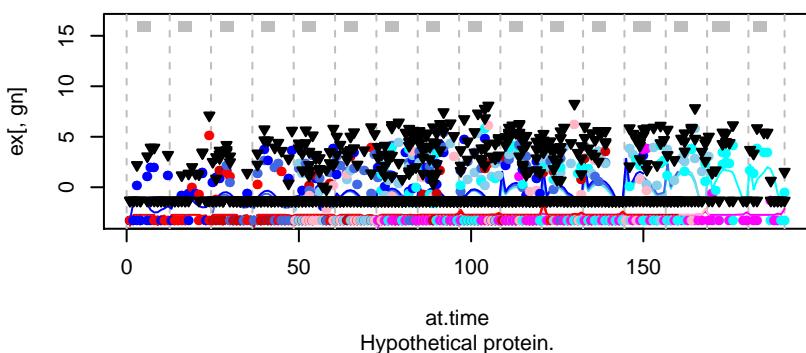
Protein of unknown function DUF1618 domain containing protein.



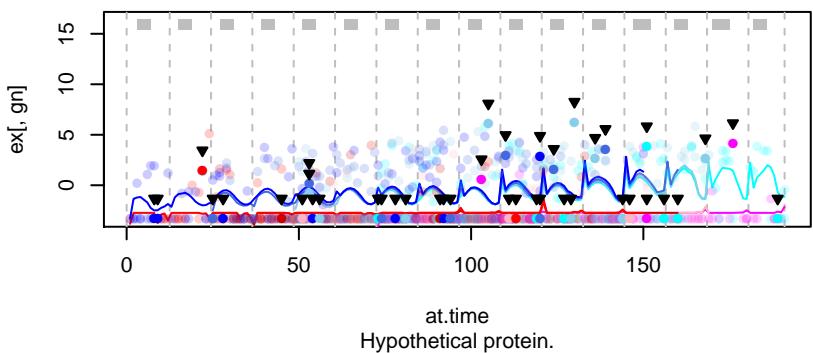
Os11g0685500 : :
kos_tak



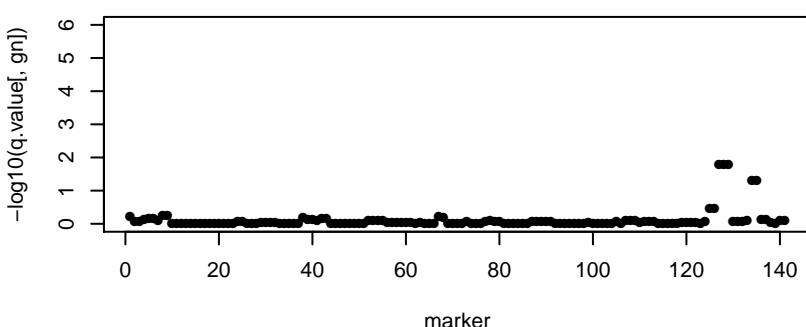
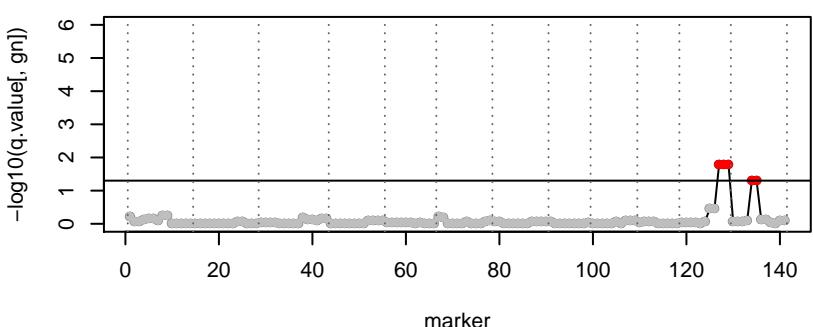
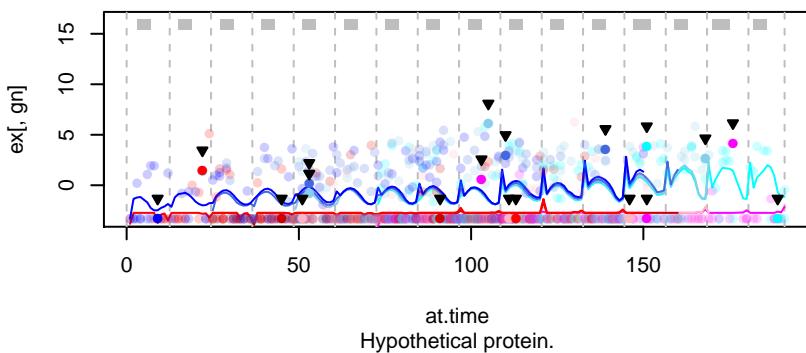
Os11g0685500 : :
Background



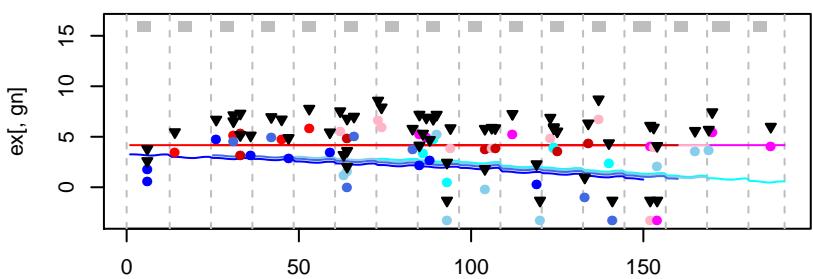
Os11g0685500 : :
1477.43806734952_Most significant each peak



Os11g0685500 : :
1459.6973042572_Most significant peak

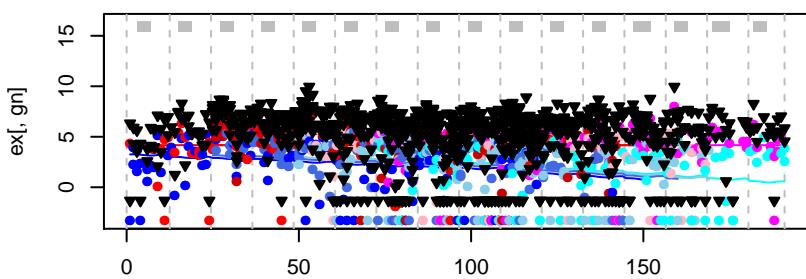


Os01g0298500 : AT4G25150.1 :
kos_tak



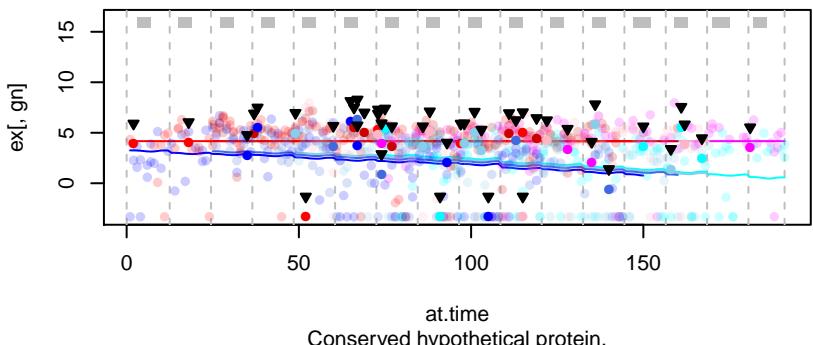
at.time
Conserved hypothetical protein.

Os01g0298500 : AT4G25150.1 :
Background



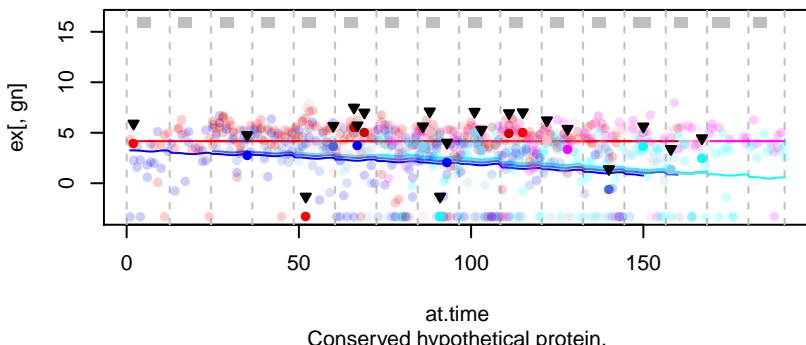
at.time
Conserved hypothetical protein.

Os01g0298500 : AT4G25150.1 :
1602.37848677405_Most significant each peak

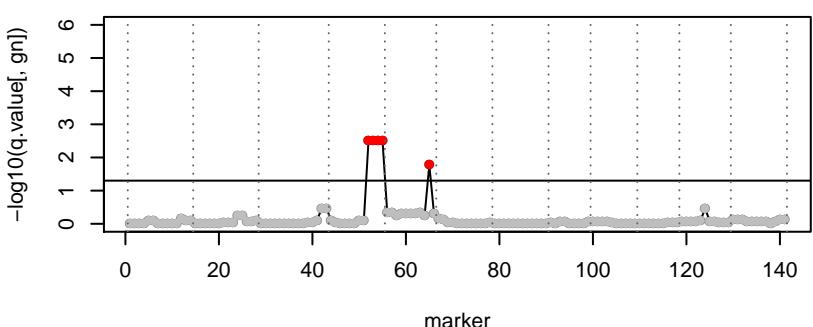


at.time
Conserved hypothetical protein.

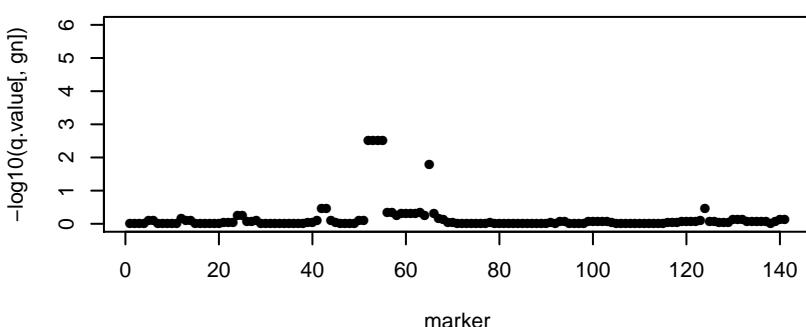
Os01g0298500 : AT4G25150.1 :
1602.6683495339_Most significant peak



at.time
Conserved hypothetical protein.

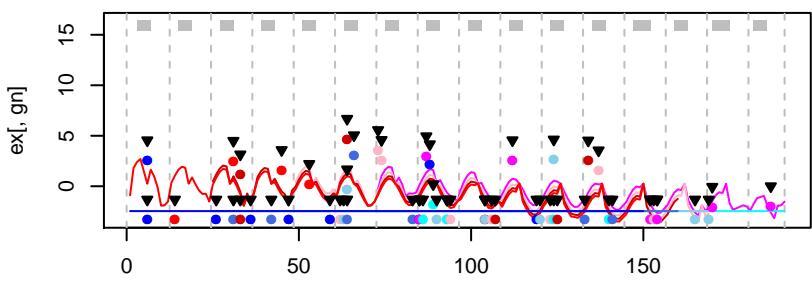


marker



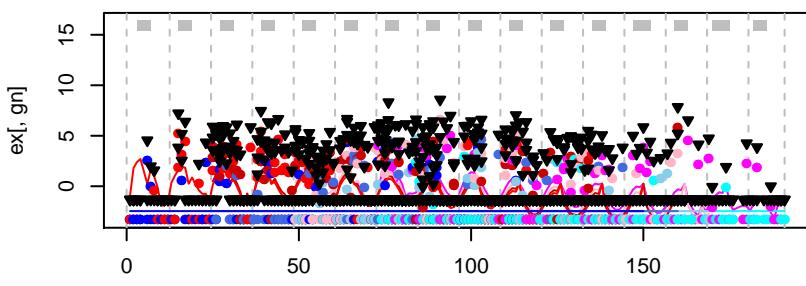
marker

Os01g0621600 : AT1G64300.1 :
kos_tak



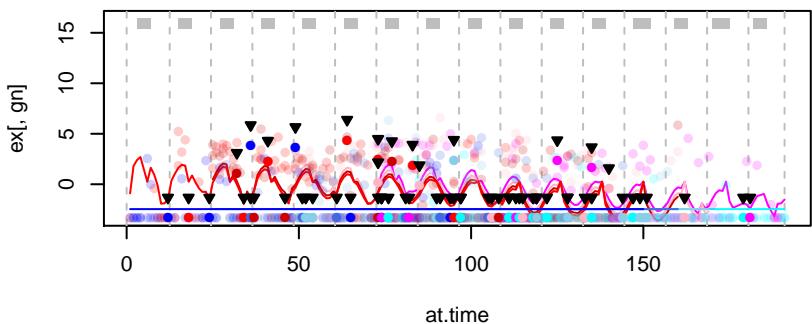
Serine/threonine protein kinase-related domain containing protein.

Os01g0621600 : AT1G64300.1 :
Background



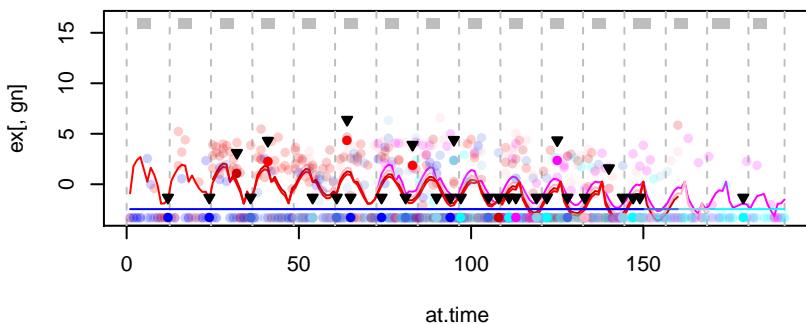
Serine/threonine protein kinase-related domain containing protein.

Os01g0621600 : AT1G64300.1 :
1557.94658747294_Most significant each peak

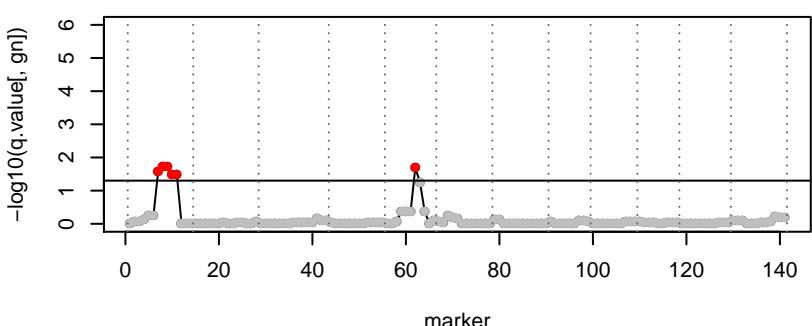


Serine/threonine protein kinase-related domain containing protein.

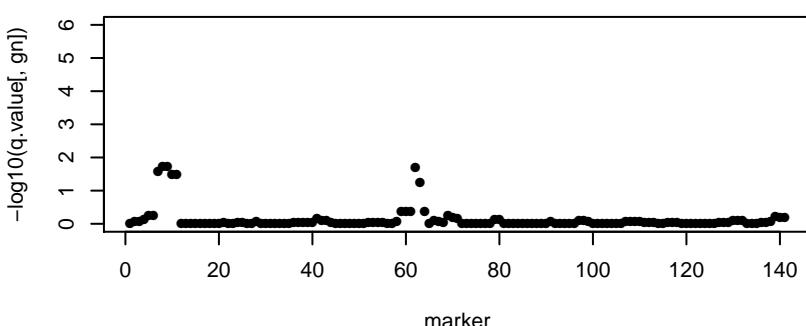
Os01g0621600 : AT1G64300.1 :
1572.98697424055_Most significant peak



Serine/threonine protein kinase-related domain containing protein.

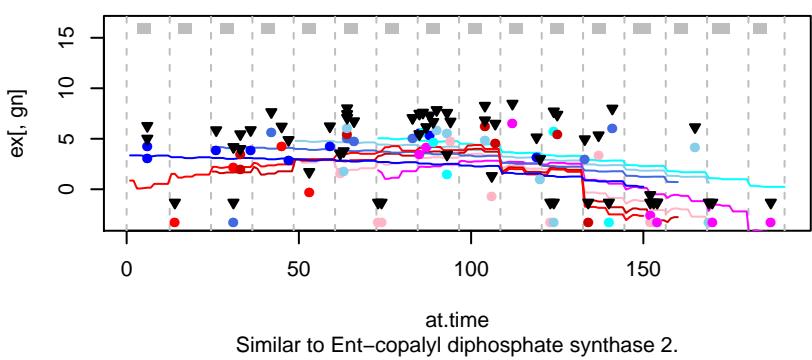


Serine/threonine protein kinase-related domain containing protein.

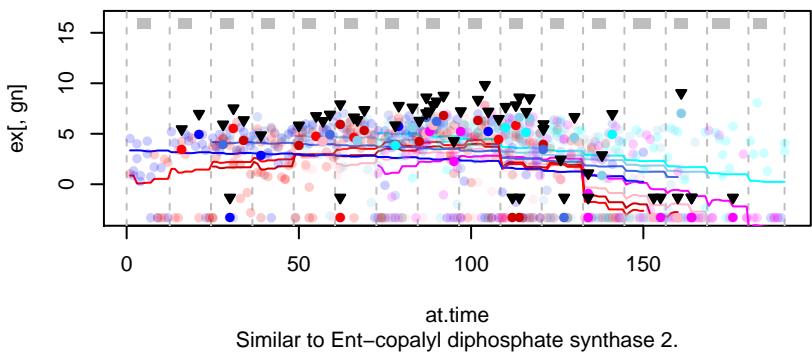


Serine/threonine protein kinase-related domain containing protein.

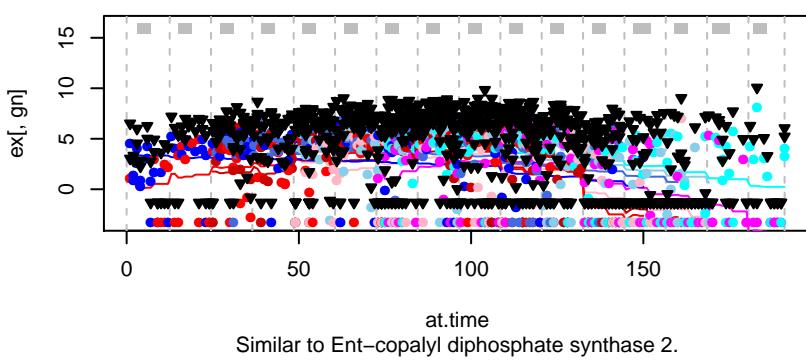
Os02g0570900 : AT4G02780.1 : ABC33,ATCPS1,CPS,CPS1,GA1
kos_tak



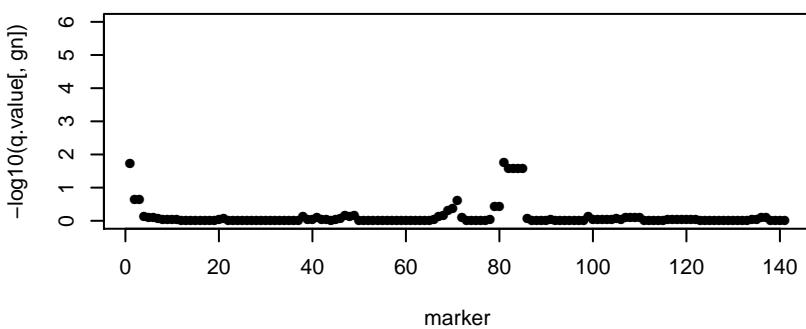
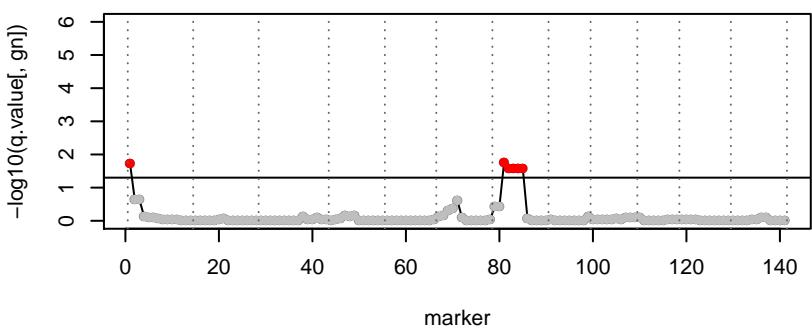
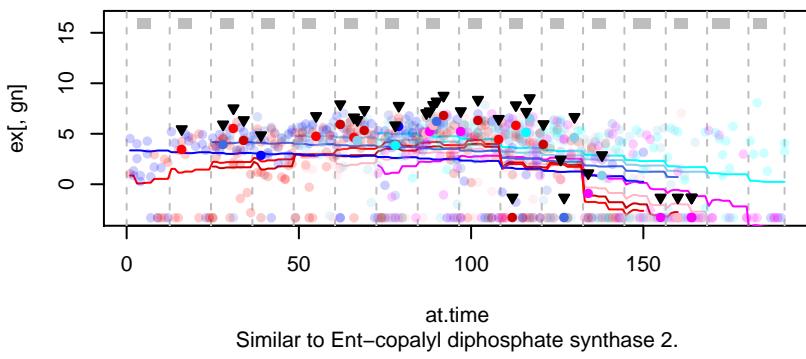
Os02g0570900 : AT4G02780.1 : ABC33,ATCPS1,CPS,CPS1,GA1
2065.55263307915_Most significant each peak



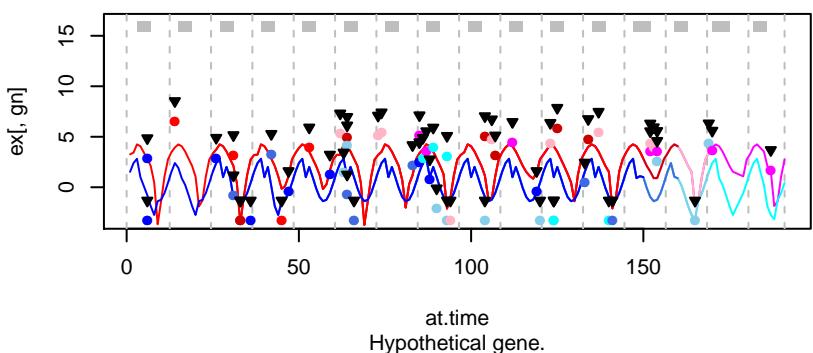
Os02g0570900 : AT4G02780.1 : ABC33,ATCPS1,CPS,CPS1,GA1
Background



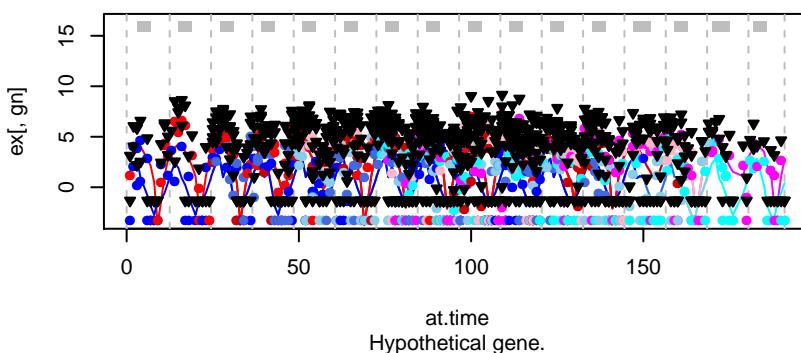
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2091.98870328852_Most significant peak



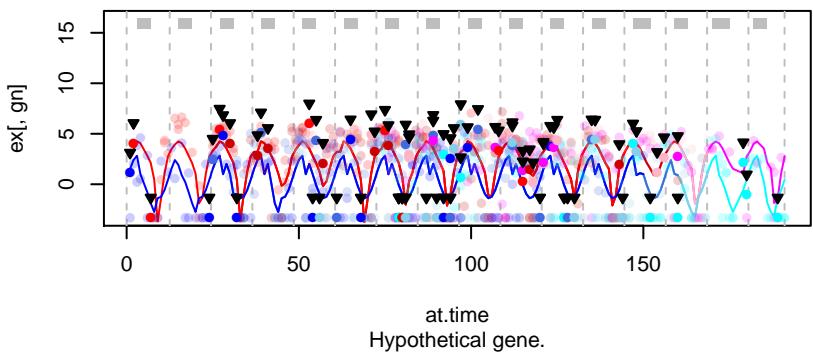
Os12g0527301 : AT1G25530.1 :
kos_tak



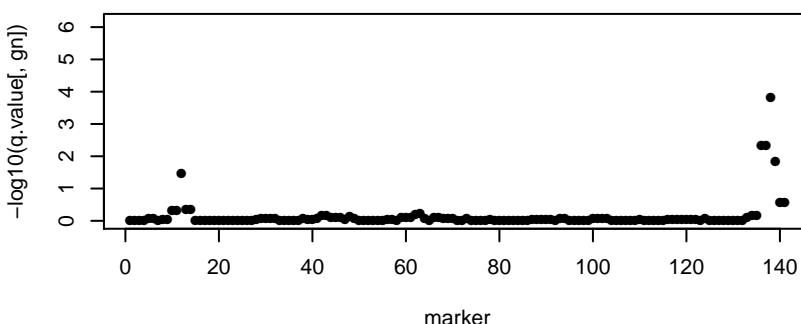
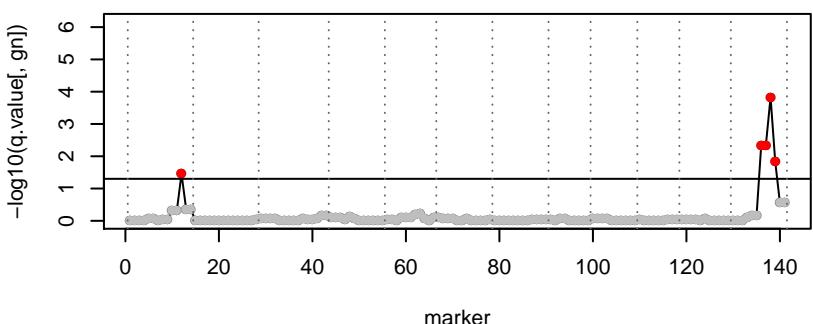
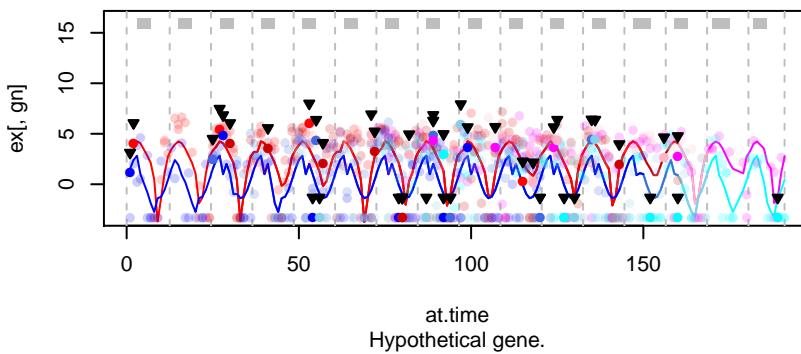
Os12g0527301 : AT1G25530.1 :
Background



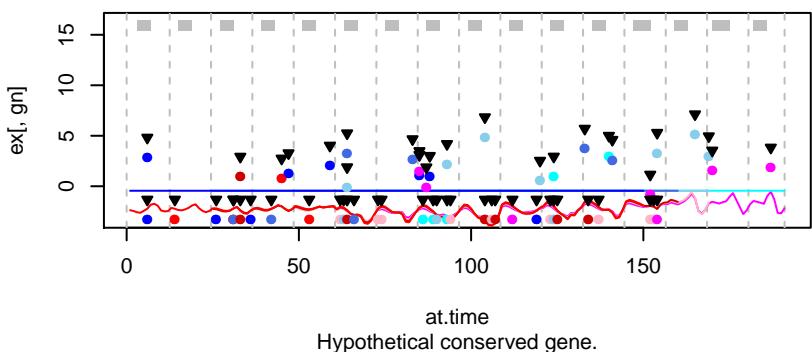
Os12g0527301 : AT1G25530.1 :
1874.24320775877_Most significant each peak



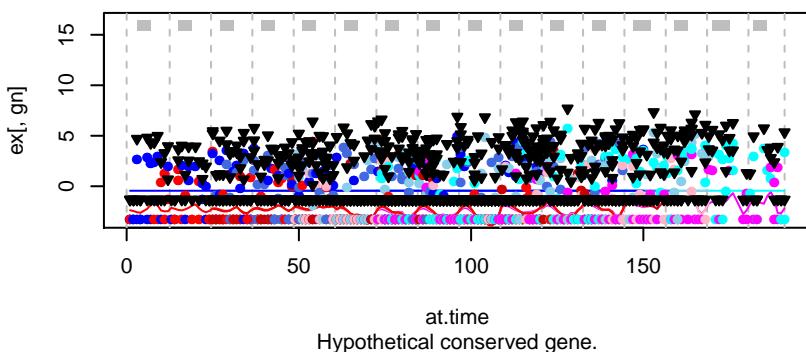
Os12g0527301 : AT1G25530.1 :
1884.85552000905_Most significant peak



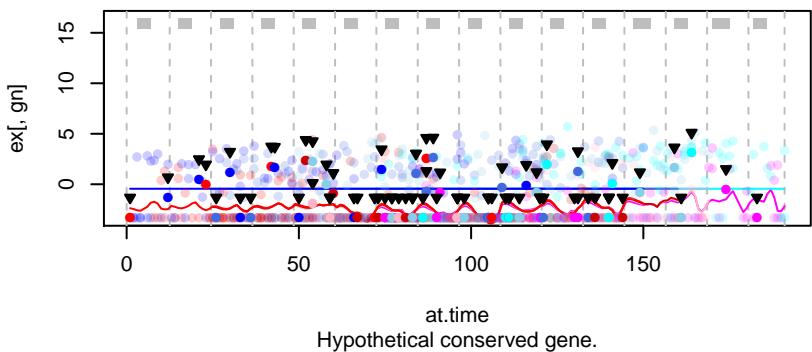
Os01g0298301 : AT2G25920.1 :
kos_tak



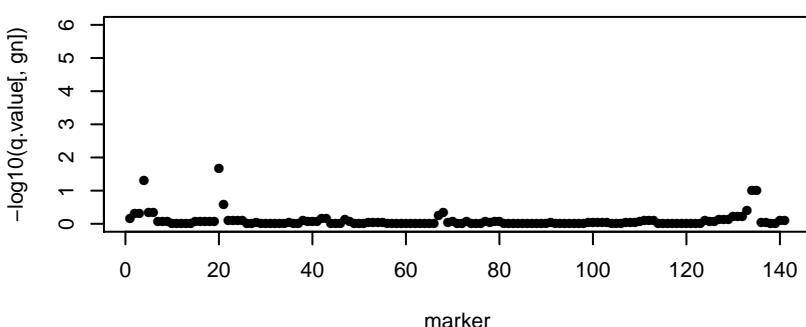
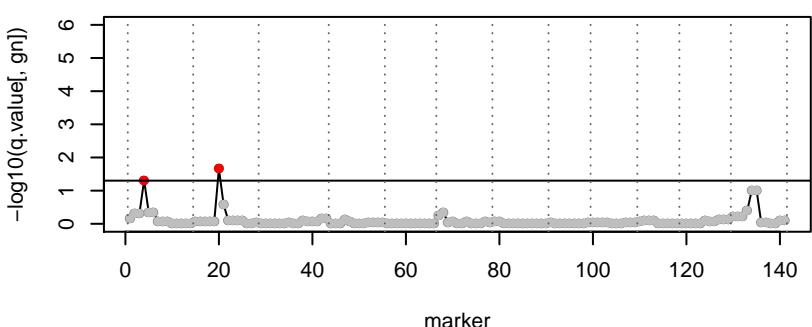
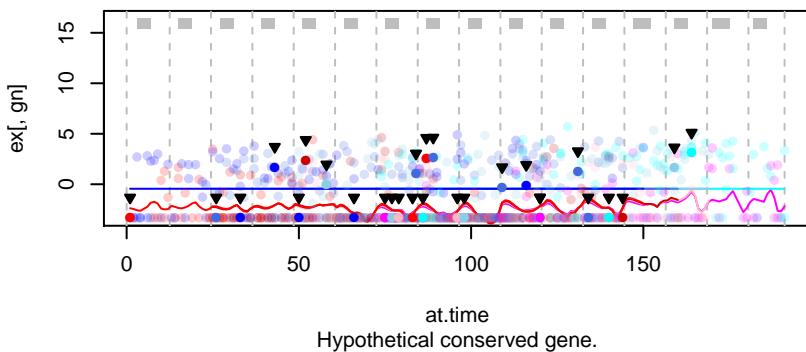
Os01g0298301 : AT2G25920.1 :
Background



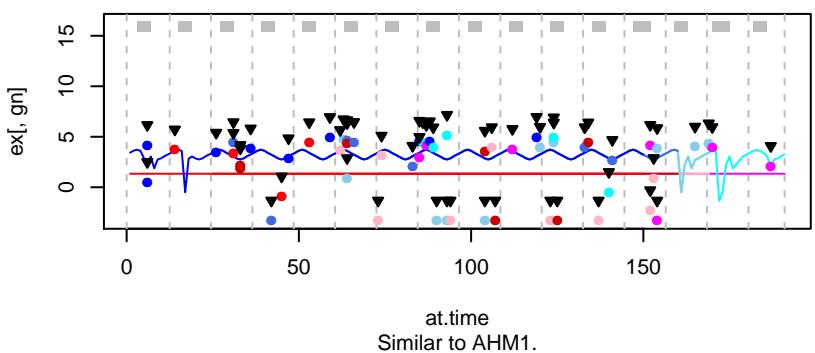
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1717.23204039277_Most significant each peak



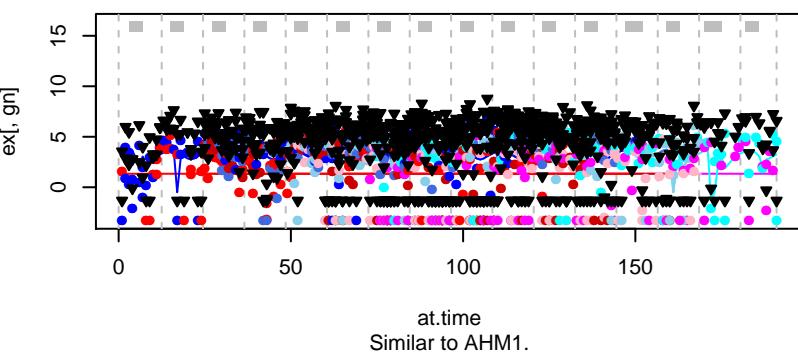
Os01g0298301 : AT2G25920.1 :
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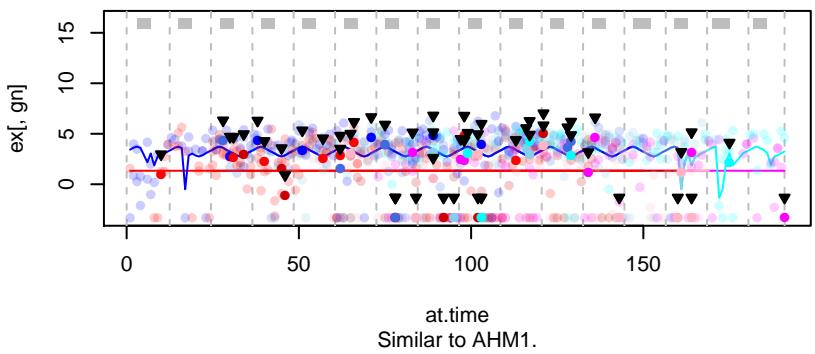
Os09g0463700 : AT5G64360.3 : EIP9
kos_tak



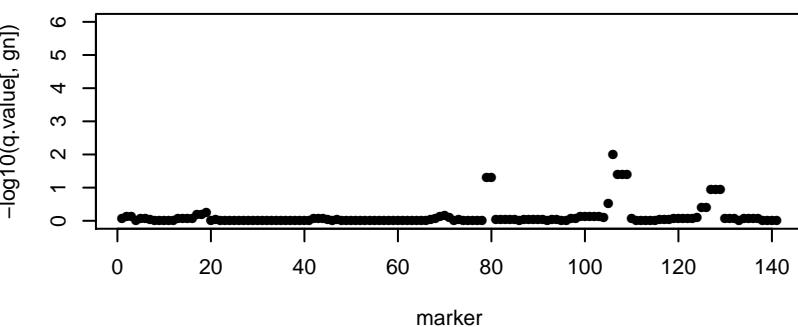
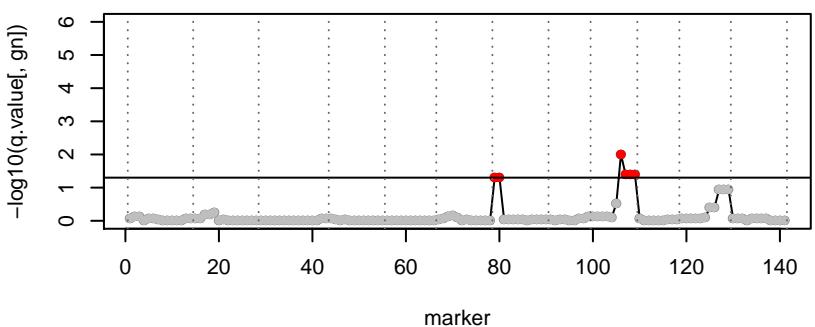
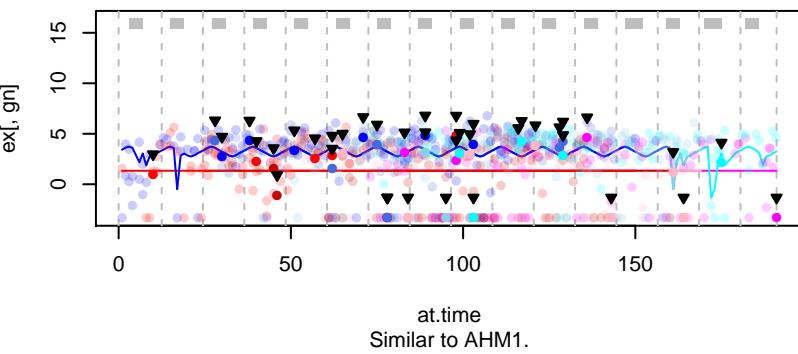
Os09g0463700 : AT5G64360.3 : EIP9
Background



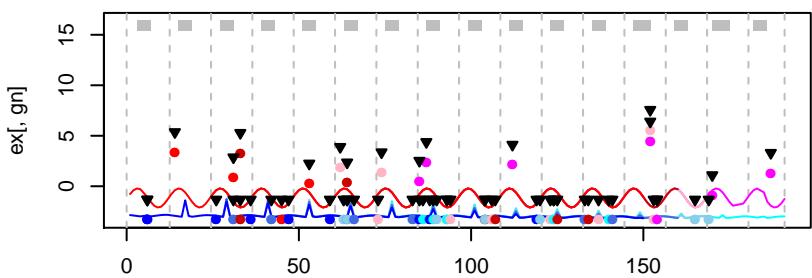
Os09g0463700 : AT5G64360.3 : EIP9
1789.35866386205_Most significant each peak



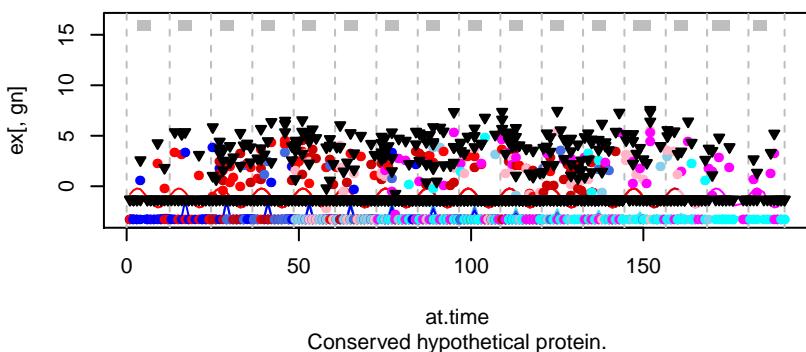
Os09g0463700 : AT5G64360.3 : EIP9
1793.49738616171_Most significant peak



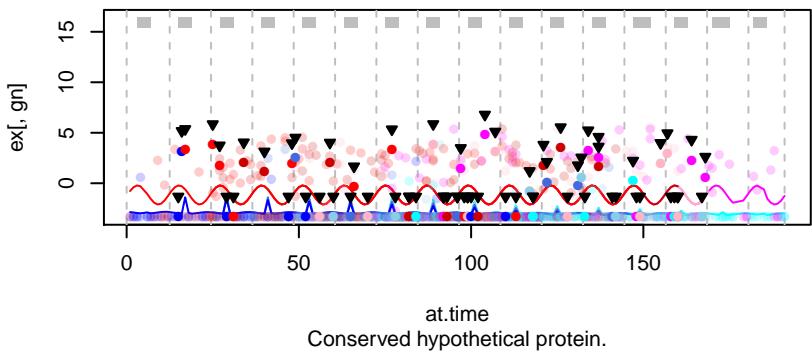
Os03g0267400 : AT5G19840.2 :
kos_tak



Os03g0267400 : AT5G19840.2 :
Background



Os03g0267400 : AT5G19840.2 :
1270.46168959638_Most significant each peak



Os03g0267400 : AT5G19840.2 :
1272.15739241147_Most significant peak

