

Supplemental Table 4. Primer Sequences.

Primer	Sequence(5'→3')
I108	CACCATGGTTAAAAAAGCGAAATGGC
I109	TCACGGCGTTCTCTCTGC
I110	CACCATGGGGAAAAAAGCTAAATGGT
I111	TCAGCTGCCTGCTCCGTT
I112	CACCATGGGTAAGAGTTGGTTTTAG
I113	TTAAGCAAGATTAGTATCTTTCTTAGTAA
I114	CACCATGGGTAAGAAGCTGTTAACATGTG
I115	TTAACTGCTACCACCATTTCTTCTC
I800	GGCCATTUATGGGAAGATCTCCAGCTTCT
I801	GGTGATTUATGCAAGCCTCTGTTTTATTGG
I118	CACCATGGGTGCTTCAGGGAAAT
I119	TTAACCTCTCGGCTTCTCGA
I120	CACCATGGGTGGGTCAGGAAATT
I121	TTAGCTTCGCTGGCTCTTG
I122	CACCATGGGTGGCTCTGGAAATT
I123	TTAGCCTCTCTGGCTCTTTGC
I124	CACCATGGGTTCTGGGAATTTGATT
I125	TCAAGCACCTGGAAATGACA
I126	CACCATGGGATCTGGATGGCTG
I127	TTATCCGGAACCAGGCTTT
I128	CACCATGGCTAAGAAGAAGGGCTTG
I129	TCATCTCAAGCTGCTCTGCTT
I130	CACCATGGCGAAGAGGAGGTGCG
I131	TCAACAATTATGTTGATATGTGGTCA
I132	CACCATGGGGAAGAAAGGAAGTTG
I133	TCACGCAAATCTGTAAAAAGCC
I134	CACCATGGTGAAGAAAGGAAGTTGGTT
I135	TCACACAAATCTGTAAATTCCTTTC
I136	CACCATGGGGAAAACCGACGGA
I137	TTAGTACTGAAAATCTTCGTGAGCA
I138	CACCATGGCTAAAAAGAACGGAACG
I139	TCACTTTAACCACCGGAGCT
I140	CACCATGGGTAAGAAGAGCGGTTCT
I141	TCATCTTAACCATCGCCTATAATC

SUPPLEMENTAL DATA. Bürstenbinder et al.

I142	CACCATGGGGAAAAAGAACGGC
I143	TCATCTAAGCCAATTCCTAAAGTCA
I144	CACCATGGGGAAAAGTAGCAAATGG
I145	TCAGTACATGTTGTTGTTTCCCTG
I146	CACCATGGCCAACTCCAAACGTT
I147	TTAATGAGAGAGAAGCTGACGAGC
I148	CACCATGGGGAAGAAAGGGAGTG
I149	CTAATGATCATGCCTCCAGC
I150	CACCATGGGAAAAGCGTCACGG
I151	TCAGTACCTATACCCAATTGGCA
I152	CACCATGGGCTTTTTTCGGGAGA
I153	CTAAACAAGAAACGAAGAATGCATC
I154	CACCATGGGTTTCTTTGGAAGACTGTT
I155	CTATTGAAAGAAAAGAGGATTAGAACT
I156	CACCATGAGAAAGAATCTCACAAAATTGAC
I157	TCACCAACGCATCCTACG
I158	CACCATGGGAAGAGCTGCGAGAT
I159	CTAATTATAGAATCTAAAATCAGTCTCG
I160	CACCATGGGCAGAGCAGCAAGAT
I161	TTAAAGCGGATCACAGGAACA
I162	CACCATGGGAAAGACTCCTGGTAAATG
I163	TCACCGTTTCCAGTCGGT
I164	CACCATGGGAAAGACTCCAAGTCCT
I165	TCACTCAGCTTTCGTTGACTCA
I873	GGTGATTUATGGGAAAGCCTGCAAGG
I874	GGCCATTUTCATCTCTTCCACTCTACCGG
I168	CACCATGGGGAAGTCTACAAAATGGTT
I169	TCACCTCTTCCGCTCTGC
I170	CACCATGGGAAGATCTCCAGCTTCT
I171	AAATCACCTCTGCCATTTTCTATCC
I172	CACCATGGGTGTTACAGGAGGATTAGTC
I173	TTAGGTGCTGCTATTTAGCTTATGTG
A027	CACCATGAAACTCACCTATCAATTCCTTCATC
A028	TTTATAAGCATCATATTTATTACTCTTCTTCT
I752	GGGGACAAGTTTGTACAAAAAAGCAGGCTTCATGTCTGCAAACTTCTATATAA
I753	GGGGACCACTTTGTACAAGAAAGCTGGGTCTTAGCAGAAGCAAACCTTCATT

SUPPLEMENTAL DATA. Bürstenbinder et al.

I1378	GGGGACAAGTTTGTACAAAAAAGCAGGCTTCATGGATACCTTAATCAAGCAGAC
I1379	GGGGACCACTTTGTACAAGAAAGCTGGGTCTCAGAAACAGCATGCATTTTC
I1374	GGGGACAAGTTTGTACAAAAAAGCAGGCTTCATGGATAATTTGGTTAAGCA
I1375	GGGGACCACTTTGTACAAGAAAGCTGGGTCTTAGTAACACCGAAAGCAGA
I192	CACCACTTCGAAGTGAACATATGGAAC
I193	TGTTGTACAACCATCACCTGC
I198	CACCATCAGAATCTCTCCACTAAACCTAAT
I199	TTTGCCAACATCACTTTCCTT
I202	CACCGTGTAACACGTGACAACACCA
I203	ATTGAATTAACGTTTTCTAAAAGCG
I220	CACCACGCAACATTTTGAGTTATTCTTTG
I221	AATGGTTTTAGTTTGGTTTCACTTG
I1250	GGGGACAAGTTTGTACAAAAAAGCAGGCTTCATGTTGCTCAAAGGAAGGAATAAT
I1251	GGGGACCACTTTGTACAAGAAAGCTGGGTGCCTCTCTGGCTCTTTGC
I1468	GGGGACAAGTTTGTACAAAAAAGCAGGCTTCatcagaatctctccactaaaccta
I1469	GGGGACCACTTTGTACAAGAAAGCTGGGTCCACAAATCTGTAAATTCCTTTC
I1380	GGGGACAAGTTTGTACAAAAAAGCAGGCTTCcagtaacgtagaactcactccatca
I1381	GGGGACCACTTTGTACAAGAAAGCTGGGTCATGAGAGAGAAGCTGACGAGC
I1382	GGGGACAAGTTTGTACAAAAAAGCAGGCTTCACGCAACATTTTGAGTTATTCTTTG
I1383	GGGGACCACTTTGTACAAGAAAGCTGGGTCCCAACGCATCCTACG
I1369	GGGGACAAGTTTGTACAAAAAAGCAGGCTTCatgcatcttaggtccataaatt
I1370	GGGGACCACTTTGTACAAGAAAGCTGGGTCCCGTTTCCAGTCGGTTCTGGT
I035	CGCTGAATCTAACGGTTTTTG
I036	TAGAAGCTCCCTATCATCGCC
A004	ATTTTGCCGATTTTCGGAAC
I053	ACGAAGTCGATTTTTCTCACG
I054	AACGCTTTTCTCGCCTTTTAC
I409	CGGAGCCGGTCTTCCCGGAATCA
I410	AGCCATCCATCGGTCAAGCCATTGT
A015	AGCCAACCTAGGACGGATCTGGT
A016	CTATCCGAACCTTCTGCCTCATT
I088	GGCTAAGAAGAAGGGCTTGTT
I089	CTTCAAGGCACGTAGAGCTTTC
I029	TTGGCAAAATGGTGAAGAAAG
I030	GCTTTGAAGAGATGGCTTG
I095	GAGCAACGAGGTGGTTCAAG

SUPPLEMENTAL DATA. Bürstenbinder et al.

I096	GCTTCCATACTCCGGAGAGTC
A005	CAAAGACCAGCTCTTCCATC
A006	CTGTGAACGATTCCTGGACCT