ATGCCTGAAGAAACCATGCAAGGTGATGTTGAGACCTTCGCCTTCCAGCTGAGTGTTGCTCAGCTTATGTCCCTGATTATCAACACATTCTACTCCAAC
M P E E T M Q G D V E T F A F Q A E | A Q L M S L | I N T F Y S N
AAAGAAATCTTCCTCCGTGAATTGATTTCCAACTCCTTGATGCCTTGGACAGGATCCGCTTATGAGTCGCTCACAGACCCGTCCCGCCTGGACAGTGGC K D L N I R L I P N K N D R T L T L I D T G I G M T K A D L V N N TTGGGTACCATCGCCAAGTCCGGCACTAAGGCCTTCATGGAGCTCCCAGGCTGGTGGTGATTGGTCAGTTTGGTTTGGTTTTCAT L 6 T I A K S 6 T K A F M E A L Q A G A D I S M I 6 0 F 6 V 6 F 1
TCTGCCTACCTGGTCGCCGACCGTGTCACAGTCCTCCAAGCACAACGATGACGAGCAGTACGTGTGGGAGTCCTCTGCTGGTGGTTCCTTCACCATC RPDTGEPIGRGTKIILHMKEDQMEYLEERKVKE ATTGTCAAGAACACTCCCAGTTCATCGGCTATCCAGCCCCCGCGAGAAGGAACGGAACTTAGCGAGGATGAAGAGGAGGAGGAGAGA IVKKHSQFIGYPIKLLVEKERDKELSEDEEEE GATGACAAGAAGGACAAGGAAGGAAGGTGATGATGACACCTCTAAGATAGAAGATGTTGGTGAGGAAGATGAGGAGAAAAGAAA D D K K D K E E E K K E G D D E T P K I E D V G E E D E E K E K K NAGAAAAAGAAGACAGTCAAGGAAAAGTACACTGAGGATGAAGAGTTGAACAAGACCAAGCCTATCTGGACTCGCAACCAGGACGATATTAGCCAGGAA KKKKT VKEKYTEDEELNKT KPIWT RNQD DISQE GAGTATGGTGAATTCTCACAAATCTCTGCAAGAGAGACATTCTCTGTGGAGGGTCAGCTTGAGTTCCGTGCGCTG L F V P R R A P F D L F E N K K K K N N I K L Y Y R R Y F I M D 1
TGTGAAGATCTCATCCTGAATACCTTAACTTTGTCAAGGGTGTGGTTGACAGGGAAGATCTTCCCCTCAACATTTCCCGTGAGATGCTCCAGCAGAAC C E D L I P E Y L N F V K **6 V V D S E D L P L N I S R E** M L Q Q N Aagattctgaaggtcatcaggaagaacttggtcaagaagtgcctggaggtcttcgaagagttggctgaggataaggagaactacaagaaatgctacgag K I L K V I R K N L V K K C L E L F E E L A E D K E N Y K K C Y E CAGTICITCAAAGAACCTGAAGCTGATGAGGTACCACACTCTGGCTCTGGGTGAC Q F S K N L K L G I H E D S T N R K K L A E L M R Y H T S A S G D GAAATGTGCTCTTTCAAGGAATATGTTAGCCGCATGAAGGAGAACAGAAGAACATTTACTACATCACTGGTGAGAGAGCCGTGAGCAGGTTGCCAACTCT GEM C S F K E Y V S R M K E N Q K N I Y Y I T G E S R E Q V A N S GCCTTTGTTGAGGGGTGGAGAGGTGGAGGGGTGAAGGAGTTTGATGGC K Q L V S V T K E G L E L P E D E E E K K K R E E D K A K F E N L TGCAAAGTCATGAAGGACATCCTTGACAAGAAGGTGGAGAGGTGTAGTTCTAACAGATTAGTTGAATCTCCTTGCTGTATTGTCACTTCACAGTAC C K V M K D I L D K K V E K V V V S N R L V E S P C C I V T S Q Y GOTTGGACTGCCAACATGGAACGATTGAAGGCTTAGAGGCTTAGAGCTCTAGACCATGGGTTACATGGCTGCCAAGAAGCACCATGGAAATCAAC G W T A N M E R I M K A Q A L R D T S T M G Y M A A K K H L E I N CCTGACCATGCCATGCCATGCCTTAGAAGCTTTAGAGAACCCTTAGAAACTGCTAGAAACTGCT P D H A I V E T L R Q K A E A D K N D K A V K D L V M L L F E T A CTGCTTTCATCTGGTTTCGCCCTTGAGGAGCCTGGTGTCCACGCTTCACGCATTCACTATCACTAGCTTAGGAATTGATGAAGGATGATTCACCLL S S G F A L E E P G V H A S R I H R M I K L G L G I D E D D V

Appendix A. Nucleotide and deduced amino acid sequences of *Fo-Hsp90* cDNA. The asterisk indicates the translational termination codon (TAA). The putative polyadenylation signal (AATAAA) is shown in a blue box. Five highly conserved regions representing signature amino acid sequences for Hsp90 homologs (Gupta, 1995) are shown in red. The characteristic pentapeptide MEEVD at the end of C-terminal is shown in a green box. Nucleotide and deduced amino acid sequences of *Fo-Hsp90* cDNA. The asterisk indicates the translational termination codon (TAA). The putative polyadenylation signal (AATAAA) is shown in a blue box. Five highly conserved regions representing signature amino acid sequences for Hsp90 homologs (Gupta, 1995) are shown in red. The characteristic pentapeptide MEEVD at the end of C-terminal is shown in a green box.