

Oligo Name	Sequence 5' to 3'	Scale (μmole)
Oligonucleotides for HindIII-FspBI digestion of <i>Zea mays</i>		
P1.1_HindIII_bar01	ACACTCTTTCCCTACACGACGCTCTTCCGATCTAACT	0.05
P1.1_HindIII_bar02	ACACTCTTTCCCTACACGACGCTCTTCCGATCTCCAAG	0.05
P1.1_HindIII_bar03	ACACTCTTTCCCTACACGACGCTCTTCCGATCTTTGCAA	0.05
P1.1_HindIII_bar04	ACACTCTTTCCCTACACGACGCTCTTCCGATCTGGTGCCA	0.05
P1.1_HindIII_bar05	ACACTCTTTCCCTACACGACGCTCTTCCGATCTAACAACGT	0.05
P1.1_HindIII_bar06	ACACTCTTTCCCTACACGACGCTCTTCCGATCTCCACCATG	0.05
P1.1_HindIII_bar07	ACACTCTTTCCCTACACGACGCTCTTCCGATCTTTGTTGCA	0.05
P1.1_HindIII_bar08	ACACTCTTTCCCTACACGACGCTCTTCCGATCTCGTGGTAA	0.05
P1.2_HindIII_bar01	AGCTAGTTAGATCGGAAGAGCGTCGTGTAGGGAAAGAGTGT	0.05
P1.2_HindIII_bar02	AGCTCTTGAGATCGGAAGAGCGTCGTGTAGGGAAAGAGTGT	0.05
P1.2_HindIII_bar03	AGCTTTGCAAAGATCGGAAGAGCGTCGTGTAGGGAAAGAGTGT	0.05
P1.2_HindIII_bar04	AGCTTGGCACCAGATCGGAAGAGCGTCGTGTAGGGAAAGAGTGT	0.05
P1.2_HindIII_bar05	AGCTTGTTAGATCGGAAGAGCGTCGTGTAGGGAAAGAGTGT	0.05
P1.2_HindIII_bar06	AGCTCATGGTGGAGATCGGAAGAGCGTCGTGTAGGGAAAGAGTGT	0.05
P1.2_HindIII_bar07	AGCTTGCAACAAAGATCGGAAGAGCGTCGTGTAGGGAAAGAGTGT	0.05
P1.2_HindIII_bar08	AGCTTTACCACGAGATCGGAAGAGCGTCGTGTAGGGAAAGAGTGT	0.05
P2.1_FspBI	GTGACTGGAGTTCAGACGTGTGCTCTTCCGATCT	0.05
P2.2_FspBI	TAAGATCGGAAGAGCGAGAACAA	0.05
Oligonucleotides for PstI-MspI digestion of <i>Hordeum vulgare</i>		
P1.1_PstI_bar01	ACACTCTTTCCCTACACGACGCTCTTCCGATCTAACTTGCA	0.05
P1.1_PstI_bar02	ACACTCTTTCCCTACACGACGCTCTTCCGATCTCCAAGTGCA	0.05
P1.1_PstI_bar03	ACACTCTTTCCCTACACGACGCTCTTCCGATCTTTGCAATGCA	0.05
P1.1_PstI_bar04	ACACTCTTTCCCTACACGACGCTCTTCCGATCTGGTGCCATGCA	0.05
P1.1_PstI_bar05	ACACTCTTTCCCTACACGACGCTCTTCCGATCTAACAACGTTGCA	0.05
P1.1_PstI_bar06	ACACTCTTTCCCTACACGACGCTCTTCCGATCTCCACCATGTGCA	0.05
P1.1_PstI_bar07	ACACTCTTTCCCTACACGACGCTCTTCCGATCTTTGTTGCATGCA	0.05
P1.1_PstI_bar08	ACACTCTTTCCCTACACGACGCTCTTCCGATCTCGTGGTAATGCA	0.05
P1.2_PstI_bar01	AGTTAGATCGGAAGAGCGTCGTGTAGGGAAAGAGTGT	0.05
P1.2_PstI_bar02	CTTGAGATCGGAAGAGCGTCGTGTAGGGAAAGAGTGT	0.05
P1.2_PstI_bar03	TTGCAAAGATCGGAAGAGCGTCGTGTAGGGAAAGAGTGT	0.05
P1.2_PstI_bar04	TGGCACCAGATCGGAAGAGCGTCGTGTAGGGAAAGAGTGT	0.05
P1.2_PstI_bar05	ACGTTGTTAGATCGGAAGAGCGTCGTGTAGGGAAAGAGTGT	0.05
P1.2_PstI_bar06	CATGGTGGAGATCGGAAGAGCGTCGTGTAGGGAAAGAGTGT	0.05
P1.2_PstI_bar07	TGCAACAAAGATCGGAAGAGCGTCGTGTAGGGAAAGAGTGT	0.05
P1.2_PstI_bar08	TTACCACGAGATCGGAAGAGCGTCGTGTAGGGAAAGAGTGT	0.05
P2.1_MspI	GTGACTGGAGTTCAGACGTGTGCTCTTCCGATCT	0.05
P2.2_MspI	CGAGATCGGAAGAGCGAGAACAA	0.05
Illumina adapter sequences		
ILLPCR1	AATGATACGGCGACCACCGAGATCTACACTCTTTCCCTACACGACG	0.05
ILLPCR2_ind01	CAAGCAGAAGACGGCATACGAGATCGTGATGTGACTGGAGTTCAGA CGTGTGC	0.05
ILLPCR2_ind02	CAAGCAGAAGACGGCATACGAGATACATCGGTGACTGGAGTTCAGA CGTGTGC	0.05

ILLPCR2_ind03	CAAGCAGAAGACGGCATAACGAGATGCCTAAGTGACTGGAGTTCAGACGTGTGC	0.05
ILLPCR2_ind04	CAAGCAGAAGACGGCATAACGAGATTGGTCAGTGACTGGAGTTCAGACGTGTGC	0.05
ILLPCR2_ind05	CAAGCAGAAGACGGCATAACGAGATCACTGTGTGACTGGAGTTCAGACGTGTGC	0.05
ILLPCR2_ind06	CAAGCAGAAGACGGCATAACGAGATATTGGCGTGACTGGAGTTCAGACGTGTGC	0.05
ILLPCR2_ind07	CAAGCAGAAGACGGCATAACGAGATGATCTGGTGACTGGAGTTCAGACGTGTGC	0.05
ILLPCR2_ind08	CAAGCAGAAGACGGCATAACGAGATTCAAGTGACTGGAGTTCAGACGTGTGC	0.05
ILLPCR2_ind09	CAAGCAGAAGACGGCATAACGAGATCTGATCGTGACTGGAGTTCAGACGTGTGC	0.05
ILLPCR2_ind10	CAAGCAGAAGACGGCATAACGAGATAAGCTAGTGACTGGAGTTCAGACGTGTGC	0.05
ILLPCR2_ind11	CAAGCAGAAGACGGCATAACGAGATGTAGCCGTGACTGGAGTTCAGACGTGTGC	0.05
ILLPCR2_ind12	CAAGCAGAAGACGGCATAACGAGATTACAAGGTGACTGGAGTTCAGACGTGTGC	0.05