

A consensus linkage map of oil palm and a major QTL for stem height

May Lee, Jun Hong Xia, Zhongwei Zou, Jian Ye, Rahmadsyah, Yuzer Alfiko, Jingjing Jin, Jessica Virginia Lieando, Maria Indah Purnamasari, Chin Huat Lim, Antonius Suwanto, Limsoon Wong, Nam-Hai Chua and Gen Hua Yue

Supplementary Figure 1 Genomic sequence of the gene for asparagine synthase-related protein and SNPs detected between a *Dura* and *Pisifera* tree

- A. 1197 bp upstream DNA sequence from the start codon of the gene for asparagine synthase of oil palm. Ten SNPs (see letters in red) between the *Dura* and *Pisifera* trees were detected.

TGAGAATTATTATGGGAAAATAATATCTTGAGTCCACATCGATTGTGAGTYRAGAGAAATTYTGATTTATATGAG
ATCGAAATCTTCTCTCTTTAGTRAAAGTATCTTTATATAGGATAACATGATGAGATCCAAACYGGCACCTCGATGT
ATTTGAGGTCAAAAAAAGATAATACCTCATCAAACAATATAGAGATTGATCCATCCGATCTATGATCCAAATC
ATCCGAACCCYGATCACAATAAGAATCCATGCAAACGTTGTATTTAATCACTCGTTCA YCR TATAGATATTGAATAT
TTGACTAAAAAAGTAGAACAAAATTTTTCATTAATCTAGATAARATTTTAGATTATTATAATTTATAATGGCAAC
AGRTGATTATGAAGTATCGAAGGGTGTTCAGCTAGGRTTTTAGTATCCAAGATACTATTAGCCTTCTCTATGTTTG
TCTGTTTTGTGTTTCTTCTCCATCTTGTAATAAATAAATAAAGGGGCGATTATGGAGACAGTCCCGCAGACAGGG
CAAGGAATCCAAGAGTGCGATTTACCGTCGTCTCCGAAACGGAGGAGGTTGACGGGGTCTTCTCGGGACGTTA
GGAGAAGGATCGGACCACACATCGTCCGCCAAAAACAACCCCGAGTGGGACCCACAGTGGTCTCTTTTTCTTT
GTTTCGTTTTCTAATTTAATTTTTGCCCTCTCAAGCTAAAAATATCCCATCCGCGGGCTGGCAGATGTCGACGTG
GCACCTCGTACACAGAGATATTTTCTTAAACCAGTGGGCGGTAGCGTGGATCCGCGTCAGCGGATAAGTTCTCCAC
AGCGGCCTCCCTTCGTACCACTGCAGAAACACCGGATCTGGCCCCGATTTACGGCCCCACCCACGAGACCAAGTC
AATTGTGTGACGTAAGGCGTGAACGGTAACCACACCCGGAAGTGTTCATATCTCGCATGTCTGGAATAGCTGCGC
AATTCTAGCATGGGGAATAGGGATCCTTATCCTCCCATCCCTATCCAATTCCACCTCCAATTGAAAAATGTA
CTATACCACCGGACACA ACTATATATACACCTTCCCTATCTCTTGCCAAACTCCATCCCCTCCGTTCCCTTTCTTC
TCCTATTGCTTTACTTTTCTTGAGTTTTAGACGCAGAGAGAATAGAGAAG-Start codon

- B. 1745 bp DNA sequence from the start codon to stop codon of the gene for asparagine synthase-related protein of oil palm. Introns were labeled with green and underlined. Three SNPs in intron 1 and one SNP in exon 2 were detected between the *Dura* and *Pisifera* trees. The SNP in exon 2 did not change amino sequence. A TTC-microsatellite (see letters in pink) in intron 2 showed polymorphism between the *Dura* and *Pisifera* trees.

ATGTTGGCGATATTCCACAAGGCGTTCGCTCACCCGCCAGGAGCTCAACAGCCCCGGTGGCGGCCGGCGTGTG
CCCAAGAACCCGGAGGAGATCCTCCGGGAGTCCACTCCCTCCACCTGGGGACTCCTTCTCCGCCACCTTCAGCG

Supplementary Figure 2 Expression of the gene for asparagine synthetase-related protein in leaf (L) and trunk (T) tissues of short and tall trees of oil palm. The expression in the trunk was significantly ($P < 0.05$) higher in short trees than that in tall trees whereas the expression was similar ($P > 0.05$) in leaf in short and tall trees.

