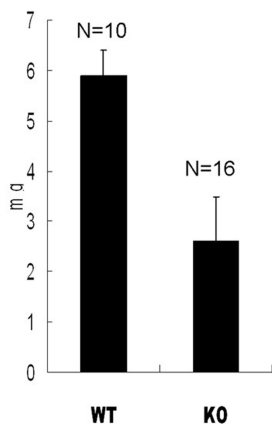
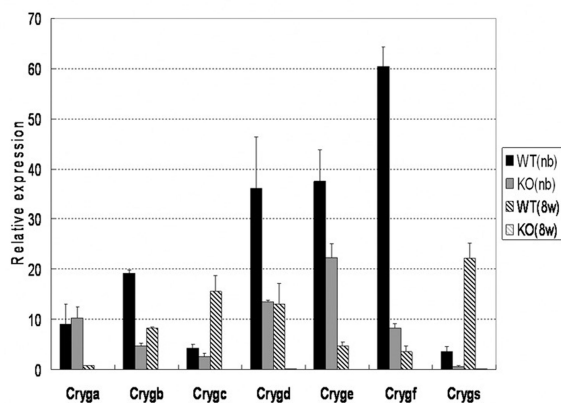


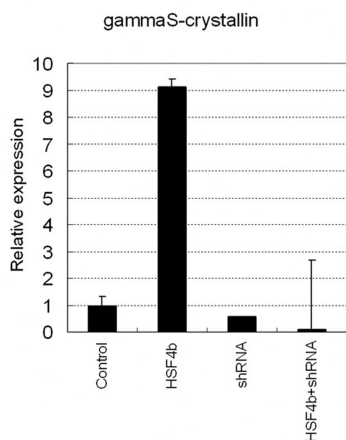
A



B



C



D

HSE : NNTTCNNGAANNITTCNNGAANNITTCNN
 * * * * *

Y A : ACAACTCGAAATGCCAGAGAGAACCGA
 -189 -163

Y B : GCAACCGAACCTACCGAAGAATGA
 -186 -160

Y C : GATCCATGAATTCCTCAGCCAGGG
 -97 -71

Y D : TCCATTCCGAAGCTCTGGAGAGTCTAA
 -176 -150

Y E : TCCATCCGAAGCTCTGGAGAGTCTAA
 -167 -140

Y F : TCCATTCCGAAGCTCTGGAGAGTCTAA
 -167 -140

Y S : CGGTACTGAAGCTTCGCTTTATTTAT
 -163 -136

E



F

