

<i>SNP</i>	<i>Count</i>	<i>MAF</i>	<i>Forward Primer Sequence</i>	<i>Reverse Primer Sequence</i>	<i>Reporter 1 Sequence</i>	<i>Repo</i>
Pf_01_000130573	10	0.40	CTTCCTCTTACCCATATTTCTCTTTCTCT	TGTGAAAATGAGAAGAGAGAAATATGTGTGA	TCTTCATTATATGCATCTTTT	TTTCT
Pf_01_000539044	10	0.40	CCAAGAAGTCCAAGAATAATAGTTACTATCACA	TTCATGTGTGGCTGCTTTCCTAT	CAACCGTTTCTTTAGCC	CAACC
Pf_02_000842803	10	0.50	ATTCCAAAATATGTTTGCTGCTTTTCA	AGAAATGCAGTGGTACTTGTGCTA	AGATTATTGCATCGTACTT	AGATT
Pf_04_000282592	10	0.30	TGGTAATTTGTTGATTGCTTACATTGTTATGT	GACAAGTTTTAATATGAACGGTCAGGTAGA	ACTCGTTCATGCTATTATA	ACTCG
Pf_05_000931601	10	0.30	CGATGACTACCATAATGATGTTTCATATTCACA	TCCTCATTTCTGTTATTAATTTGCATCATCTTCA	TTCGGGTGGTTGATCA	TCGGG
Pf_06_000145472	9	0.44	TGTTTCATGTGCCCTCTTTTTCTG	GGAAGGTAGAATATCAAATACAAGGGATGT	AAGCATAGTAAGAAACATT	AAGCA
Pf_06_000937750	11	0.36	CTAGCTCAGCTTCCAATTTGTCAAA	GTCCACATCCATATACTGTGTACAA	TGCATGTAGCAATACCCCA	CATGT
Pf_07_000277104	11	0.36	AATTAATATTATGAATATGAGTACTCAGCATTATCTT	ACGTCCATATGTTGAAGATGATTGCA	CCACCGAACTTATATTT	CACCG
Pf_07_000490877	12	0.42	CGTAAACCAGGAGTAAGACGTAACCT	AGATGAGTTAGCAACGAAACCATCAT	TTCCATTAGCTAAATTC	CATTA
Pf_07_000545046	10	0.30	GGATGTATGACAGCTTTTCTAATTTCTCTCT	CCATCCTTTTATGAATGCTAAGAGAATACGT	TCTTAAGCCGTTTAGTG	TCTTA
Pf_07_000657939	9	0.22	CGTCCACTTATGATAATGACCTGAAA	TGCTTTCGTTTATATTGCAACATTTCTTCA	ACGATCGTCAAATTC	ACGAT
Pf_07_000671839	10	0.20	GGAACCATTCATAGTTTAGTAGAATCAACCA	AGTAGTCATGTGATATATGATTTTGTGGCAA	CTTTTTCATTTTCATTTAATAAA	TTTTC
Pf_07_000683772	7	0.43	CTTTTATCGTACCACCATTAACTTTTGGGA	ATAAAGATTTTATCATTCTGTTACAAATAACAATGAA	CATCATCATGATGGTTCAG	CATCA
Pf_07_000792356	9	0.22	TCGTTTAATGGGACTCTCTGTTTTCG	ACTAAGTCCAATAGTTTGTATATTCCTTGTTTT	ACCATACCAAATTC	CCATA
Pf_07_001415182	16	0.50	AGGTACTGATTATTGGAATGATTGAGCAA	ACTTCCACCCTCATCACGAAA	AGCTTATCATTTTTTTTATTCC	CTTAT
Pf_08_000613716	8	0.38	CACCTCAAATCTTTTCCAGTTATTTTCTATCCAT	TGGGAAAGAATTAACAATACTTATAAGTGCATAAATA	AAAACTTCATTTAAAATCA	ACTTC
Pf_09_000634010	7	0.43	AATGTTAATTTTATTTTCATCATCCGCATTTAATTT	AAATACCATGTGTACCTGTTGTGGAT	CCCTCAATCCACTTTTATA	CCCTC
Pf_10_000082376	4	0.50	TGTCGTTATTAATAATTTAAAGTTGACT	AGAAAGTATACATTTGTCCGCCGAA	ACGACATTGAAATTTTAT	ACGAC
Pf_10_001403751	9	0.33	ACAAGAACCCTGTTACCTACATTA	CTGGTTTAGTGGATTACCTTCAGA	AATGAAAACGTTACTCCC	ATGAA
Pf_11_000117114	7	0.43	ATTTTCATAAGAAGCTTCCATCTCCTT	AAGTATCTTGCTCGTTTGTATAAAGCA	AAATTTTCATCCAAAAGTAATT	TTTCA
Pf_11_000406215	12	0.42	ACGAATGAACAAAATATGATACACAGGATGA	CTGATGACTTTTTTCAAGTTCAATTTTGGATT	CTACAGCTAGTAATATTTT	CAGCT
Pf_13_000158614	7	0.29	AGTGTGAGCTTTCGTAATATTTTAGTGAAGA	AGAGGATATCCAATAGGGTGTGTT	ATTTTTGGACCCAAAGCTA	ATTTT
Pf_13_001429265	12	0.42	CACACATAFCAAAATCGCATTCATCCT	GAAAGGTATTCAAACCATAAACAAATGGATGA	CCTCCAGCAGTATATAC	CCTCC
Pf_14_000755729	10	0.40	CGTAACAAATCATCAATTAAGTCATCCGAAA	CAAATCTTGAGGAATAGGTTTATATGCTGAT	TTTTAAATTTTCAATTTTTTTG	AAATT