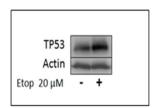
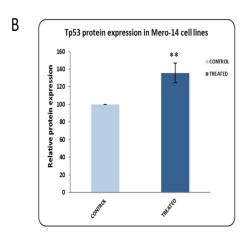
Α





P53 protein levels and stabilisation induced by DNA damage in Mero-14 cells A. Detection of the Total p53 protein levels in the Mero-14 cell lines with and without DNA damage by using SDS PAGE and Western Blotting. The cell lines were incubated for 24 hrs with or without 20 μ M etoposide. B. Quantification of the wild-type p53 protein levels in the Mero-14 shown in Figure S2A, which was generated by using ImageJ software. Data is representative of three experiments; band intensity was normalised to β -actin.

^{*}indicates p<0.05.

(C) Sequence alignment of the coding sequence of the TP53 gene (exons 1–11) from Mero-14 cell line, as described in Methods. TP53, RefSeqNG 017013.2 P53,NC 000017Exon1 Mero14,P53Exon1 TP53, RefSeqNG 017013.2 P53,NC 000017Exon1 Mero14.P53Exon1 ······ 5030 5040 5020 TP53, RefSeqNG 017013.2 GATGGGATTGGGGTTTTCCCCTCCCATGTGCTCAAGACTGGCGCTAAAAGTTTTGAGCTT P53,NC 000017Exon1 Mero14,P53Exon1 TP53, RefSeqNG 017013.2 P53,NC 000017Exon1 Mero14,P53Exon1 5230 TP53, RefSeqNG 017013.2 TCCTCTCTGAGTCACGGGCTCTCGGCTCCGTGTATTTTCAGCTCGGGAAAATCGCTGGGG P53,NC 000017Exon1 Mero14,P53Exon1 15910 15920 15930 15940 15950 15960 1 TP53, RefSeqNG 017013.2 TGCTGGATCCCCACTTTTCCTCTTGCAGCAGCCAGACTGCCTTCCGGGTCACTGCCATGGAGGAGCCGCA P53,NC 000017Exon2 -........... Mero14, P53Exon2 16050 16040 16010 16020 16030 TCAGGAAACATTTTCAGACCTATGGAAACTGTGAGTGGATCCATTGGAAGGGCAGGCCCACCACCCCCAC TP53, RefSeqNG 017013.2 P53,NC 000017Exon2 Mero14, P53Exon2 16120 TP53, RefSeqNG 017013.2 P53,NC 000017Exon2 Mero14, P53Exon2

16120

TP53, RefSegNG 017013.2 P53,NC_000017Exon3 Mero14, P53Exon3

3120 16130 16140 16150

GGGAAGCGAAAATTCCATGGGACTGACTTTCTGCTCTTGTCTTTCAGACTTCCTGAAAACAACGTTCTGG

16150

TP53,RefSeqNG_017013.2 P53,NC_000017Exon4 Mero14,P53Exon4	16310 16320 16330 16340 16350
TP53,RefSeqNG_017013.2 P53,NC_000017Exon4 Mero14,P53Exon4	16410 16420 16430 16440 16450
TP53,RefSeqNG_017013.2 P53,NC_000017Exon4 Mero14,P53Exon4	16510 16520 16530 16540 16550
TP53,RefSeqNG_017013.2 P53,NC_000017Exon4 Mero14,P53Exon4	16610 16620 16630 16640 16650
TP53,RefSeqNG_017013.2 P53,NC_000017Exon5,6 Mero14,P53Exon5,6	17310 17320 17330 17340 17350
TP53,RefSeqNG_017013.2 P53,NC_000017Exon5,6 Mero14,P53Exon5,6	17410 17420 17430 17440 17450
TP53,RefSeqNG_017013.2 P53,NC_000017Exon5,6 Mero14,P53Exon5,6	17510 17520 17530 17540 17550 GAGCAGCTGGGGCTGGAGAGACGACAGGGCTGGTTGCCCAGGGTCCCCAGGCC
TP53,RefSeqNG_017013.2 P53,NC_000017Exon5,6 Mero14,P53Exon5,6	17610 17620 17630 17640 17650 TTATCCGAGTGGAAGGAAATTTGCGTGTGGAGTATTTGGATGACAGAAACACT
TP53,RefSeqNG_017013.2 P53,NC_000017Exon5,6 Mero14,P53Exon5,6	17710 17720 17730 17740 17750

	18210 18220 18230 18240 18250 18260 182
TP53,RefSeqNG 017013.2	GCTTGCCACAGGTCTCCCCAAGGCGCACTGGCCTCATCTTGGGCCTGTGTTATCTCCTAGGTTGGCTCTG
P53,NC_000017Exon7	
Mero14,P53Exon7	
	18310 18320 18330 18340 18350 18360 183
TP53, RefSeqNG_017013.2	GTAACAGTTCCTGCATGGGCGGCATGAACCGGAGGCCCATCCTCACCATCATCACACTGGAAGACTCCAG
P53,NC_000017Exon7 Mero14,P53Exon7	
TP53,RefSeqNG_017013.2 P53,NC_000017Exon7 Mero14,P53Exon7	18410 18420 18430 18440 18450 18460 184
	GCCTGCTGTGCCCCAGCCTCTGCCTCTGACCCCTGGGCCCACCTCTTACCGATTTCTTCCATACTA
	18710 18720 18730 18740 18750 187
TP53,RefSeqNG_017013.2 P53,NC_000017Exon8,9 Mero14,P53Exon8,9	CTATCCTGAGTAGTGGTAATCTACTGGGACGGAACAGCTTTGAGGTGCGTGTTTGTGCCT
METOT4, F33EXONO, 9	
	18810 18820 18830 18840 18850 188
TP53,RefSeqNG_017013.2 P53,NC 000017Exon8,9	CAAGAAAGGGGAGCCTCACCACGAGCTGCCCCCAGGGAGCACTAAGCGAGGTAAGCAAGC
Mero14,P53Exon8,9	
TP53,RefSeqNG_017013.2 P53,NC_000017Exon8,9 Mero14,P53Exon8,9	18910 18920 18930 18940 18950 189
	CCTCAGATTCACTTTTATCACCTTTCCTTGCCTCTTTCCTAGCACTGCCCAACAACACCA
	19010 19020 19030 19040 19050 190
	19010 19020 19030 19040 19050 190
TP53,RefSeqNG_017013.2 P53,NC_000017Exon8,9 Mero14,P53Exon8,9	ATATTTCACCCTTCAGGTACTAAGTCTTGGGACCTCTTATCAAGTGGAAAGTTTCCAGTC
	21810 21820 21830 21840 21850 21860 218
MDE2 D-50NG 017012 0	**************************************
TP53,RefSeqNG_017013.2 P53,NC 000017Exon10	ATATACTTACTTCTCCCCCTCTCTGTTGCTGCAGATCCGTGGGCGTGAGCGCTTCGAGATGTTCCGAGA
Mero14,P53Exon10	
	21910 21920 21930 21940 21950 21960 219
TP53,RefSeqNG 017013.2	CCAGGCTGGGAAGGAGCCAGGGGGGAGCAGGGCTCACTCCAGGTGACCTCAGCCCCCTTCCTGGCCC
P53,NC_000017Exon10	CCAGGCTGGGAAGGAGCAGGGGAGCAGGGGCTCACTCCAGGTGAGTGA
Mero14,P53Exon10	
	22010 22020 22030 22040 22050 22060 220
TP53,RefSeqNG 017013.2	ATAGGATTCCATCCTGCCTTCATGGTCAAAGGCAGCTGACCCCATCTCATTGGGTCCCAGCCCT
P53,NC_000017Exon10	
Mero14,P53Exon10	

TP53,RefSeqNG_017013.2	22710 22720 22730 22740 22750 22760 227
P53,NC_000017Exon11 Mero14,P53Exon11	
TP53,RefSeqNG_017013.2 P53,NC_000017Exon11 Mero14,P53Exon11	22810 22820 22830 22840 22850 22860 228
TP53,RefSeqNG_017013.2 P53,NC_000017Exon11 Mero14,P53Exon11	22910 22920 22930 22940 22950 22960 229
TP53,RefSeqNG_017013.2 P53,NC_000017Exon11 Mero14,P53Exon11	23010 23020 23030 23040 23050 23060 230
TP53,RefSeqNG_017013.2 P53,NC_000017Exon11 Mero14,P53Exon11	23110 23120 23130 23140 23150 23160 231
TP53,RefSeqNG_017013.2 P53,NC_000017Exon11 Mero14,P53Exon11	23210 23220 23230 23240 23250 23260 232