

Supplementary Table 1 General information of SBP-box genes selected for phylogenetic analysis.

Gene	Accession no. or locus ID	Protein sequence of SBP-domain
AtSPL1	At2g47070	CQVENCEADLSKVVDYHRRHKVCEMHSKATSATVGGILQRFCCQCSRFLHLLQEFDEGKRSCRRRLAGHNKRRRK
AtSPL2	At5g43270	CQVEGCNLDLSSAKDYHRKHRICENHSKFPKVVVSGVERRFCQCSRFLHCLSEFDEKKRSCRRRLSDHNARRRK
AtSPL3	At2g33810	CQVESCTADMSKAKQYHKRHKVCQFHAKAPHVRISGLHQRFCQCSRFLHCLSEFDEAKRSCRRRLAGHNERRRK
AtSPL4	At1g53160	CQVDRCTADMKEAKLYHRRHKVCEVHAKASSVFLSGLNQRFCQCSRFLHDLQEFDEAKRSCRRRLAGHNERRRK
AtSPL5	At3g15270	CQVDRCTVNLTEAKQYYRRHRVCEVHAKASAATVAGVRQRFCQCSRFLHELPEFDEAKRSCRRRLAGHNERRRK
AtSPL6	At1g69170	CQVYGCSKDLSSSKDYHRKHRVCEAHSKTSVVIVNGLEQRFCQCSRFLHLLSEFDDGKRSCRRRLAGHNERRRK
AtSPL7	At5g18830	CQVPDCEADISELKGYHKRHRVCLRCATASFVLDGENKRYCQCGKFHLLPDFDEGKRSCRRKLERHNNRRKR
AtSPL8	At1g02065	CQAEGCNADLSHAKHYHRRHKVCEFHSAKASTVVAAGLSQRFCQCSRFLHLLSEFDNGKRSCRKRLADHNRRRK
AtSPL9	At2g42200	CQVEGCGMDLTNAKGYYSRHRVCGVHSTPKVTVAGIEQRFCQCSRFLHLLPEFDLEKRSCRRRLAGHNERRRK
AtSPL10	At1g27370	CQIDGCELDLSSSKDYHRKHRVCEHSHKCPKVVVSGLERRFCQCSRFLHAVSEFDEKKRSCRKRLSHHNARRRK
AtSPL11	At1g27360	CQIDGCELDLSSAKGYHRKHKVCEHSHKCPKVSVSGLERRFCQCSRFLHAVSEFDEKKRSCRKRLSHHNARRRK
AtSPL12	At3g60030	CQVDNCGADLSKVVDYHRRHKVCEIHSKATTALVGGIMQRFCQCSRFLHLLSEFDEGKRSCRRRLAGHNKRRRK
AtSPL13	At5g50570	CLVDGCDSDFSNCREYHKRHKVCDVHSTPKVVTINGHKQRFCQCSRFLHLLSEFDEGKRSCRKRLDGHNNRRRK
AtSPL14	At1g20980	CQVDNCTEDLSHAKDYHRRHKVCEVHSKATKALVGKQMQRFCQCSRFLHLLSEFDEGKRSCRRRLAGHNRRRK
AtSPL15	At3g57920	CQVEGCRMDLSNVKAYYSRHKVCCIIHSKSSKVVIVSGLHQRFCQCSRFLHLLSEFDLEKRSCRRRLACHNERRRK
AtSPL16	At1g76580	CQVDNCKEDLSIAKDYHRRHKVCEVHSKATKALVGKQMQRFCQCSRFLHLLSEFDEGKRSCRRRLDGHNNRRRK

MdSBP1	MDP0000861601 or MDP0000156994	CQVERCGADLVDAKRYRRHKVCEFHSKAAVVIVSGIQRFCQQCSRFHELIEFDEAKRSCRRRLAGHNERRRK
MdSBP2	MDP0000149339	CQVDNCKEDLSNAKDYHRRHKVCEVHSKSTRALVAKMQRFCCQCSRFHPLSEFDEGKRSCRRRLAGHNRRRRK
MdSBP3	MDP0000263766	CLVDGCNSDLSTCRDYHRRHKVCELHSKTPQVTINGNKQRFCQQCSRFHAPPEFDEGKRSCRKRLDGHNRRRRK
MdSBP4	MDP0000146640 or MDP0000778465	CQVYGCNKDLSSCKDYHRRHKVCEVHSKTAKVIINGIEQRFCQQCSRFHLLGEFDDGKRSCRKRLAGHNERRRK
MdSBP5	MDP0000803116	CQAEGCAADLSHSKHYHRRHKVCEFHSKASTVIANGLTQRFCQQCSRFHLLSEFDNGKRSCRKRLADHNRRRRK
MdSBP6	MDP0000262141	CQVEGCHVALLNAKEYHRRHKVCAMHSKASRVTVLGQDQRFCQQCSRFHVSEFDESKRSCRRRLAGHNERRRK
MdSBP7	MDP0000170630	CLVDGCRADLSRCREYHRRHRVCELHSKAPVVVVVRGEQKRFCQQCSRFHSLVDFDKVKRSCRKRLNGHNQRRRK
MdSBP8	MDP0000865739	CQADRCTADLSEEKQYHRKHKVCDLHKSQVVLVSGLHQRFCQQCSRFHQLPEFDDTKRSCRRRLAGHNERRRK
MdSBP9	MDP0000158607	CQAERCGADLVDAKRYHRRHKVCEFHSKAAVVIVSGTRQRFCQQCSRFHELIEFDEAKRSCRRRLAGHNERRRK
MdSBP10	MDP0000171877	CQVHGCNMDLTFSKDYHRRHRVCDAHSKTAVVIVNGIKQRFCQQCSRFHLLAEFDDVKRSCRRRLAGHNLRRRK
MdSBP11	MDP0000246046	CQVHGCNMDLTFSKDYHRRHRVCDAHSKTAVVIVNGIKQRFCQQCSRFHLLAEFDDVKRSCRRRLAGHNLRRRK
MdSBP12	MDP0000155354	CQVEGCNLDLSSVKDYHRKHRICANHSKSPKVVVDGVERRFCQQCSRFHGLSEFDENKRSCRRRLSDHNARRRK
MdSBP13	MDP0000919693	CQVEDCKADLSNAKDYHRRHKVCDMHSKATKAVVGNVLQRFCQQCSRFHGLQEFDEGKRSCRRRLAGHNRRRRK
MdSBP14	MDP0000919694	CQVEDCKADLSNAKDYHRRHKVCAMHSKATKALVGSVMQRFCQQCSRFHALQEFDEGKRSCRRRLAGHNRRRRK
MdSBP15	MDP0000180408	CQVEDCKADLSNAKDYHRRHKVCDMHSKATKALVGNVMQRFCQQCSRFHALQEFDEGKRSCRRRLAGHNRRRRK
MdSBP16	MDP0000180409	CQVEDCKADLSNAKDYHRRHKVCDMHSKATKAPVGNVLQRFCQQCSRFHVLQEFDEGRRSCRRRLAGHNRRRRK
MdSBP17	MDP0000271587	CQVPSCGIDIKELKGYHRRHRVCLACANAITVVIDGETKRYCQCQGFHVLPDFDEGKRSCRKLERHNNRRRR

MdSBP18	MDP0000297978	CQVEGCQVDLSAKAYYSRHKVCGLSKTPTVIVAGLEQRFCCQCSRFLHLLPEFDQGKRSCRRRLAGHNERRRK
MdSBP19	MDP0000210138	CLVDGCNCDLSTCRDYHRRHKVCELHSTPQVTINGNKQRFCCQCSRFLHLLPEFDQGKRSCRKRLDGHNRERRRK
MdSBP20	MDP0000589558	CQVYGCNKDLSSYKDYHRRHKVCEVHSKTAKVIVNGIEQRFCCQCSRFLHLLGEFDDGKRSCRRRLAGHNERRRK
MdSBP21	MDP0000193702	CQVHGCNMDLTFSKDYHRRHVCDAHSKTAVVIVNGIEQRFCCQCSRFLHLLAEFDDVKRSCRKRLAGHNLRRRK
MdSBP22	MDP0000119458	CQVHGCNMDLTFSKDYHRRHVCDAHSKTAVVIVNGIEQRFCCQCSRFLHLLAEFDDVKRSCRKRLAGHNLRRRK
MdSBP23	MDP0000249364	CQVEGCNLDLSSAKDYHRKHRCNSHSPKVVVDGVELRFCQCSRFLHGLSDFDENKRSCRKRLSDHNARRRK
MdSBP24	MDP0000176265	CLVDGCNSDLKCRDYHRRHKVCELHSTKSVTIKQERRFCQCSRFLHLLVEFDEGKRSCRRRLDGHNKRRRK
MdSBP25	MDP0000195461	CQVPSCGVDIKELKGYHRRHVRCLACANASTVDIDGESKRYCQCGKFHVLSDFDEGKRSCRRKLERHNNRRRK
MdSBP26	MDP0000142582	CLVDDCRADLSRCREYHRRHVRCELHSTPQVVVKGQRFCCQCSRVLHLLVEFDDGKKSCRKRLNGHNQRRRK
MdSBP27	MDP0000255162 or MDP0000538106	CQADRCTADLSDEKKYHRKHVCDLHSSQVVLVSLGHQRFCCQCSRFLHLLSEFDDTKRSCRSRLSGHNERRRK
OsSPL1	LOC_Os01g18850.1	CQVDGCTVNLSSARDYNKRHKVCEVHTKSGVVRIKNVEHRFCQCSRFLHLLQEFDEGKKSCRSRLAQHNRRRK
OsSPL2	LOC_Os01g69830.1	CSVEGCAADLSKCRDYHRRHKVCEAHSKTAVVTVAGQQRFCCQCSRFLHLLGEFDEEKRSCRKRLDGHNKRRRK
OsSPL3	LOC_Os02g04680.1	CQVEGCNVDLSSAKPYHRKHVRCEPHSKTLKVVIVAGLERRFCQCSRFLHLLAEFDQKKRSCRRRLHDHNARRRK
OsSPL4	LOC_Os02g07780.1	CQVEGCGVELVGKDYHRKHVRCEAHSKFPRVVVAGQERRFCQCSRFLHLLSEFDDQKKRSCRRRLYDHNARRRK
OsSPL5	LOC_Os02g08070.1	CQAEGCKADLSAAKHYHRRHKVCDFAHAKAAVLAAGKQQRFCQCSRFLHLLAEFDEAKRSCRKRLTEHNRRRK
OsSPL6	LOC_Os03g61760.1	CQVEGCTADLTGVRDYHRRHKVCEMHAKATTAVVGNTVQRFCCQCSRFLHLLQEFDEGKRSCRRRLAGHNRRRK
OsSPL7	LOC_Os04g46580.1	CQVEGCDITLQGVKEYHRRHKVCEVHAKAPRVVVGTEQRFCCQCSRFLHLLAEFDDAKKSCRRRLAGHNERRRK
OsSPL8	LOC_Os04g56170.1	CQAEGCKADLSSAKRYHRRHKVCEHHSKAPVVVVTAGLHQRFCCQCSRFLHLLDEFDDAKKSCRKRLADHNRRRK

OsSPL9	LOC_Os05g33810.1	CQVPGCEADIRELKG YHRRHRVCLRCAHAAAVMLDGVQKRYCQQCGKFHILLDFDEDKRSCRRKLERHNRRRRK
OsSPL10	LOC_Os06g44860.1	CQAE GCKADLSGAKHYHRRHKVCEYHAKASVVAASGKQQRFCQQCSR FHVLT E FDEAKRSCRKRLAEHNRRRRK
OsSPL11	LOC_Os06g45310.1	CQVEGCGLELGGYKEYYRKHRVCEPHTKCLR VVVAGQDRRFCQQCSR FHAPSEFDQEK RSCRRRLSDHNARRK
OsSPL12	LOC_Os06g49010.1	CQVEGCKVDLSSAREYHRKHKVCEAHSKAPKVIVSGLERRFCQQCSR FHGLAEFDQKKKSCRRRLSDHNARRK
OsSPL13	LOC_Os07g32170.1	CQVERCGVDLSEAGRYNRRHKVCQTHSKEPVVLVAGLRQRF CQQCSR FH E L T E F D D A K R S C R R R L A G H N E R R R K
OsSPL14	LOC_Os08g39890.1	CQVEGCGADLSGIKNYYCRHKVCFMHSKAPRVV VAGLEQRF CQQCSR FH L L P E F D Q G K R S C R R R L A G H N E R R R K
OsSPL15	LOC_Os08g40260.1	CQVDDCRADLTNAKD YHRRHKVCEIHGKT T K A L V G N Q M Q R F C Q Q C S R F H P L S E F D E G K R S C R R R L A G H N R R R R K
OsSPL16	LOC_Os08g41940.1	CAVDGCKEDLSKCRDYHRRHKVCEAHSKTPLVVVSGREMRF CQQCSR FH L L Q E F D E A K R S C R K R L D G H N R R R R K
OsSPL17	LOC_Os09g31438.1	CQVEGCGVDLSGVKPY YCRHKVCYMHAK EPIVV VAGLEQRF CQQCSR FH Q L P E F D Q E K K S C R R R L A G H N E R R R K
OsSPL18	LOC_Os09g32944.1	CAVDGCKADLSKHRDYHRRHKVCEPHSKTPVVVVSGREMRF CQQCSR FH L L G E F D E A K R S C R K R L D G H N R R R R K
OsSPL19	LOC_Os11g30370.1	CSVDGCRSDLSRCRDYHRRHKVCEAHAKTPVVVVAGQEQRFCQQCSR FHNLAEFDDGKKSCRKRLDGHNRRRRK
CNR	Solyc02g077920	CQVDQCTADMADAKPYHRRHKVCEFHSKSPIVLISGLQKRFCQQCSR FHLLAEFDDAKRSCRRRLAGHNERRK
SlySBP2	Solyc04g045560	CQVDSCNLDLSSAKQYHRKHRVCHHSKCPKV IILGRHRRFCQQCSR FHSLSDFDENKRSCRT RLSDHNARRK
SlySBP3	Solyc10g009080	CQVEECTADMVNAKTYHRRHKVCEFHAKAPEVLIDGLRQRF CQQCSR FH Q L A E F D D A K K S C R R R L A G H N E R R R K
SlySBP4	Solyc07g053810	CQAEKCNVDLSDAKQYHKRHKVCEYHAKSQVVVVAGLRQRF CQQCSR FH E L T E F D E S K R S C R R R L A G H N E R R R K
SlySBP6a	Solyc03g114850	CQVQCGCKDLSPCKDYHKRHKVCEVHSKTAKVIVNGIEQRFCQQCSR FHLLAEFDDGKRSCRKRLAGHNERRK
SlySBP6b	Solyc05g012040	CQVHGCNKDLSSSKDYHKRHKVCNEHSKTAIVIVNGIEQRFCQQCSR FHLLAEFDEGKRSCRKRLAGHNERRK
SlySBP6c	Solyc12g038520	CQVQCGCKDLTSC KDYHKRHKVCEIHSKTAKVIVNGIQRF CQQCSR FH L L A E F D D G K R S C R K R L A G H N E R R R K

SlySBP7	Solyc01g080670	CQVPDCEADISELKGYHKRHRVCLRCANATSVVLDGHSKRYCQQCGKFHILSDFDEGKRSCRRKLERHNNRRRR
SlySBP8a	Solyc10g018780	CQAEGCNADLTHAKHYHRRHKVCEFHSKASTVIAAGITQRFCCQCSRFHLLSEFDNGKRSCRKRLADHNRRRRK
SlySBP8b	Solyc01g090730	CQAEGCNADLTHAKHYHRRHKVCEFHSKASTVIAAGLTQRFCCQCSRFHVLSEFDNGKRSCRKRLADHNRRRRK
SlySBP10	Solyc05g015510	CQVEGCNLDLSSAKEYYRKHRVCDSHSKSPKVIVAGVARRFCQCSRFHSVSEFDDKKRSCRRRLSDHNARRRK
SlySBP12a	Solyc01g068100	CQVDDCGTDLKAKDYHRRHKVCEMHSKASRALVGNVMQRFCCQCSRFHALQEFDEGKRSCRRRLAGHNKRRRK
SlySBP12b	Solyc05g053240	CQVQDCRADLSSAKDYHRRHKVCEVHСКАAKALVGNVMQRFCCQCSRFHVLEEFDEGKRSCRRRLAGHNKRRRK
SlySBP13	Solyc05g015840	CLVDGCNADLSECREYHRRHKVCEVHSKTAKVTIAGRDQRFCCQCSRFHSLVEFDDGKRSCRKRLDGHNNRRRK
SlySBP15	Solyc10g078700	CQVEGCQADLSDAKAYYSRHKVCGMHSKSPTVVVAGLEQRFCCQCSRFHQLTEFDQGKRSCRRRLACHNERRRK
VvSBP1	XM_002273498.1	CQVHGCNMDLSSSKDYHKRHKVCDVHSKTPKVIVNGIEQRFCCQCSRFHLLAEFDDGKRSCRKRLAGHNERRRK
VvSBP2	XM_002271276.1	CQVEGCNLDLSTAKDYHRKHRVCESHTKCPKVIVGGLERRFCQCSRFHSLSEFDEKKRSCRRRLSDHNARRRK
VvSBP3	XM_002270190	CLVDGCKSDLKCRDYGRRHKVCELHSKTAKVTIGGHEQRFCCQCSRFHSLGEFDEGKRSCRKRLDGHNNRRRK
VvSBP4	NW_002239818.1	CQADDCGVDLRAAKRYHRRHKVCERHAKAAFVFLGGIEQRFCCQCSRFHEISQFDDTKRSCRKRLAGHNQRRRK
VvSBP5	XM_002274898	CQVEDCRADLGNAKDYHRRHKVCDMHSKASKALVGNVMQRFCCQCSRFHLLQEFDEGKRSCRRRLAGHNRRRRK
VvSBP6	XM_002277003.1	CQVTGCEADISELKGYHRRHRVCLRCANASVVILDGQNKRYCQQCGKFHILSDFDEGKRSCRRKLERHNNRRRR
VvSBP7	XM_002273192.1	CQVEDCGADLSKAKDYHRRHKVCEMHSKAGCALVGNDMQRFCCQCSRFHVLQEFDEGKRSCRRRLAGHNKRRRK
VvSBP8	XM_002278476.1	CQVEGCKVDLSDAKAYYSRHKVCGMHSKSPTVIVAGLEQRFCCQCSRFHQLAEFDQGKRSCRRRLAGHNERRRK
VvSBP9	XM_002280016.1	CQVDNCTADMSEAKRYHKRHRVCEHHAKAPVILIAGIQRFCCQCSRFHELSEFDDTKRSCRRRLAGHNERRRK
VvSBP10	XM_002267188.1	CQVEGCNLDLKSADYHRRHRICENHSKSPKVIVAGLERRFCQCSRFHELTEFDDKKRSCRRRLNDHNARRRR

VvSBP11	XM_002275692.1	CQAERCTADLTDKQYHRRHKVCEHHAKAQVVVVGGIRQRFQCQCSRFHELSEFDEAKRSCRRRLAGHNERRRK
VvSBP12	XM_002274324.1	CLVDGCTSDLRNCREYHRRHRVCERHSTPVIIGGQEKRFQCQCSRFHSLGEFDEVKRSCRKRLDGHNRERRRK
VvSBP13	XM_002274466.1	CQVEGCHVALVNAKDYHRRHKVCEMHSKAPKVVVLGLEQRFCQQCSRFHAVSEFDDSKRSCRRRLAGHNERRRK
VvSBP14	XM_002277147.1	CQAEGCNADLTHAKHYHRRHKVCEFHSTKASTVFAAGLTQRFQCQCSRFHLLSEFDNGKRSCRKRLADHNERRRK
VvSBP15	XM_002280124.1	CLVDGCNSDLSNCREYHRRHKVCELHSTAKVTIGGHTQRFQCQCSRFHSLLEEFDEGKRSCRKRLDGHNRERRRK
VvSBP16	XM_002265167.1	CQVHGCNKDLSSSKDYHRRHKVCEVHSTAKVIVNGIEQRFCQQCSRFHLLAEFDDGKRSCRKRLAGHNERRRK
VvSBP17	XM_002273748.1	CQVDNCREDLSNAKDYHRRHKVCEMHSKSTKALVGKQMQRFCQQCSRFHPLSEFDEGKRSCRRRLAGHNERRRK
VvSBP18	XM_002282562.1	CQAEKCGADLTDKRYHRRHKVCEVHAKAAMVEVAGLRQRFQCQCSRFHELSEFDEAKRSCRRRLAGHNERRRK
ZjSBP01	CM003114.1	CQADNCNADLTDKHYHRRHKVCEFHAKAPVVLVAGHQRFQCQCSRFHGLLEFDENKRSCRRRLAGHNERRRK
ZjSBP02	CM003117.1	CQVDNCKEDLSNAKDYHRRHKVCELHSTKALVAKQMQRFCQQCSRFHPLTEFDEGKRSCRRRLAGHNERRRK
ZjSBP03	CM003117.1	CQAERCADLSADKRYHRRHKVCEFHFKAPVIVAGLPQRFQCQCSRFHELSEFDEAKRSCRRRLAGHNERRRK
ZjSBP04	CM003118.1	CLVDGCKADLSTCRDYHRRHKVCELHSTPQVTIGGNTQRFQCQCSRFHSLLEEFDEGKRSCRKRLDGHNRERRRK
ZjSBP05	CM003118.1	CQVYGCNKDLSSSKDYHRRHKVCELHSTAKVIVNGIEQRFCQQCSRFHLLAEFDNGKRSCRKRLAGHNERRRK
ZjSBP06	CM003118.1	CQVQGCNLDLKLAKDYHRRHRICEIHSKSPKVVIVAGMERRFCQQCSRFHELSEFDGKKRSCRRRLSDHNARRRK
ZjSBP07	CM003120.1	CQVPTCGADIRELKGYHRRHRVCLRCANASTVVLDGETKRYCQCGKFHLLSDFDEGKRSCRRKLERHNNRRRK
ZjSBP08	CM003121.1	CQADECGLDLKLAKPYHRRHKVCEHAKAAVVLVNGLRQRFQCQCSRFHEISEFDGNKKSCRERLAGHNLRERRK
ZjSBP09	CM003122.1	CQVEDCGADLSNAKDYHRRHKVCEMHSKASRALVGNVMQRFCQQCSRFHVLQEFDEGKRSCRRRLAGHNKRRRK
ZjSBP10	CM003123.1	CQVHGCNMDLSSSKDYHRRHKVCDVHSTAKVIVNGIEQRFCQQCSRFHLLAEFDDGKRSCRRRLAGHNERRRK

ZjSBP11	CM003123.1	CQVEGCNLDLSSAKDYHRKHRICESHSKSPKVIVGGLERRFCQQCSRFHPLSEFDEKKRSCRRRLSDHNARRRK
ZjSBP12	CM003123.1	CLVDGCISDLSKCRDYYRRHKVCELHSKTPKVTIGGHEQRFCQQCSRFHSLEEFDEGKRSCRKRLDGHNRRRRK
ZjSBP13	CM003125.1	CQVEGCKVDLSDAKAYYCRHKVCGMHSTPKVIVAGLEQRFCQQCSRFHQLPEFDQGKRSCRRRLAGHNERRRK
ZjSBP14	CM003125.1	CQVEDCRADLSNAKDYHRRHKVCDMHKASKALVGNVMQRFCQQCSRFHVLKEFDEGKRSCRRRLAGHNRRRRK
ZjSBP15	KN813464.1	CQAEKCTADLTDAKQYHRRHKVCEIHAKAQVVVVGGLRQRFCQQCSRFHELSEFDETKRSCRRRLAGHNERRRK
ZjSBP16	KN815747.1	CLVDDCKADLSSCREYHKRHRVCERHKTPTVMVKGEEKRFCQQCSRFHALGEFDEVKRSCRKRLDGHNRRRRK

Supplementary Table 2 Primer sequences of ZjSBP in quantitative real-time PCR.

Gene	Forward primer (5'-3')	Reverse primer (5'-3')
<i>ZjSBP01</i>	GAAGAGGAGGAGGATGAAG	TGGAACCTGCTACATTGT
<i>ZjSBP02</i>	GATGTTGTTGATGCCTTGA	AACCTTGTGAACCTGGAA
<i>ZjSBP03</i>	GTGCTGATCTGAGTGATG	AGTCTGTGTAGTCTGGATT
<i>ZjSBP04</i>	ACCATAGACGCCATAAGG	GAGGAATATACAAGTGAAGTAGAA
<i>ZjSBP05</i>	GTGAGAACAGTAGTCAATATCC	TGCCGTAACAAGTATGGT
<i>ZjSBP06</i>	TAGGAGACAGCAGAGGAA	CATTATCGGTTGACAGAAGAG
<i>ZjSBP07</i>	GATTGAGCAGAGACTTGATG	ACTACTTGACTTCGTGTTATTC
<i>ZjSBP08</i>	AAGAAGAAGAAGAAGAAGAAG	ACCATTGACCAAGACTACG
<i>ZjSBP09</i>	CTTA ACTCCACTCCACATTG	CACTGCTGCTGATGATTC
<i>ZjSBP10</i>	ATATACTTCCAGGTGCCATT	AGGTTCCAGATACAGATGATT
<i>ZjSBP11</i>	AGAGATGTAAGTCTTCTTGTC	AGAGCCTTGTTGGATTCA
<i>ZjSBP12</i>	ATAGCCATTCTCAGTTAGGT	CCAATCTCCTTAGTATTATCAGTT
<i>ZjSBP13</i>	TGAGAGTCCTCCATCTGA	TCTTGTGAACTGCTGCTA
<i>ZjSBP14</i>	ATAGTCGCAGAGCAAGAA	CACCACAGAACACCAATC
<i>ZjSBP15</i>	GAGAGGAATAGGCAGAGAAT	CCACAACCACAACCTTGAG
<i>ZjSBP16</i>	AACCTCAATCTTGGCATCT	GCTTCTCTTCACCTCATCA
<i>Actin</i>	GAGGAAGCAACTGGCAACTAAGG	TACGAGCAAGCTGGATATCCTTC