



Figure S3. Identification of candidate genes that showed relatively lower nucleotide diversity (π) in all cultivated ginseng accessions and high genetic differentiation (F_{ST}) relative to wild ginseng. Red solid lines are the cutoff ($\pi_{wild/SCG} > 2$; $F_{ST} > 0.05$). Each green dot represents a single protein-coding gene. Barplot distributions of Tajima's D values are showed for wild and cultivated ginseng separately. X- and Y-axis are Tajima's D value and number of genes, respectively.