

Appendix B. The information of 263 SSR primer pairs used in this study

Primer Name	Forward primer sequence (5' - 3')	Reverse primer sequence (5' - 3')	Repeat Motif
LBSSR0001	TTTCTATGCTGGATAGGATTCCTAG	ACCCAAATACTGCCTTCTTAACC	(AGA)17
LBSSR0002	AACATAATTATCCGAACAACCTCTCAG	GCCTTAAAAGTAGGAATGGTGCT	(AAT)22
LBSSR0003	TGACGCAATATGTGTAATCGG	GTTGCAGTCTAAAAAAGTAGGTTCC	(GAA)27
LBSSR0004	TATCTTCCATTATTCAAGCTCAGC	CTACTGTGCATGTTTTTCCATC	(ATA)15
LBSSR0006	CGTAAATCGGAATTGGGAAC	CTTCTTTAACCTAGCCTCCCT	(GAA)11
LBSSR0007	AAGATGGGTTGATTCATTTCTTG	GCACAGTGGAGAGTGTTGCAC	(AGA)13
LBSSR0008	TTTAGGGTTGGACACAAGTCG	AACAGGACCTCCTTGCAATGT	(AAG)13
LBSSR0010	CGAGAGGCTACCAGACACTAGG	TCTCCAATATGTTGTACTCCGCT	(TTTC)11
LBSSR0011	GAAGGTATTAGGCAATCAAGC	TGTAATAATCAAGTCAAAGCATGAG	(ATT)21
LBSSR0012	AGGAACTAAAAGGAAAAGGCTC	CATCACATAAATTAACACTGATGGAG	(TTA)14
LBSSR0013	ATCAAAATCATAAGCCTTGGAAAC	CTAGAAGTTCATGATTCTCAAAGTTC	(ATC)11
LBSSR0016	GTGTCAGTGAGAGGACAAATGGT	TTCCAGTTAAGAACGTAACATATGC	(AAT)16
LBSSR0018	CATCGTTTTCCATTATCCGACT	TTTAGGAATAGATTTTGGGCTCA	(ATT)20
LBSSR0019	TTCCTTGCACAAAGATGGATTAG	CGTACACTTTACCCTCCCCAG	(ACA)18
LBSSR0020	CAGTTGAGTTGCCTTAATGGCT	ATGCATGAATTGTGAAAACCTGG	(ATA)14
LBSSR0022	TGTCTGATATTGATGGACAATGG	CGAATCTCAGGAGGAGGCTG	(TTA)23
LBSSR0023	ATCAACCTGTACCTTCGGCTG	AGATGCCTAATGAGGCCAATG	(TATG)12
LBSSR0024	CAATATTCTATTACGGTCACTAGCTGT	ATGCTGAAATGATAACAATGCG	(ATT)14
LBSSR0026	AAATGTCCGTTTATCTGTGTTAGG	GTCGTTTTAAAAGTTCAAGGCAC	(ATA)18
LBSSR0027	CCTCACTTCAAGGTATGTTAAGGC	ATTGGTTGGTTCATATCATAAGGTC	(TCTT)11
LBSSR0029	AAACCAAAGAGTTGAACATAGATGA	AAAAGAATGTGATATACACAAAGGC	(AAT)20
LBSSR0030	TAATCCTAGTCCACGCAGTACG	CATCTTGTAACCTATTTTCTCCCTTG	(TTA)23
LBSSR0031	GACCCACAAGGGTAAAACCTGG	GGGTAATTAAGGATTTTGAGTCCT	(AAT)17
LBSSR0032	TGTCTGATATTGATGGACAATGGT	ACGAATCTCAGGAGGAGGCTG	(TTA)23
LBSSR0033	TGAGCATCAACCTGTACCTTCG	GAAAGATGCCTAATGAGGCCA	(TATG)12
LBSSR0034	TGCTGGATAGGATTCCTAGAATG	CCTTTAACCCAAATACTGCCTTC	(AGA)17
LBSSR0035	CATAATTATCCGAACAACCTCTCAG	TAGGAATGGTGCTATTCCTTACC	(AAT)22
LBSSR0037	TAGGGTTGGACACAAGTCGTG	CAGGACCTCCTTGCAATGTTC	(AAG)13
LBSSR0038	CTTCGAGAGGCTACCAGACACT	TCTCCAATATGTTGTACTCCGCT	(TTTC)11
LBSSR0039	CCAAGATGTGGGATTTCACTG	AGTCCGCGATAAAAATATACATGG	(GTT)11
LBSSR0041	GACCCACAAGGGTAAAACCTGG	GGGTAATTAAGGATTTTGAGTCCT	(AAT)17
LBSSR0042	TGTCTGATATTGATGGACAATGGT	CGAATCTCAGGAGGAGGCTG	(TTA)23
LBSSR0043	TCAACCTGTACCTTCGGCTG	GATGCCTAATGAGGCCAATG	(TATG)12
LBSSR0044	GGTCTTTCGGAAACAGTCGTC	TGATTCTCCTACATTCCATAAGGC	(GTT)11
LBSSR0045	TCGGGTAGTGTATTGGGCTAG	GGCGTTCACTATTCTATTACCG	(ATA)18
LBSSR0046	TTCGAGTCCATGTTACTAAGAGC	TTCATAAGCTTCGGAGTCTCTG	(ATG)13
LBSSR0047	TCCACATGCTCATATGCTTTGTA	TGAGAGATCATATAGCCGACTCA	(ATA)13
LBSSR0049	TGATGTCTATTTTATCCCACACA	CTGTCTAATGAATGACTAGCTGTCTG	(ATT)25
LBSSR0050	GAAATTAACAAGCCCCAGATG	CATCTTCGATTGGCTACACCTAC	(TTA)16
LBSSR0051	GAAAATAACCATAGCCCCGAGT	CCCCTTGTCAGCTACATCATCT	(ATA)14
LBSSR0052	AGTTTTATCGTGCCCACTCGT	TCCTCTCTGCCAATTGGCT	(TTC)16
LBSSR0055	TATGATCGACTCAAACCTCGTCC	CATCTTCATTCTGAAGCGACTCT	(GAA)18

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LBSSR0064	TGCCTTAAATGTCCATGCCTC	TGACCCTTGACACCTCCACAG	(ATT)15
LBSSR0066	TTCCATTACACTTCGCTGCTC	CAGATGAAGAGGGACGAGACG	(TCT)17
LBSSR0067	AGCCAAGGCTGTGATTGACTC	CCTCTTGAAGCGCCAGTAGC	(AAG)17
LBSSR0070	CAGACCATAATTTATCCCCTCTTG	TCTGAGACGAAATCCACATGC	(AAG)21
LBSSR0076	ACAATCAAAGCTTGGAAATAGCTC	CGTCGCTAAATTGCTCATAACT	(AAG)22
LBSSR0077	TCACGCATCTTTGAAATGTCTAG	AAAGAGCGTGGAGGTTACAG	(TTA)23
LBSSR0082	TTGACTCACCATTGCACTTAGTC	TTAGCCTAATTAAGAGTGTGGGC	(TAT)14
LBSSR0083	TTCTCCATTTTGACCAATCTGAC	AGGTGAAAGCCTTCTCCGAC	(TTA)14
LBSSR0084	TAGCCTCTTATCAATACCAGCG	TGGGTAAATCCGAAAATGTTG	(TAT)14
LBSSR0085	TACCGGCAGTTCATTATCCTG	TACGATGCGTATGCAGAAGGT	(ATT)15
LBSSR0087	ATGTCGAAACGCAAATGTGA	CCGACATTCATTATCCTGTCAT	(ATA)15
LBSSR0088	TCGACTTGGGGTTGGATACA	AACAGGACCTCTTGGTCCATG	(AAG)16
LBSSR0090	TATGGTCTCCTTTGTCAAAGC	GCCTATGACTACTGATTCCCATG	(AAG)17
LBSSR0093	GCTCAGGTGCCTTAGCTAGTTG	CCCTCAGCATTATGAGCAACC	(TTC)15
LBSSR0095	TCTATTTAGTTCCTGAGCTAAGCC	CTTTAGGCCCCACTTAACCA	(TTA)19
LBSSR0097	GGCCTAGAGTAAGGTGACATCC	ACCATCTTTGTCAACTCACGG	(AAG)20
LBSSR0098	TTGGAACATAGAAGAGCACGG	GAGTCGATGCTGAGGTACCG	(TCT)21
LBSSR0100	ATTACTCGACTCATTCCGCCT	AACCCAGAAACCAAGGTCCG	(TTC)29
LBSSR0101	TGCCTTAAATGTCCATGCCT	AAGTTTGCGTAGGTGGTTGAC	(ATT)15
LBSSR0103	GGTTCAACATACAGACAGTCCTACA	ACTAGAACTTCCGATCCTTCCAG	(ATA)16
LBSSR0107	AGATCAAATGGTTGAGGTCAAGA	CTGCATCTCTATCTTCCCACTGT	(GAA)18
LBSSR0108	AAAACCCTCTGCTCGCCGT	TCGATGTACAGTGACGGTCCG	(TTA)19
LBSSR0109	GTCCCAAACCTCTAATTAACCGACT	GCCAAGTGACAGCTCTTGCTC	(ATT)19
LBSSR0110	GTGAGAGTGCAACTGAGAATTCCG	ATCCTCTTCACCAACCGTAGTTC	(AAG)19
LBSSR0112	TGAGACCTCTACCCAAAGTTATCC	TCAAGAGCTTTCTTCATCGAGTC	(TATT)15
LBSSR0118	TACGAGGATTCCATGCTAGTTG	CGATATCGGAGAACTCCATCT	(AGA)18
LBSSR0119	TTCCGTCAAGTGAAACGTCG	GAAGTATCGACGGTCAGCTC	(TTA)20
LBSSR0120	TACTCAAACAAGGCCTTCCGT	AAGGTGTGTACAGATTGGGGC	(TCT)20
LBSSR0121	TGAATTGAGCCATGAATTGTTT	TGTCGGTCATACCACTGATAATG	(TTC)18
LBSSR0122	AGCTTTGTCTCTCAAGATGAAGC	CAGCTCTAGTGGTGAGGATTTTG	(TTC)14
LBSSR0204	CCACTGAAACAAAACCAAGAAAC	TAGTGAATTGCAAATATCAGATGC	(ATT)24
LBSSR0206	TGCCGTTACCTTTTTGACACTC	AGATGATTTCTTTCCATCTGGC	(AAT)23
LBSSR0208	AATTCATAAAGCGCCTGCTATC	CGTTCGAAGGAAACACATGTC	(ATT)18
LBSSR0210	AGGAATTGTTGTTACGAGGTG	CGTCTGTAGGTGACTGAAAAGC	(TTC)18
LBSSR0211	TGATTCTGCTCCATCTTCATACG	GCAGTCACCTACTCACCTCACG	(CTT)34
LBSSR0215	TACCGATCTTACGTTACTCCGA	ACCTAGCCGACTATAGCTCAGC	(TG)17
LBSSR0221	ATTCTGCTCCATCTTCATACGTC	TTCAGCAGTCACCTACTCACCTC	(CTT)34
LBSSR0231	CTTGGGTTCACAACTTAATAAAG	TCAATCATTCGTAAGCATTATGATC	(AGA)24
LBSSR0234	AAGTGATGGCAATATGATGACAAC	ATGGTGGGATTATACTGGGCT	(ACA)38
LBSSR0236	CAGACCATAATTTATCCCCTCTTG	TCTGAGACGAAATCCACATGC	(AAG)21
LBSSR0239	GCCTCTGAGGGTGTAAACTAAGC	TAAGGACGGATAGCGTATATTGAC	(TA)19
LBSSR0242	TTCTTACCACACCACCGACG	ATTTGGCACCTTTATTGTTGC	(TTC)26
LBSSR0245	CCATGGAAGTATTCTCGTCACC	TGATGAAGTTGGAGCTACTAGTGG	(ATT)11
LBSSR0248	TGAGAACATCGTATTTTTCAAGTAAC	CCATTCCTTTGCCTTTCCTATTC	(AAT)15
LBSSR0249	CTCTGAGTGCGTCCACTTGAG	ACAATGGTTGAACGAGTGAGAAG	(TTC)14

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LBSSR0253	GATGTAACAGTAACGCCCTCTG	TTCATTTTGCTATCTCGAATCC	(GTT)11
LBSSR0254	ATGGATGAATGAATGCTCGTG	ACTCCTCTCCCTTCAAATACC	(AAC)12
LBSSR0255	GTGAGAGTGCAACTGAGAATTCC	ATCCTCTTACCAACCGTAGTTC	(AAG)19
LBSSR0256	TCATCTTCTCACTTGAATTACAG	TGATGAACTGCTAGTGAGCTTGA	(TTC)21
LBSSR0258	CCAAAGTCAATTCCCATAATGTC	CCAAAGTCAATTCCCATAATGTC	(ATA)14
LBSSR0259	TTGAGGACGATGCTATTGGTG	CCTATTTATATCCCTAGCCCGA	(TGT)16
LBSSR0260	ACCGCCATTGCATTATTCTATC	GCGATGCGTATGTGTAAGCTG	(ATT)20
LBSSR0266	CAAGACCCATGTCAGCTTCATC	GCCCCGAAAAATCATCATTAG	(TTA)16
LBSSR0270	AGGAGTCAGGTCTGCGTACACT	CGAGGTATTTACTCCAGTTGGTG	(GTT)14
LBSSR0276	GAAGATAAAGGGAAAGACATGGT	TTTACTTCCAATTCACGCTTCA	(GAA)16
LBSSR0277	AGGAAATTCGTCCAAACAAGAC	TGGTATCAGAAGCCGATCATAG	(ATT)14
LBSSR0281	AGAGAGCAAGGCATTTGGAAG	GAACCGATGAACGAACCAGC	(AAG)15
LBSSR0285	ATAGCGTCTAAGTTCGATTGC	GGCAGATCCAGGATTTTACTC	(ATT)18
LBSSR0289	ATCCTTTTCCAAATCCAGCC	TCCCGACTTTCTTTTCTAGACG	(TTA)16
LBSSR0290	AGAGCTTGCCCTCTATCCAAG	CTCAAAGTTGGTGCTAACATGG	(ATGT)10
LBSSR0291	AAGGCTTTGGTCTTCTTACTGTA	GCAAGCGACTGTGTATCTTCAA	(TATT)14
LBSSR0294	CCTACCTAGCCATATAAGTTCGG	TGATGTTGACGTGACTTAAGCG	(AT)14
LBSSR0295	TTGAGTCCCTGACTAGAGGGC	GACTCGGTGGATAAAATCATGTC	(TA)17
LBSSR0297	TGCTGCAGAGAACAACATTAGG	TTCTTATCTCCCATTCAATCCAGT	(AG)11
LBSSR0301	AATACCGGCGTTTTTCGTT	ATGCGGATTGTCATAATACCAGT	(ATT)20
LBSSR0302	CCATCTATTGTTCTCATCACTCTTG	GAAGAGCGGGTCATAAACTACTG	(GTT)17
LBSSR0303	ATGGTGGGATTATACTGGGCT	CGAACCATCCCCTTAATGAAG	(TTG)12
LBSSR0305	CTGTTAAATTCTGGCATAAGGGA	AAAAGTTCACAAAAATCACTACAG	(TAT)23
LBSSR0306	GATTTGGGGTGAATACAAGC	GCTTTCTCATTTTAAACAGGACCTC	(AGA)17
LBSSR0308	TAAGTGGTGGCATAAGTGTGGAC	AGCCTAACTGGTCAATGAAATGT	(TTA)11
LBSSR0309	GTAGTAACCACGCCGATTGC	CTGAGGTGGATACACTACGG	(GCCTGT)12
LBSSR0313	GCTGAACCATTTGCTGATGAG	GAAGACAGGGTCTGGCAAG	(TTC)15
LBSSR0322	CCTCACAGAAGGTCAATCATAACAG	GAACACTCTTGATCATGGCTAG	(AT)20
LBSSR0323	ATCAAAGGTCAATGGTCAGGC	TAGTAAACTGGTCCACATGGCA	(AT)11
LBSSR0324	CCGTACCTATATAGTCGGGCAG	ATGAGGGGATATCAAATGAAGAAC	(AT)17
LBSSR0332	TGAGAGTGCAACTGAGAATTCC	TCCTCTTACCAACCGTAGTTC	(AAG)19
LBSSR0333	CCAAGCAATCAATTCGTAAGC	CACCCATCATTGGGTTTACAAC	(TTC)17
LBSSR0335	TCGTTGTGGGTGTTACAGGTG	AAACCTGAAAAATCCAATTCAGC	(TTG)11
LBSSR0338	AAGATTTGGGGAAGACGACG	CATGGCAGCATATTAGCGAAG	(GAA)13
LBSSR0339	GAAATTAACAAGCCCCAGATG	CATCTTCGATTGGCTACACCTAC	(TTA)16
LBSSR0341	AGCCCAAATTGCTCTATCTAGC	TTAACAATCCTAAACCTCTTCC	(AGA)11
LBSSR0342	ATATGCCTAGCCCCGAAATCAC	GGGATGCTTTATGGAGACTACTG	(AAC)11
LBSSR0344	TTCTGGAATTGGCACTGCTG	GGGCTATCAAATTAACCCGTC	(TTC)12
LBSSR0349	CTGTAATCCGCTGGCATAACG	CCAAAGGTGTGAAGACACGG	(AAT)20
LBSSR0353	CGATTTTACGAAACAGCTATC	TTACTACTGAAAAGGCAAGGCA	(ATA)18
LBSSR0356	ACACTACGTAAGCCCACCCG	GACCCTGGACTGACATTGGAC	(TAT)22
LBSSR0358	CCCCTATTTATATGCCTAGTCCG	GGAGAAACATCGTGTGGGATG	(ACA)11
LBSSR0360	CTGTTGGAAGTGATTAGCGGA	GGGGCATTATTAGAAGCTTATCC	(AAT)13
LBSSR0363	GTAGTAACCACGCCGATTGC	CTGAGGTGGATACACTACGG	(GCCTGT)12
LBSSR0365	AAGTTCAGGGTAATTTTGGC	AAGTAGCGATGTACGAGGCATT	(TTA)20

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LBSSR0370	TCTGAATCTGCAACTTTGGAGG	CCAATTCAGATGAGATAGCCAGT	(AT)14
LBSSR0373	ATACATAAGCCCTTCGGCTATC	GAGGCTCATAGATACGCAAGTG	(AT)14
LBSSR0382	CAGGAGCACCCATTATTCTG	GAAATATGCACGGGACAGGAC	(TA)16
LBSSR0384	GCTATAATGACACCCCGAGCT	AGGGTTATCCGTACATCTGCAG	(TA)13
LBSSR0386	TCTTTGGGCTCTCCTACATGC	ATTGATCAGGTGGATAGCGTCT	(TA)13
LBSSR0387	CTGTATTTGTTACCGAATCGCTC	CATCATAACGCCATAACACAGC	(TA)13
LBSSR0390	CAGACCTGTGACCCCTCAGG	TCATCGTCGACGTAATAATCAGG	(TA)15
LBSSR0391	ACCCCAAATGAAAGTAGAATGC	TTCGGTGACTATAAATCATCTCACT	(AAG)11
LBSSR0393	AGCGACATTACCCAGAGAAGTC	ACTTGAAATGACCCCTCTTGCC	(AAT)11
LBSSR0394	GTTGCTTAATTGAATGGGTGC	ATCATCCAAGGTATTGTCGTCC	(GA)15
LBSSR0397	TACATGTCCACCGATAACAATGAC	TTGGATAGGCCTGAATAGCGT	(AT)17
LBSSR0398	GAAACTGAAACCTGATTATGAGCA	ACGCCACATACTAACATGATGG	(AT)14
LBSSR0399	TGGCCTAAGAGAGGTGTCACA	CCCCACATAAAATGAAGACTC	(AT)11
LBSSR0400	CTTAAGGCCAACTCAGGAACG	TACGATGTCCTACGAGTAAGCG	(AT)15
LBSSR0402	CCTCACAGAAGGTCAATCATAACAG	GAACTCACTCTTGATCATGGCTAG	(AT)20
LBSSR0403	ATCAAAGGTCAATGGTCAGGC	AGTAAACTGGTCCACATGGCAG	(AT)11
LBSSR0404	CCGTACCTATATAGTCGGGCAG	ATGAGGGGATATCAAATGAAGAAC	(AT)17
LBSSR0405	TCGGTTTGCAGAGTTTACGC	CTCAACCGAGTCCTTTTCACC	(TC)11
LBSSR0406	GCCCAAAGTTATTCTATGAGT	CACAATGGGTGTTGTACCCAG	(AC)11
LBSSR0407	CGAACATCTGGAAGTGGCAC	GAGCATGGATACTATCTGTCACG	(TA)19
LBSSR0413	GAGGACCGATAAGGTAATAGTGC	ATCATGGTCTAGCAGTGTAGTAAGG	(AT)20
LBSSR0415	TGAATTATGTTTGGGCCGTAC	GACAAGGCTGCCACTACACG	(AT)16
LBSSR0416	TATGCATATATGACGCCTCTCC	TTTGGGCTCTCATGTATGCTG	(TA)11
LBSSR0418	GGGATATAGCGATGAGGATAATTC	CATAACAAGGCTCGGTGTCAG	(AT)15
LBSSR0419	ACCAACCTTATGTCACTCCGAT	CTATAGCGGCTCGGTGTCAG	(TA)13
LBSSR0421	ATCCACCCCATACATACTCATC	GTTGCAACTAAAGGAGTTGTCCA	(TTC)13
LBSSR0422	CGTAAACTCTACCCTCCCCAG	TCGGATCCATGAAATAGTCCG	(TTG)11
LBSSR0423	GGATGAATGAATGCTCGTGC	TCCTCTTCCCTTCAAATAACCC	(AAC)12
LBSSR0425	GAACAGGAAACCTCTCAAGCAG	CGAGCTTCAAAGAAGCAGT	(ATG)13
LBSSR0426	TAGGGAACCTGATATATGCGG	CGCCACACCATTCTATCTCTG	(TTG)11
LBSSR0428	GCTCTCCTTCATTGTCACTCG	TGCTAATGAATAGGCTGAGATTG	(TTC)12
LBSSR0429	CCATGAACTGATCAATGACCTCT	CACCTCCACTTGAGTCACTTTTC	(AAG)14
LBSSR0430	AAGTCTAATGGAGCTACAACCCG	CATTACTTCAAAAATGGTAGGC	(ATA)11
LBSSR0431	GATTACACGACACAGTTTCAGGA	ATCGCAACTCTATTTCTGGGAC	(AT)11
LBSSR0440	TACGGGCACAATTATCACACG	CTAGCCTTTCGGCTCACTCTG	(TA)11
LBSSR0443	CATGGCAGTAGTGTGCACCTC	GGAATCTTATGTTGCCCATAGG	(TA)11
LBSSR0446	TCAGGTTATCCGTACCAACTGC	GAAACCCCGAGCTCATATGG	(TA)15
LBSSR0448	TTCCGGAGGTGCTAATCGAG	AGTACTAGACACATGGACGGGAC	(TA)11
LBSSR0450	AGACCCGAGACACCTCAGGC	GATCAGGTGGTAGTGCGTACATG	(TA)16
LBSSR0457	AGTCACCTCAAGTTGTGCGC	TTATACTAAACTGGTGCATTGGC	(TA)11
LBSSR0461	ACTTCACTCCAACAGATTCAGGA	ATACCTCCGAACATCGTTATGAG	(TA)13
LBSSR0464	GGTGCTTATGATCACCCAGT	CTAAACATCATACCCAACCCG	(TG)16
LBSSR0465	TCGGTTATGTCTGTGATTTGCA	GGAGCCTCAAAGACTGTCGTAG	(TA)16
LBSSR0467	ATTTAGCAGGGATGTTCACTC	TATAGATCAACTGGTCGGGACAG	(TA)14
LBSSR0472	GGGAGTATTTTGTGGGCTAGTTG	AGCATTGGTTGCTGGGACTC	(TA)16

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LBSSR0480	CCAGTTGTACAGGTAGCTATGCG	GACTTTCCTTGGGCTCTCATG	(TA)11
LBSSR0483	AGCTGCAGTTGGATTTGACG	TCAGGTCTGCGTACACTCTACC	(AAC)10
LBSSR0484	TACTTCGGATACATCAATGGCA	CATAGTCCTCGGTATTAGCTGGA	(GAA)12
LBSSR0485	TCAATAGAGTCGAGTTGTCCCTC	CTGCTAAAAATCCACCGACTG	(ATT)10
LBSSR0488	TTTGTAAGCGGAGATGGTAGTG	CGTATGGATATCATTATGCGCT	(TA)13
LBSSR0490	TGCTGCAGAGAACAACATTAGG	TCTCCCATTCAATCCAGTATGC	(AG)11
LBSSR0494	AGCCGATAAGGAATAACGACG	GGCTACAGCAATTCTCACTCCA	(AT)12
LBSSR0498	AGTCAGCAACGAATCTAGCATG	ATGGTCAAGGTGCGTATGTTC	(AG)13
LBSSR0501	GTGGTATCCGTACCCAATGTC	ATGAAACCTCGAGCTTATATGGC	(TA)16
LBSSR0505	GTAGCCATGAGAGCTCACACG	CTCATCCTGCTAGTCATCACACC	(TA)15
LBSSR0511	TTGAATATCAGCGTTTGGCC	CGTACACTCTACCCTCCCCAG	(ACA)12
LBSSR0512	CCCATACTACTCATCACCTCTGC	GTTGTCCAAACAACACACAGTGC	(TTC)13
LBSSR0513	GGTCTGCGTACACTCTACCCTC	CAACTTACACGCATCTCAACTCA	(GTT)12
LBSSR0515	AGGTGAGGGTAAGGTCTGCG	TGTAGGTTGTCTGATCTCGGAGT	(TGT)11
LBSSR0516	GCTCTCCTTCATTGTCACTCG	TGCTAATGAATAGGCTGAGATTG	(TTC)12
LBSSR0517	AAATAGCCTAAGAGCCAACCACT	CAGTCCCTCCTAGTCATGTGC	(AT)11
LBSSR0519	GCCTTATTTAGTGTGGTCGTATG	CCATTTTGGGAGCTGCTGTAC	(AT)14
LBSSR0525	TCACCGAGTCCATCACTAGAGG	GTAATCATATGCCATCCTGGC	(TA)16
LBSSR0526	GGAACCGAAACCTGGTCATG	AGTGTCTGCTGGTAGGACGC	(AC)12
LBSSR0527	TTTCAGGTTATGGGTACGTCG	ACATGATATGACTCACTTGTGCG	(TA)11
LBSSR0530	AGCAATAAGGGAGTCAGTACGG	TCAGTCAACCAATCAGCTTATGA	(AT)17
LBSSR0532	CCTTGTCGGCTTATGTACTGTG	ACACCCAAACATAATATGGAACC	(AT)14
LBSSR0539	ATTGGGAGCTCAGAGTCGACA	CGATAGGTGACCTGCACTTCC	(AT)14
LBSSR0541	CTTAGCTGAGCCGAGGGTCTT	TCCTCATGATTCTCCTACATTCT	(GTT)11
LBSSR0542	CGGACTGGTCACTAATAGCGTA	TACCCTCTCCAGACCCCAAG	(ACA)12
LBSSR0543	GTCAGGTCTGCGTACACTCTACC	GGCCTCAGAAATCTCTTTATCAG	(GTT)12
LBSSR0544	CCCATGTTAGTCAACCTGGATCT	TATGCCACCTGGCTTAATCTTAG	(TAT)23
LBSSR0545	ATCACATTTTTTCATCACCTTATC	TGATACATCTTCGTGGAACCTG	(ATT)12
LBSSR0546	CCACATCTTGACGAATAGCTTG	AGCTGCAGGAAGACTTTGATAAG	(TTC)12
LBSSR0553	CAGGTAAATAGCCTAAGAGCCAA	AGTCCCTCCTAGTCATGTGCA	(AT)11
LBSSR0554	ATTCACATGCTCTCCAATCCTC	GTCGTTTTAGGCATGGACTCTG	(AT)11
LBSSR0555	CAGTCGGTTGGGTAGTTCACG	TAACCTCTAACTTGCCGCTGA	(TA)15
LBSSR0556	TACCGATCTTACGTTACTCCGA	ACCTAGCCGACTATAGCTCAGC	(TG)17
LBSSR0558	ACATCACCTTGCTCTATGCCTAACT	AGATCAAGTCCCTTCTGAGTTC	(TA)11
LBSSR0559	CCCCAATACAAGTAGCATTCACT	GCTCCTGCTTATCCAGGTAGAA	(TA)11
LBSSR0560	GCCGAGATACATATACCCATTGA	GGTGAGCTTGAATAGTGTGGA	(TA)20
LBSSR0561	AGTTCAGTGTTATCCGTACCCAA	GATGATGATTGTGATGTTGACGA	(AT)12
LBSSR0564	TTTGGTATTCTCCTGAAAGCCT	CCCAAATACAGAAATATGCAACA	(AT)16
LBSSR0565	ATCACTAGCTCATTCTCTTGCGT	TCCTTTTACTACCTATGGCTCA	(TG)14
LBSSR0566	TCTTTGTCGCTACTGCAGGA	ACCCCTCTTCTGGAGATTGTC	(TA)14
LBSSR0567	TTACAGGAGCACTCCACCATT	TTAAGACATAGGGCTGGGACAG	(TA)19
LBSSR0568	GCTTCCAGGTTATTCAAGAGTTTC	TACCTGGCCAATATAGCGTG	(TA)14
LBSSR0570	GGCCGTTGTCATTAAATCACT	TGCTACTTCAGGAGCATATCGA	(TA)15
LBSSR0572	CTTCGGATACATCAATGGCAT	CATAGTCCTCGGTATTAGCTGGA	(GAA)12
LBSSR0574	CATACTTCATTCCCAGACATCCT	GGATTGAAATACACAAAAGCCAG	(ATT)12

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LBSSR0575	ATCGATTAAGGGTGA CTCTGTTC	CCAAGTGAGGTTCA CATCTACCA	(ATT)13
LBSSR0589	AAAGCCGATAAGGAATAACGAC	AGGGCTACAGCAATTCTCACTC	(AT)12
LBSSR0594	AGATGTTTATGGGCTTTGACGT	CGTACCCAAATACAGTGGTGC	(TA)20
LBSSR0596	GGGATGTTGTGTGTAGTTTCAGC	GTCCCAGCTATACACAAGCCA	(AT)12
LBSSR0791	GCAGAACATTCCACTATGAA	GCTATGGAAGAGGAGGTTAA	(TA)26
LBSSR0792	AATGCTTTCCTACTTACCA	GGCACGTTACCACCTTAT	(TA)8
LBSSR0795	GGAGATGAATAATATGCCTTGC	ATTGCCTTAAGCCACAAGT	(TA)29
LBSSR0796	ACATAAGCCGTTTCGTCTATC	GCTTCGTGATTCCCTCAT	(TA)21
LBSSR0797	TTAGGGATCGTTTGGTAGGA	CGGACTATGGTTTGCTATGT	(TA)15
LBSSR0811	TTCACATGCACATCACAC	GTCTCGCACATACTCTT	(CA)11
LBSSR0812	AGTTCACAGGATTTGATGGA	ACTGTTGGATTTGTGGGAA	(CAA)5
LBSSR0814	GTGGACAAGGGATCTACC	TGCATGTGCCTCTTACTAT	(CA)7
LBSSR0816	GAGGCAAGGATTGTCCAA	GCTCAAGATGTTTAGTTAGAGG	(CA)6
LBSSR0820	CTATAAGTCCGAAACGAAGT	CGTAAGGTCTGCGTACAC	(CAA)6

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