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Genome-wide Analysis of the *WRKY* Family Genes and their Responses to Cold Stress in Watermelon

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Electronic Supplementary Material (ESM)

Group 1N

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AtWRKY20N : ADDGYNWRKYGQKHVKGSEFPRSYYKCTHPNCEVKKLFERSHDGQITDIIYKGTHDHPKP : 60
ClWRKY10N : SEDGYNWRKYGQKLVKGNVFVRSYYRCTHPTCMVKKQLERTHDGKITDTVYFGQHDHPK- : 59
ClWRKY13N : TEDGFNWRKYGQKVVKGSENPRSYYKCTYPNCPVRKQVERSLNGQITEIVYKSKHNHPK- : 59
ClWRKY14N : SDDGYNWRKYGQKQVKGSENPRSYYKCTFPNCPTKKKVERSLDGQITEIVYKGSHNHPK- : 59
ClWRKY17N : SEDGYNWRKYGQKQVKGSEYPRSYYKCTHSNCQVKKKVERSLDGQITEIIYKGAHNHAR- : 59
AtWRKY20N : ADDGYNWRKYGQKHVKGSEFPRSYYKCTHPNCEVKKLFERSHDGQITDIIYKGTHDHPKP : 60
ClWRKY10N : SEDGYNWRKYGQKLVKGNVFVRSYYRCTHPTCMVKKQLERTHDGKITDTVYFGQHDHPK- : 59
ClWRKY13N : TEDGFNWRKYGQKVVKGSENPRSYYKCTYPNCPVRKQVERSLNGQITEIVYKSKHNHPK- : 59
ClWRKY49N : SDDGYNWRKYGQKVVKGSEFPRSYYKCTHPNCEVKKLFERSHDGQITDIIYKGTHDHPK- : 59
ClWRKY53N : SYDGYNWRKYGQKQVKGSEYPRSYYKCTHPSCPVKKKVERSLDGKVAEIVYKGEHNHPK- : 59
AtWRKY20N : ADDGYNWRKYGQKHVKGSEFPRSYYKCTHPNCEVKKLFERSHDGQITDIIYKGTHDHPKP : 60
ClWRKY10N : SEDGYNWRKYGQKLVKGNVFVRSYYRCTHPTCMVKKQLERTHDGKITDTVYFGQHDHPK- : 59
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Group 1C

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AtWRKY20C : -DDGYRWRKYGQKVVRGNPNPRSYYKCTAHGCPVRKHVERASHDPKAVITTYEGKHDHDVP : 60
ClWRKY10C : VNDGYRWRKYGQKLVKGNPNPRSYYRCSSPGCPVKKHVERASYDPKVVLTTYEGQHDHDM- : 60
ClWRKY13C : LDDGYWWRKYGQKVVKGNSNPRSYYKCTYAGCGVRKHIERASHDIRAVITTYEGKHNHEV- : 60
ClWRKY14C : LDDGYRWRKYGQKVVKGNPNPRSYYKCTNPGCPVRKHVERASHDLRAVITTYEGKHNHDV- : 60
ClWRKY17C : LEDGYRWRKYGQKVVKGNPNPRSYYKCTSAGCLVRKHVERASHDLKCVITTYEGKHNHEV- : 60
ClWRKY29C : LDDGYRWRKYGQKIVKGNPYPRSYYKCTTPGCNVRKHVERASTDPKAVITTYEGKHNHDV- : 60
ClWRKY32C : SGDGYRWRKYGQKMVKGNPHPRNYYRCTSAGCPVRKHIESAVENPNAVIITYKGVHDHDM- : 60
ClWRKY48C : LDDGYRWRKYGQKVVKGNPNPRSYYKCTNPGCTVRKHVERASHDLKSVITTYEGKHNHDV- : 60
ClWRKY49C : LDDGYRWRKYGQKVVRGNPNPRSYYKCTNVGCPVRKHVERASHDPKAVITTYEGKHNHDV- : 60
ClWRKY53C : SGKGIRWRKYGQKVVKGNLYPRSYYRCTGLKCKARKYVERASEDPDSFITTYEGKHNHGI- : 60
ClWRKY54C : LPDGYRWRKYGQKVVKGNPNPRSYYKCTSLGCPVRKHIERAANDMRAVITTYEGKHNHEV- : 60
ClWRKY57C : LDDGYRWRKYGQKVVKGNPNPRSYYKCTSAGCNVRKHVERSSTDSKAVVTTYEGKHNHDV- : 60
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Group 2a

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AtWRKY40 : VKDGYQWRKYGQKVTRDNPSPRAYFKCACAPSCSVKKKVQRSVEDQSVLVATYEGEHNHPMP : 62
ClWRKY27 : VKDGYQWRKYGQKVTRDNPSPRAYFKCSSAPNCPVKKKVQRSLEDPTILVATYEGEHSHAS- : 61
ClWRKY43 : VKDGYQWRKYGQKVTKDNPSPRAYYKCSFAPSCPVKRKVQRSVEDPSYLIATYEGEHNHAK- : 61
ClWRKY52 : VKDGFQWRKYGQKVTRDNPCPRAYFKCSFAPSCPVKKKVQRSVEDQSVLVATYEGEHNHPH- : 61
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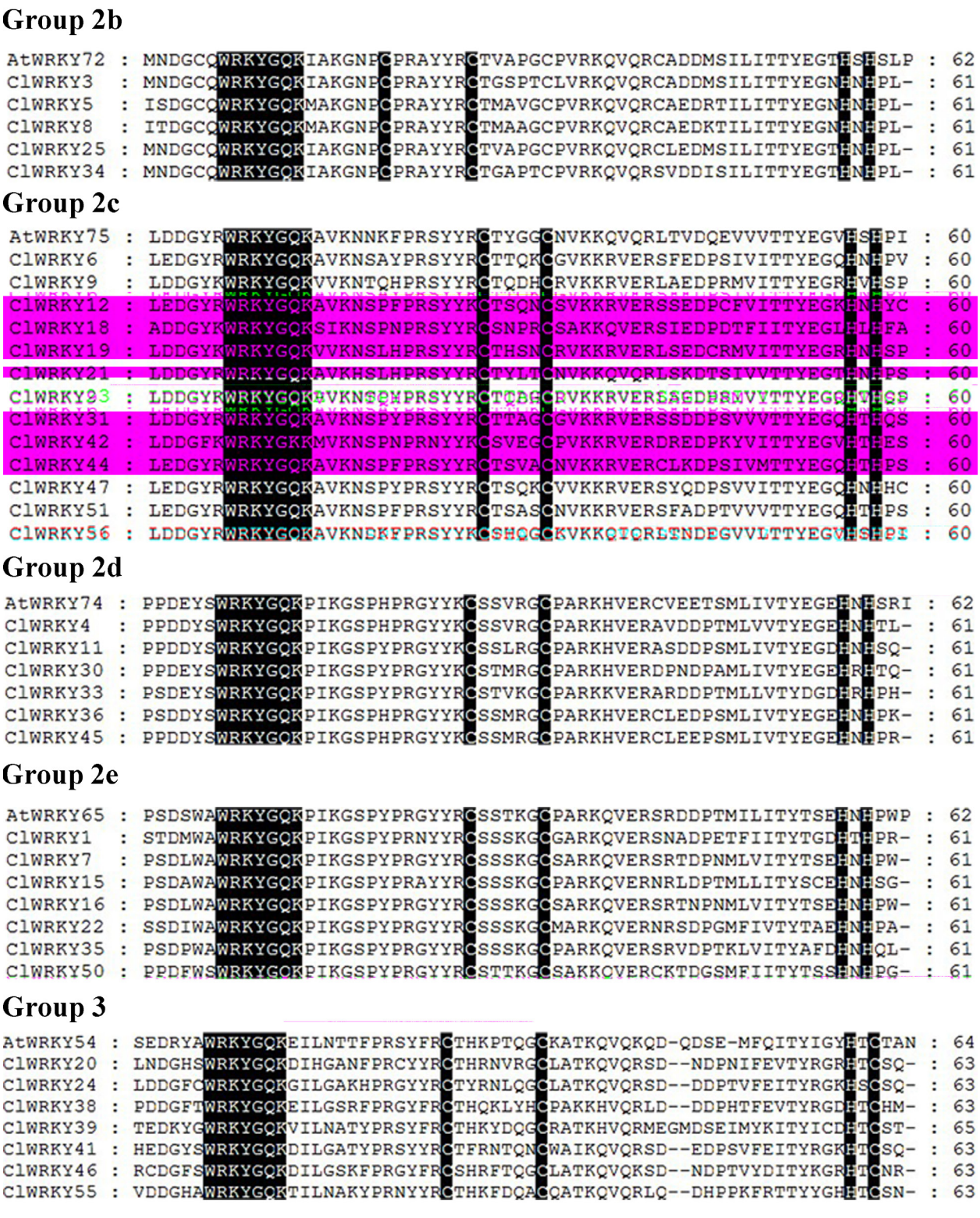

Figure S2. Alignment of the WRKY domains from multiple ClWRKY and selected AtWRKY transcription factors; 1C and 1N indicate the N- or the C-terminal WRKY domain of Group 1 WRKYs