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HSP90.1 -----
HSP90.7 ---MRKRTLVSVLFVFLSLLFLLPDQGRKLDHAN-----AEESSDDVTDPPK
HSP90.5 MAPALSRSLYTSPLTSVPIPTVSSRLSHLRSS-----FLPHGGALRRTG
HSP90.6 MIRLSKRSLVSTLLRSGNQSERIAAAASTSRSSPSATDVKRSDESRWYSSLTNGQSKNSG

HSP90.1 -----MADVQMADETFAFQAEINQLLSLIINTFYSNKEI
HSP90.7 VEEKIGGGHGLSTDSVHRESESMKKTLRNSAEKFEFOAEVSRLMDIINSLYSNKDI
HSP90.5 VSCSNLEKRCNR-FAVKCDAVAEKETTEEGSGEKFEYQAEVSRLDLIVHSLYSHKEV
HSP90.6 SFAQLNMKTNWFMGYRNESAAASDSSSQAPPAEKFEYQAEVSRLMDLIVNSLYSNKEV

HSP90.1 FLRELISNSD DALDKIRFESLTDKSKLD--GQPELFLRLVDPKSNKTLSTIDSGIGMTKA
HSP90.7 FLRELISNASDALDKIRFALTDKDVLCGGDTAKLEIQIKLDKAKKILSIRDRGIGMTKE
HSP90.5 FLRELVSNASDALDKLRFLSVTEPSLLG--DGGDLERIKKPDNDGTITITDRGIGMTKE
HSP90.6 FLRELISNASDALDKLRFLSVTNPELSK--DAPDLDIRIYADKENGITITLTDSGIGMTRQ

HSP90.1 DLVNNLGTIARSGTKEFMFALQ---ACADVSMIGQFGVGFYSALVAEKVVVTTKHN-D
HSP90.7 DLTKNLGTIARSGTSAFVEKMQ---SSGDLNLIGQFGVGFYSALVADYIEVLSKHN-D
HSP90.5 ELIDCLGTIAQSGTSKFLKALKENKDLGADNGLIGQFGVGFYSALVAEKVVVSTKSPKS
HSP90.6 ELVDCLGTIAQSGTAKFMKALKDSDKAGGDNNLIGQFGVGFYSALVADRVVSTKSPKS

HSP90.1 DEQYVWESQAGG-SFTVTRVDVDE-PLGRGKTIITFLKDDQL-EYLEERRLKDLVKKHSE
HSP90.7 DSQYVWESKANG-KFAVSEDTWNE-PLGRGTEIRLHLRDEAG-EYLEESKLELVKRYSE
HSP90.5 DKQYVWESVADSSSYLIREETDPPNLRRGTOITLVLREDDKVEFAESTRIKLVKNYSQ
HSP90.6 DKQYVWEGEANS SSFTIQEDTDPQSLPRGTRITLHLKQEAKE-NFADPERIQKLVKNYSQ

HSP90.1 FTSYPIYLWTEKTTKEKISDDEDE--DEPKKENEGEVEVEDEKKEKDGKKK--KKIKEV
HSP90.7 FLNFPISLWASKEVETEVEVEEDESADDETETTSTEEKEEDAEEEDGKKQKTKKVKET
HSP90.5 FVGFPIYTWQEK--SRTIEVEED-----EPVKEGEEGEPKKKTKTK--TEK
HSP90.6 FVSFPIYTWQEKGYTKEVEVEDDP-----TETKKDDQDDQTEKKKTKKVVVER

HSP90.1 SHEWELINKQKPIWLRKPEEITKEEYAAFYKSLTNDWEDH--LAVKHFSVGEQLEFKAIL
HSP90.7 VYEWELLNDVKAIWLRSPKEVTEEEYTKFYHSLSKDFTEKPMWASHFNAGEGVFKAVL
HSP90.5 YWDWELANETKPIWLRNSKEVEKGEYNEFYKKAFFNEFLDP--LAHTHFTTEGEVEFRSIL
HSP90.6 YWDWELTNETQPIWLRNPKEVTTAEYNEE RKAFFNEFLDP--LASSHFTTEGEVEFRSIL

HSP90.1 FVPKRAPFDLFDTRKKNL--NIKLYVRRVFI MDNCE-ELIPEYLSFVKGVVSDDDLPLNI
HSP90.7 YVPPKAPHDLYESYVNSNKAANKLKYVRRVFISDEFD-ELLPKYLSFLKGLVDSDDLPLNV
HSP90.5 YLPGMGPLNNEVTNPKTK-NIRLYVRRVFISDDFDGELFPRYLSFVKGVVSDDDLPLNV
HSP90.6 YVPPVSPSGKDDIVNOKTK-NIRLYVRRVFISDDFDGELFPRYLSFVKGVVSDDDLPLNV

HSP90.1 SRETLQONKILKVIKKNLVKKEIEMFNETAENKED-----YTK
HSP90.7 SREMLQOHSLSKTIKKKLRKALDMLRKLAEEDPDEIHDDEKDVKEKSGENDEKKGQYTK
HSP90.5 SREILQESRIVRIMKKRLRKTDFDMIQETISESENKE-----DYKK
HSP90.6 SREILQESRIVRIMKKRLVRKAFDMI LCTLSLENRE-----DYEK

HSP90.1 FVEAFSKNLLKGIHEDSONRGKIADLLRYHSTKSGDEMTSFKDYVTRMKEGQRDIFYITG
HSP90.7 FWNDFGKSVKLGITIEDAANRNRLAKLLRFETTKSDGKLTSLDQYIKRMKKSQKDIIFYITG
HSP90.5 FWNDFGRFLKLGCIEDTGNHKRITPLLRFFSSKNEBELTSLDDYIENMNGOKAIYIYLAT
HSP90.6 FWDNFGKHLKLGCIEDRENHKRIAPLLRFFSSQSENDMISLDEYVENMKPEQKAIYFIAS

HSP90.1 ESKKAVENSFFLERLKKRGEVLYMVDALDEYAVGOLKEYDGGKLVSATKEGLKLEDETE
HSP90.7 SKKEQLEKSPFLERLIKKGVEVLIFFTDPVDEYLMQVLMQYEDKQFQNSKEGLKVKGKDS-
HSP90.5 DSLKSAKSAFLEKLIQKDI EVLYLVEPIDEVALQNLQTYKEKFFVDISKEDLELGDDEDE
HSP90.6 DSITSAKNAFLEKMLEKGL EVLYLVEPIDEVAVQSLKAYKEKDFVDISKEDLDLGDKNE

HSP90.1 EKKKREKKSFEENLCKTIKEILG-DKVEKVVVSDRIVDSPCCLVTGEYGWANMERIM
HSP90.7 ---KDKELEKFAFKELTKWKGNLASENVDDVKISNRLADTPCVVVTSKFGWSANMERIM
HSP90.5 V---KDREAKQEFNLLCDWIKQQLG-DKVAKVQVSNRLLSSSPCVLVS GKFGWSANMERIM
HSP90.6 E---KEAAVKKEFGQTCDWIKKRLG-DKVASVQISNRLSSSPCVLVS GKFGWSANMERIM

HSP90.1 KAQALRDSMSGYMSSKKTMEINPDNGIMEELRKRRAEADKNDKSVKDLVMLLYETALLTS
HSP90.7 QSOTLSDANKQAYMRGKRVLEINPRHPIIKELKDRIASDPEDSVKETAQLMYQTALIES
HSP90.5 KAQALGDTSSLEFMRGRRLLEINPDHPIIKDLNAACKNAPPESTEATRVVDLTYDTALISS
HSP90.6 KAQSTGDTTSLDYMKGRRVFEINPDHSIIKNINAAYNSNPNDDEDAMRAIDLMDALVSS

HSP90.1 GFLSDEPNTFAARIHRMLKGLS--IDEDENVEEDGDMPE-ELEEDAEEESKMEVVD----
HSP90.7 GFLITDPKDFARIYNSVKSGLN--ISPDVADEEIEAAEPEPETSEATETKSLDLAGLN
HSP90.5 GFTPDSPAELGNKIYEMMAMAVGGRWRVVEE-EESSTVNEGDDKSGETEVEVPESE----
HSP90.6 GFTPDNPAELGGKIYEMMDVALSGKWSPEVQPPQQQMAHSHDAETFEAEVPEVPE----

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Supplemental Figure 2: Amino acid sequence alignment of *Arabidopsis* cytoplasmic (HSP90.1) and organellar (HSP90.5, HSP90.6, HSP90.7) Hsp90 proteins. The peptide used in CyP40 interaction experiments is underlined in red.