Supporting Information

for

Site-Selective Tyrosine Phosphorylation in the Activation of the p50 Subunit of NF-KB for DNA Binding and Transcription

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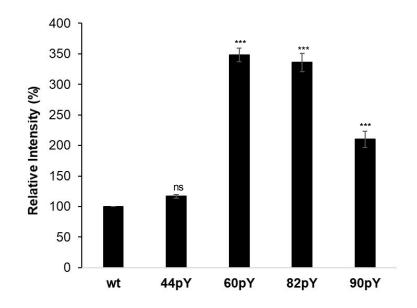


Figure S1. Statistical analysis of IL-2 promoter DNA binding of wild-type NF- κ B in comparison with each modified NF- κ Bs. Statistical significance was calculated using the Student's t-test: p < 0.001 (***), p > 0.05 (not significant, ns).

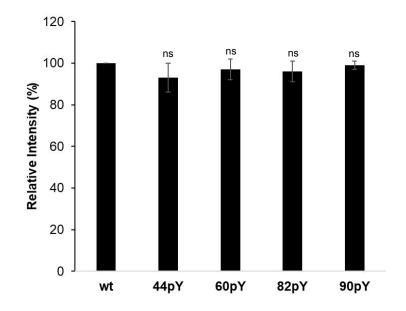


Figure S2. Statistical analysis of IL-2 promoter DNA binding of wild-type p50 in comparison with each modified p50. Statistical significance was calculated using the Student's t-test: p > 0.05 (not significant, ns).

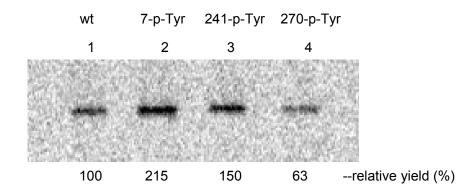


Figure S3. Cell free expression of modified p50 subunits of NF- κ B containing phosphorylated tyrosine at positions 7, 241 and 270, each synthesized by suppression of nonsense codon UAG in an *E. coli* protein biosynthesizing system using phosphotyrosyl-tRNA_{CUA}. Visualization was accomplished using the ³⁵S-methionine incorporated into the proteins. Following *in vitro* synthesis, each protein was purified on a Strep-tactin column. The purified proteins were analyzed by 15% SDS-PAGE. The isolated yields of the proteins relative to wild type are listed below each lane.

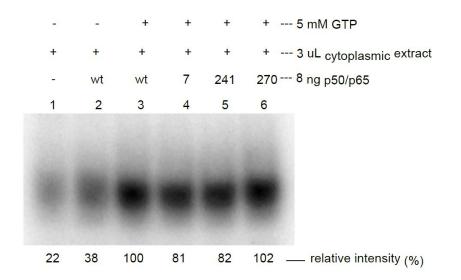


Figure S4. DNA binding assay of modified NF- κ Bs expressed in a cell free *E. coli* protein biosynthesizing system following treatment with a cytoplasmic lysate from activated Jurkat cells in the presence GTP. The cytoplasmic lysatewas incubated at 37 °C for 6 h. The extract was then cooled on ice and combined with cold samples of NF- κ B and IL-2 promoter DNA. The incubation was carried out at 4 °C for 10 min. Controls included the cytoplasmic extract alone (lane 1), wild-type NF- κ B in the absence and presence of GTP (lanes 2 and 3), and modified NF κ Bs containing p50 subunits having pTyr at positions 7, 241 or 270 (lanes 4, 5 and 6, respectively). The analysis of samples was carried out by 5% native PAGE.

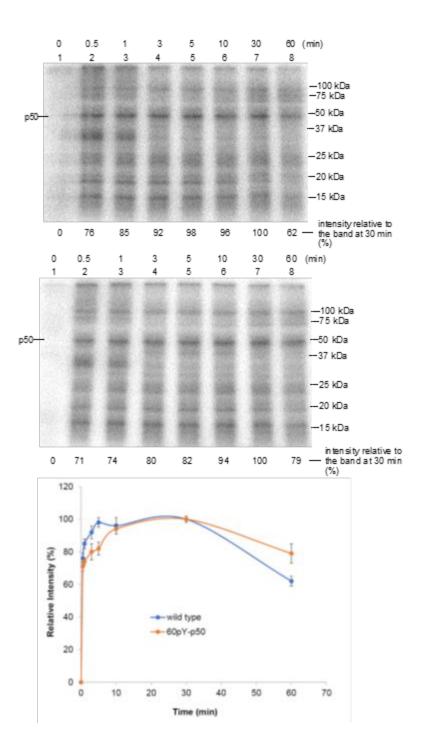


Figure S5. Phosphorylation of wild-type p50 (upper panel) and modified p50 (containing pY60) (middle panel) in the presence of Jurkat cell extracts. The *in vitro* expressed p50 proteins (80 ng) were added into the solution containing 15 μ L of activated Jurkat cell extract in the presence of 80 nM γ -³²P-GTP (6000 Ci/mmol). The reaction mixture was incubated at room temperature. The results were analyzed by 15% SDS-PAGE. 0 min: no Jurkat cell extract was added; 0.5 – 60 min after Jurkat cell extract addition. (lower panel) Time course of phosphorylation of the wild-type and modified p50s in the presence of cell extract from activated Jurkat cells. The band at 30 min was defined as 100%.

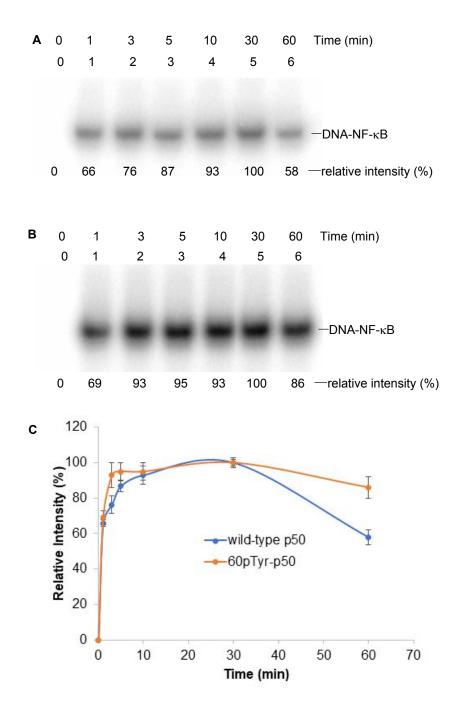


Figure S6. Time dependent DNA binding of phosphorylated wild-type p50 (**A**) and modified p50 (**B**, p50 containing pY60) in the presence of Jurkat cell extracts. The *in vitro* expressed p50 protein (8 ng) was mixed with 10 ng of wild-type p65 protein and added into $(1.5 \,\mu\text{L})$ extract from activated Jurkat cells in the presence of 80 nM GTP. The reaction mixture was incubated at room temperature. At the determined time, ³²P-labeled DNA (IL-2, 2 pmol) was added to the reaction mixture, which was incubated at room temperature. The results were analyzed using 5% native PAGE. 0 min: no activated Jurkat cell extract was added; 1 - 60 min after Jurkat cell extract addition. (**C**) Time course of binding of the wild-type and modified p50s to IL-2 promoter DNA in the presence of cell extract from activated Jurkat cells. The band having the highest intensity for each protein was defined as 100.

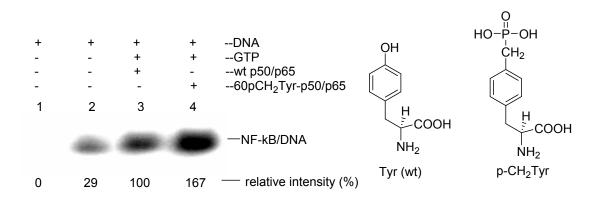


Figure S7. The *in vitro* expressed wild-type or modified (60-pCH₂Tyr) p50 protein (8 ng) was combined with 10 ng of wild-type p65 protein and was preincubated with 1.5 μ L of lysate from activated Jurkat cells in the presence of 5 mM GTP. ³²P-labeled IL-2 DNA (2 pmol) was added to the reaction mixture, which was incubated at 0 °C for 5 min. The results were analyzed using 5% native PAGE.