95 S SVPSQKTY-Q
92 SFVPSQKTY-Q
64 STVPETSDY-P
69 CAVPSTDDY-
163 PAIPSNTDY-P
113 PVIPSNTDY-P
188 NTLPKLENHN
*

Figure S4. Sequence alignment of DNA-binding domains of p53 family members used in this study. DNA-contact residues in Hsp 53 are marked with an asterisk, DBD interface residues with a hash. The flexible loop around the DNA-contact residue K120 is highlighted with a black bar. Colour coding is according to ClustalX [1].

1. Thompson JD, Gibson TJ, Plewniak F, Jeanmougin F, Higgins DG: The CLUSTAL_X windows interface: flexible strategies for multiple sequence alignment aided by quality analysis tools. Nucleic Acids Res 1997, 25(24):4876-4882.
