

Supplementary Table S1: *meq* gene sequencing results with AH/1807 and DH/18

Stain	Sequence of meq
AH/1807	ATGTCTCAGGAGCCAGAGCCGGGCGCTATGCCCTACAGTCCCGCTGAC GATCCGTCCCCCCTCGATCTTTCTCTCGGGTCGACTTCGAGACGGAAAA AAAGGAAAAGTCACGACATCCCCAACAGCCCCTCCAAACACCCCTTC CCTGACGGCCTATCTGAGGAGGAGAAACAGAAGCTGGAAAGGAGGA GAAAAAGGAATCGTGACGCCGCTCGGAGAAGACGCAGGGAGCAGAC GTACTATGTAGACAAACTCCATGAAACATGTGAAGAGCTGCGGAGGGC CAATGAACACCTACGTAAGGAAATTCGAGATCTAAGGACTGAGTGCAC GTCCCTGCGTGACAGTTGGCTTGTCATGAGCCAGTTTGCCCTATGGCG GTACCCCTAACGGTGACCCTTGACTGCTTACCGCCCCGCACGATCCC GTTCTGAACCTCCCATTTGCACTCCTCCACCTCCCTACCGGATGAAC CTAACGCTCCACATTGCTCCGGTTCCCAACCTCCTATCTGTACCCCCGT CCTCCCGATACGGAGGAACTTTGCGCCAGCTCTGCTCGACCCCACCA CCTCCCATCTCTACTCCCCATATTATCTACGCTCCGGGGCCTTCCCCCT CCAACCTCCTATCTGTACCCCCGCTCCTCCCGATGCGGAGGAGCTTTGC GCCCAGCTCTGCTCGACCCACACCTCCCATCTGTACTCCCCATTCCC TCTTCTGCCCTCCCCAGCCTCCATCTCCGGAGGGCATCTTCCCTGCATTG TGTCCTGTTACCGAGCCGTGTACCCCTCCATCGCCGGGGACGGTTTACG CTCAGCTTTGTCTGTTGGCCAGGCTCCCTTTTTTACCCCATCTCCCCCA CATCCGGCTCCGGAGCCGGAGAGGCTTTATGCTCGTCTTACCGAGGATC CCGAACAGGATTCTTGTATTCTGGGCCAGATTTATATTCAGTTTCCCTCG GATACTCAGTCTACGGTATGGTGGTTTCCAGGTGACGGGAGACCTGA
DH/18	ATGTCTCAGGAGCCAGAGCCGGGCGCTATGCCCTACAGTCCCGCTGAC GATCCGTCCCCCCTCGATCTTTCTCTCGGGTCGACTTCGAGACGGAAAA AAAGGAAAAGTCACGACATCCCCAACAGCCCCTCCAAACACCCCTTC CCTGACGGCCTATCTGAGGAGGAGAAACAGAAGCTGGAAAGGAGGA GAAAAAGGAATCGTGACGCCGCTCGGAGAAGACGCAGGGAGCAGAC GTACTATGTAGACAAACTCCATGAAGCATGTGAAGAGCTGCAGAGGGC CAATGAACACCTACGTAAGGAAATTCGAGATCTAAGGACTGAGTGCAC GTCCCTGCGTGACAGTTGGCTTGTCATGAGCCAGTTTGCCCTATGGCG GTACCCCTAACGGTGACCCTTGACTGCTTACCGCCCCGCACGATCCC GTTCTGAACCTCCCATTTGCACTCCTCCACCTCCCTACCGGATGAAC CTAACGCTCCACATTGCTCCGGTTCCCAACCTCCTATCTGTACCCCCGT CCTCCCGATACGGAGGAACTTTGCGCCAGCTCTGCTCGACCCCACCA CCTCCCATCTCTACTCCCCATATTATCTACGCTCCGGGGCCTTCCCCCT CCAACCTCCTATCTGTACCCCCGCTCCTCCCGATGCGGAGGAGCTTTGC GCCCAGCTCTGCTCGACCCACACCTCCCATCTGTACTCCCCATTCCC TCTTCTGCCCTCCCCAGCCTCCATCTCCGGAGGGCATCTTCCCTGCATTG TGTCCTGTTACCGAGCCGTGTACCCCTCCATCGCCGGGGACGGTTTACG CTCAGCTTTGTCTGTTGGCCAGGCTCCCTTTTTTACCCCATCTCCCCCA CATCCGGCTCCGGAGCCGGAGAGGCTTTATGCTCGTCTTACCGAGGATC CCGAACAGGATTCTTGTATTCTGGGCCAGATTTATATTCAGTTTCCCTCG

	CAGTTTCCCTCGGATACTCAGTCTACGGTCTGGTGGTTTCCAGGTGACG GGAGACCCTGA
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