

Supporting Information

Modulating the Folding and Binding of Peptides

Using a Stimuli-Responsive Molecular Tweezer

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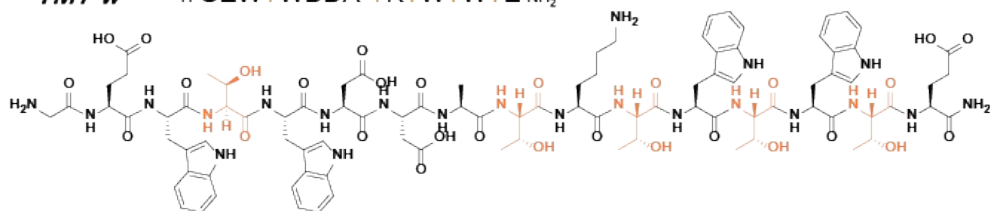
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KEYWORDS: β -hairpin-based molecular tweezer, Stimulus-responsive peptide folding, α -helix stabilization, Controlled capture-and-release, Multivalent binding

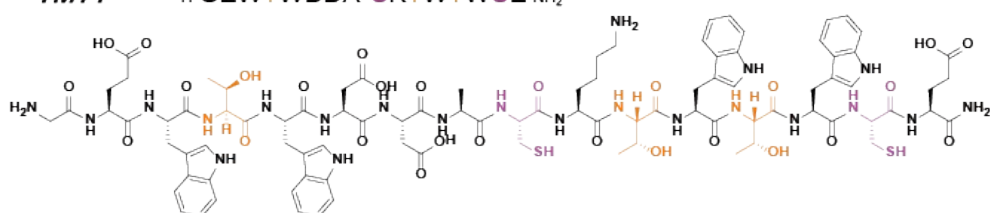
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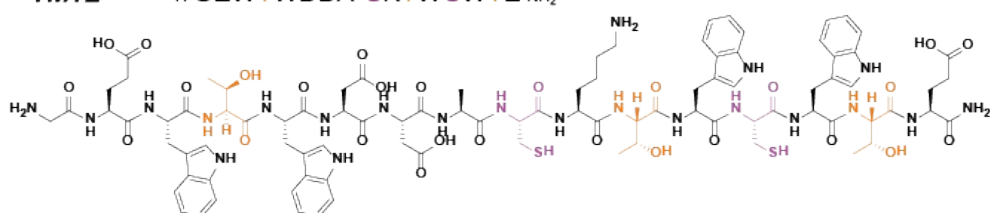
TMT-w H-GEWTWDDA-TKTWTWTE-NH₂



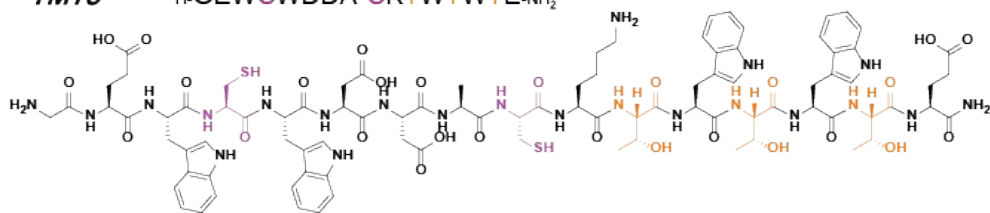
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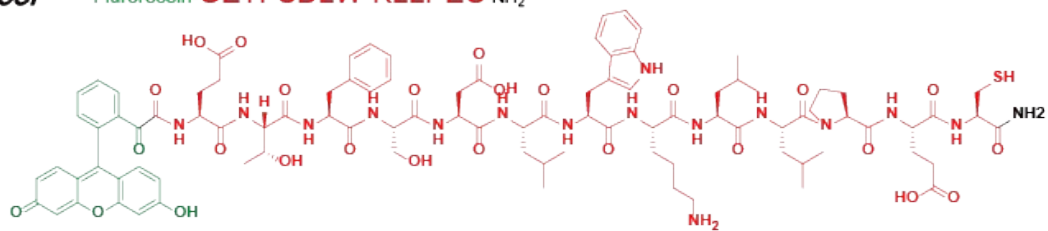
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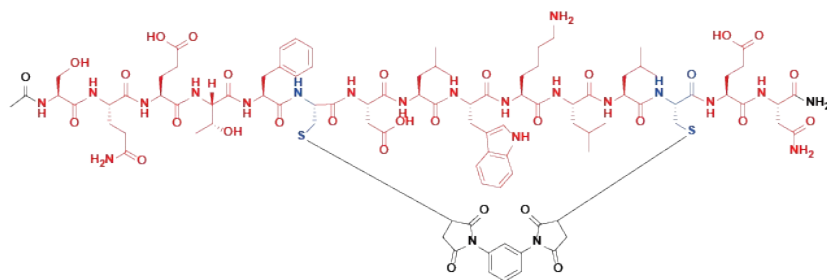
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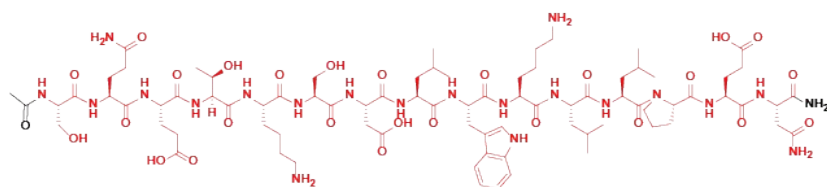
p53f Fluorecein-GETFSDLW-KLLPEC-NH₂

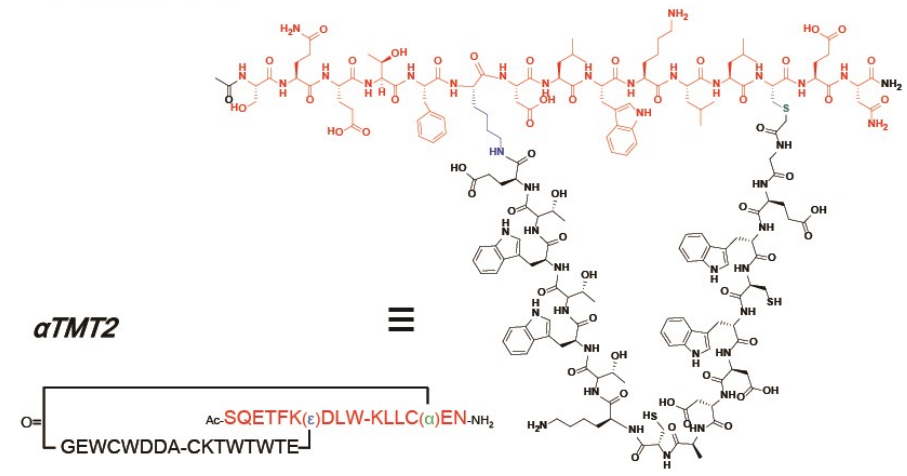
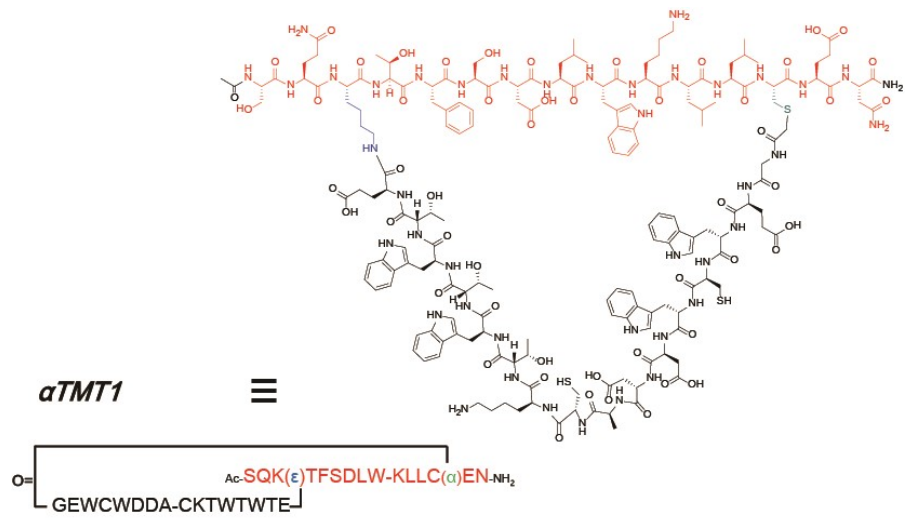


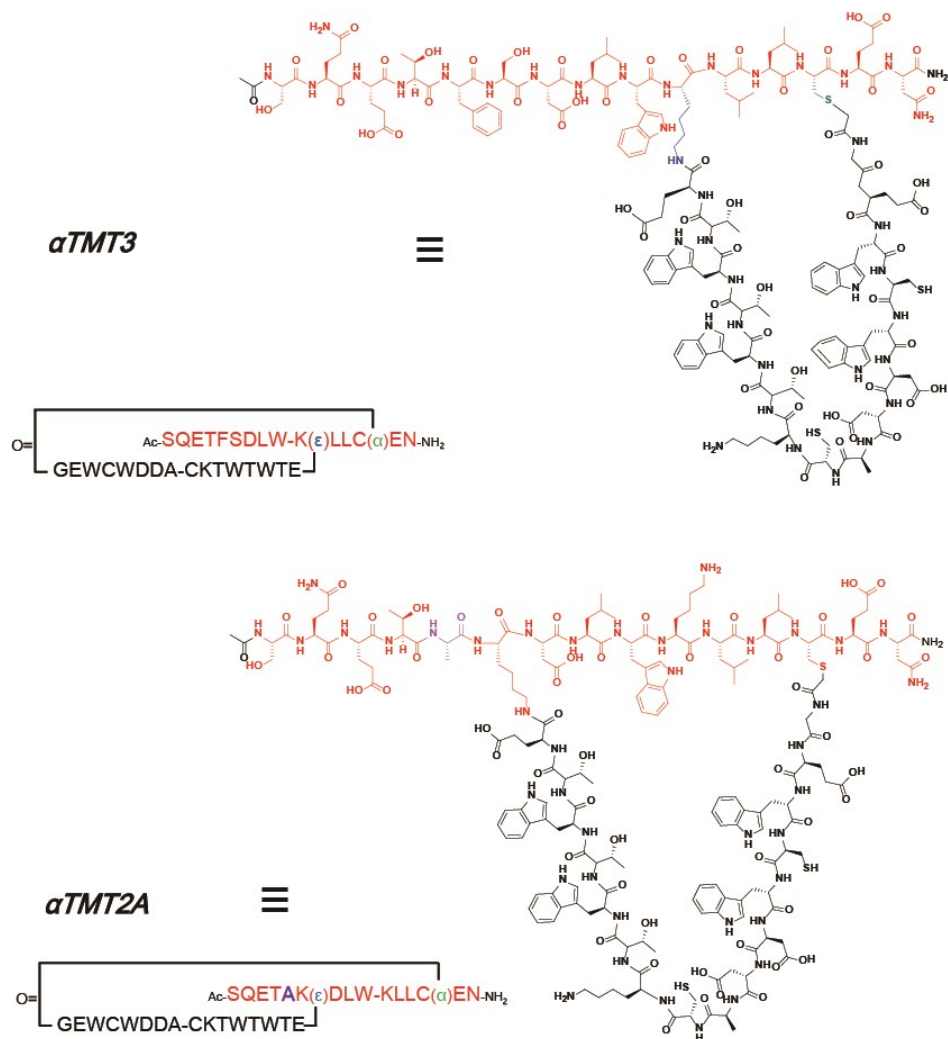
Stapld p53a Ac-SQETFCDLW-KLLCEN-NH₂



Native unstapled p53a Ac-SQETFSDLW-KLLPEN-NH₂







hDM2

histag TEV cleavage site
 HHHHHHSSGENLYFQSQIPASEQETLVRPKPLLLKLLKSVGAK
 DTYTMKEVLFYLGQYIMTKRLYDEKQQHIVYCSNDLLGLDFGVPS
 FSVKEHRKIYTMIRNLVVVNQQESSDSGTSVSEN

Fig. S1 Chemical structures of peptides and sequence of the expressed hDM2 proteins.

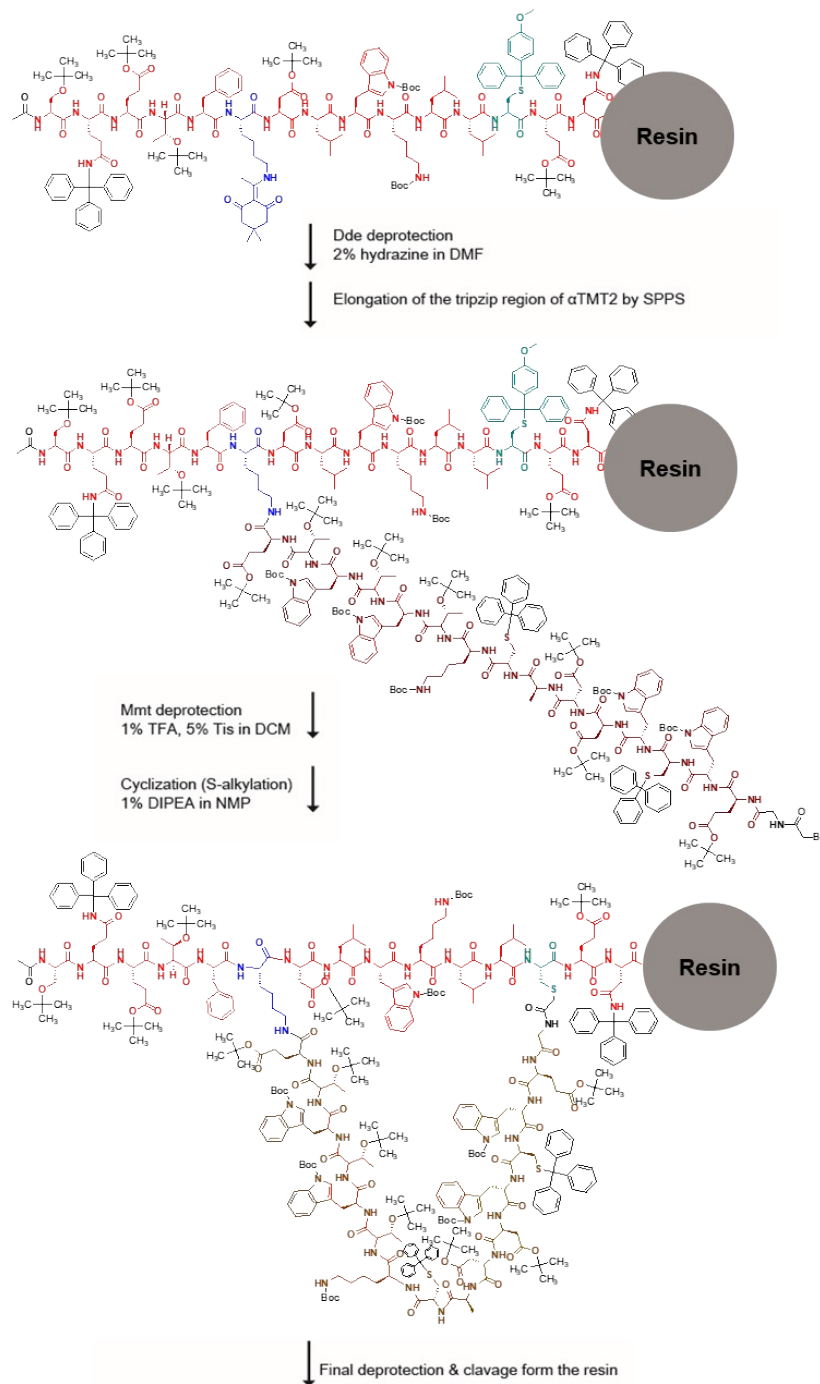


Fig. S2 Reaction scheme of the α TMT2 synthesis.

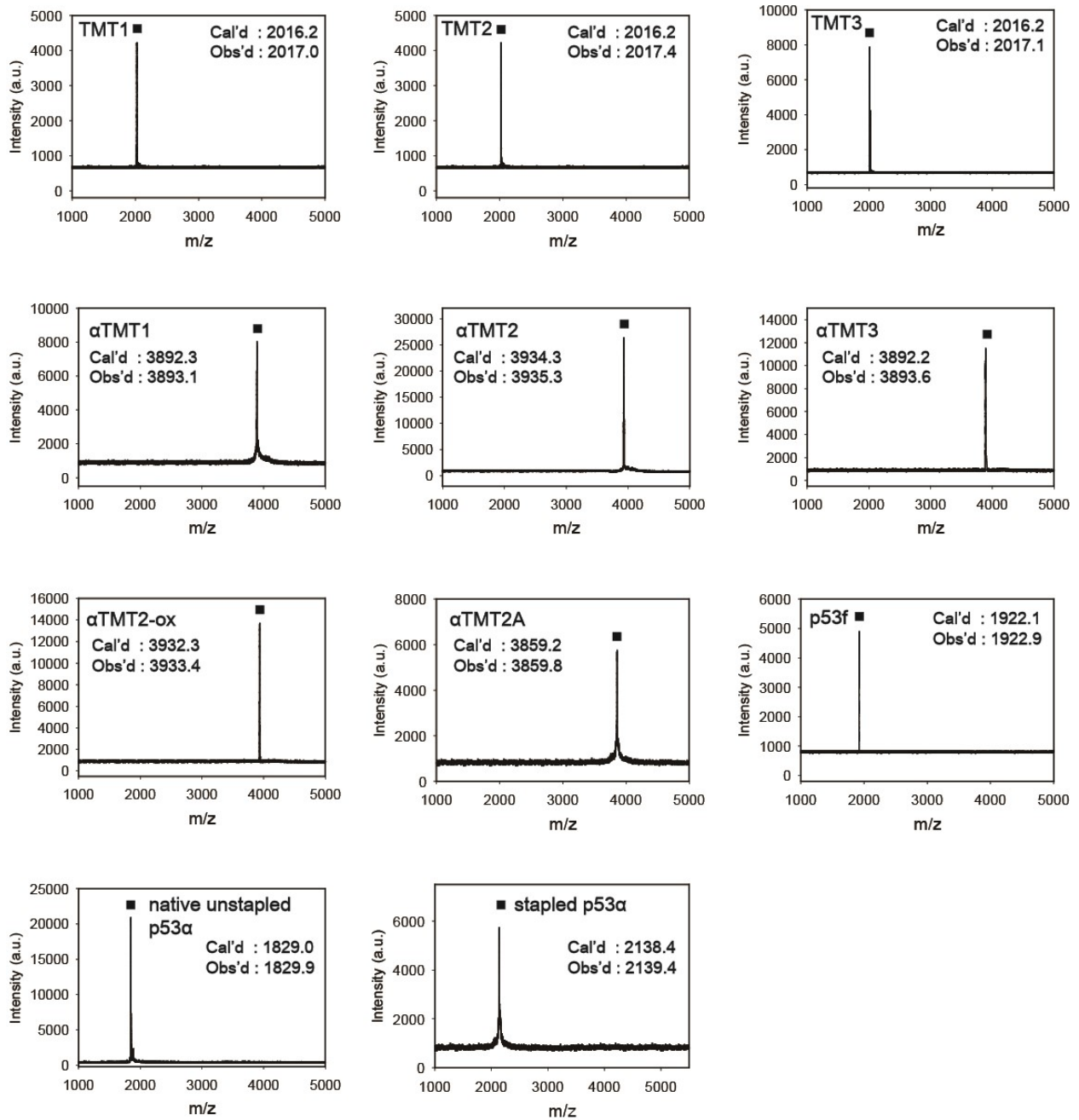


Fig. S3 MALDI-TOF MS spectra of peptides.

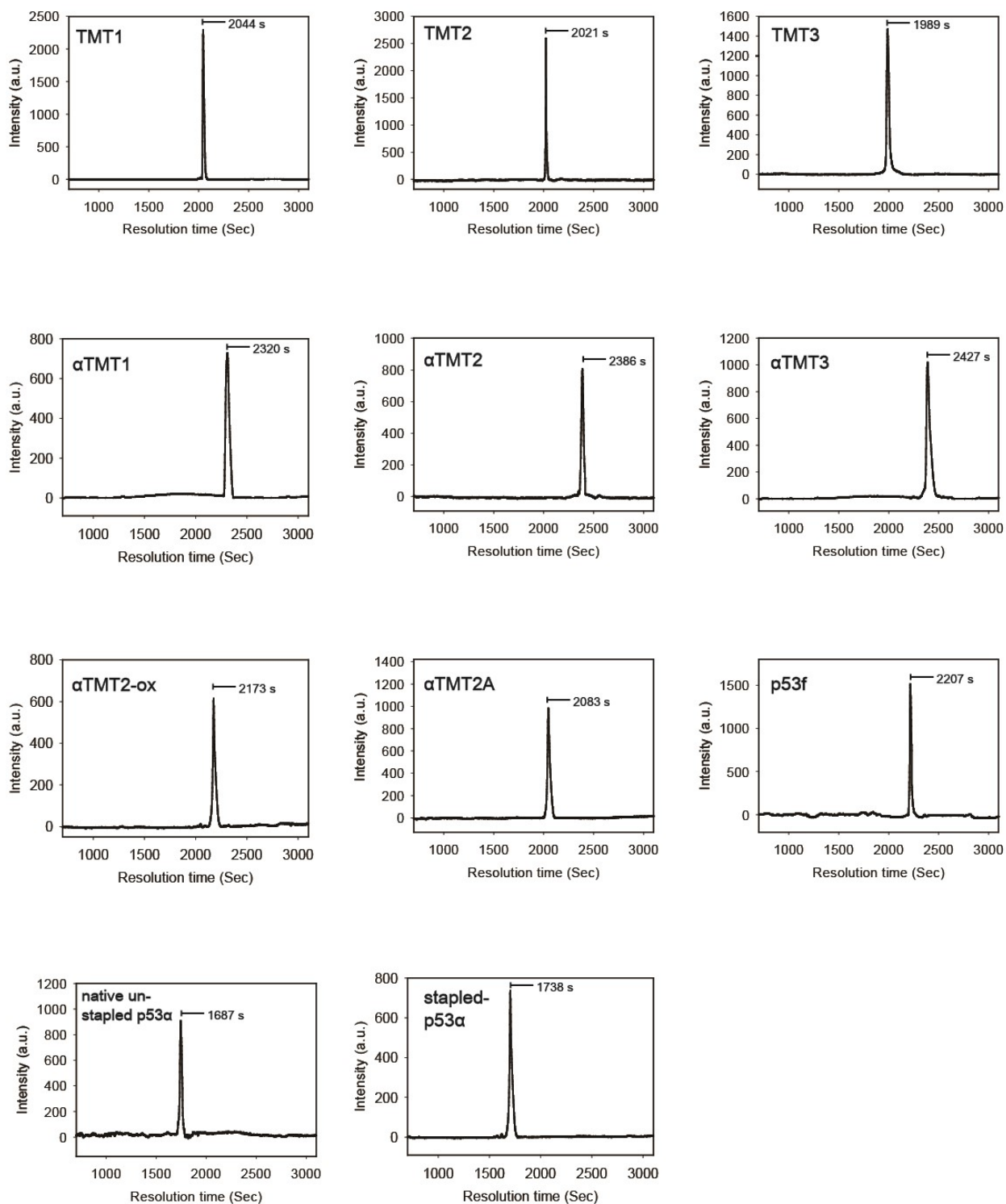


Fig. S4 HPLC chromatograms of peptides. Reverse-phase (RP) HPLC was performed using a Vydac C18 column (10 × 250 mm, 300 Å) with a gradient of 30% to 50% MeCN with 0.1% TFA.

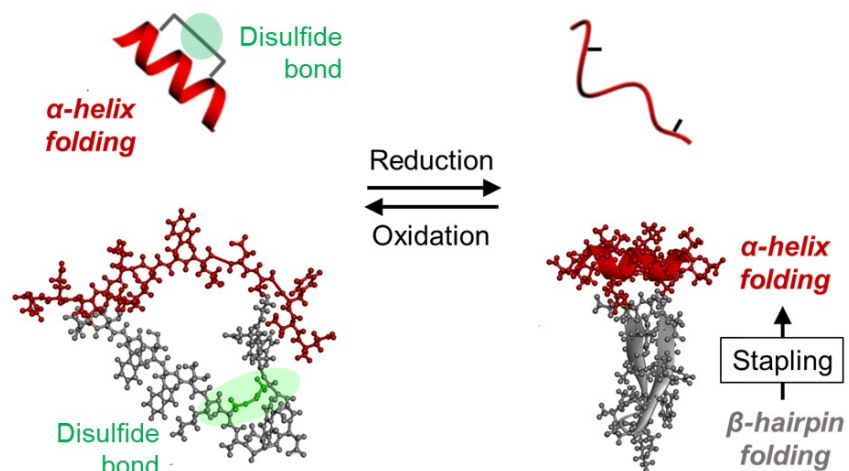


Fig. S5 Schematic illustration of redox-responsive α -helix stabilization behaviors of the typical disulfide stapling (top) and α TMT (bottom) strategies.

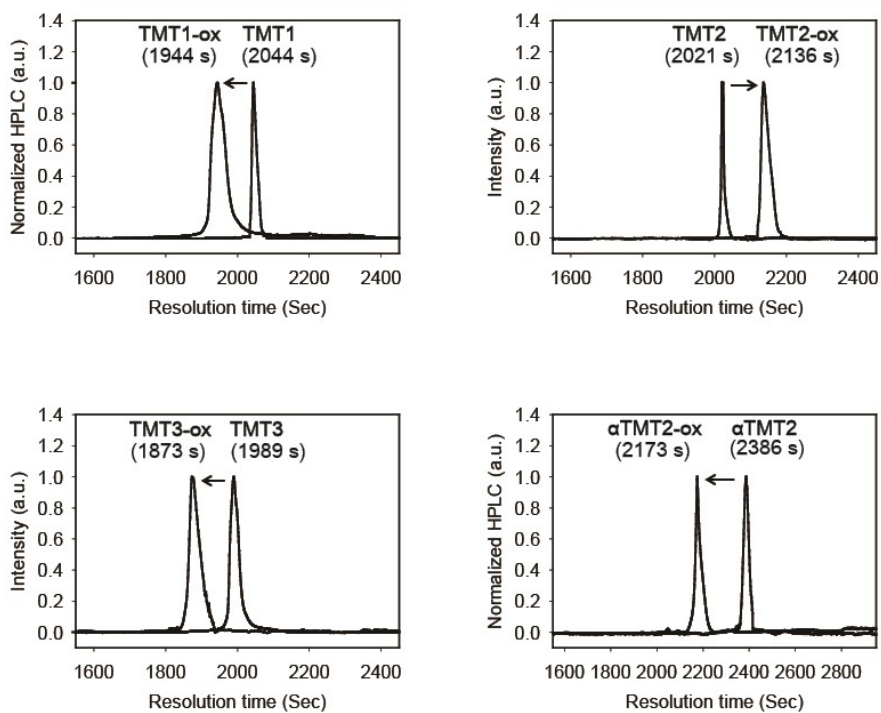


Fig. S6 HPLC peak shift after oxidation (disulfide bond formation) of peptides.

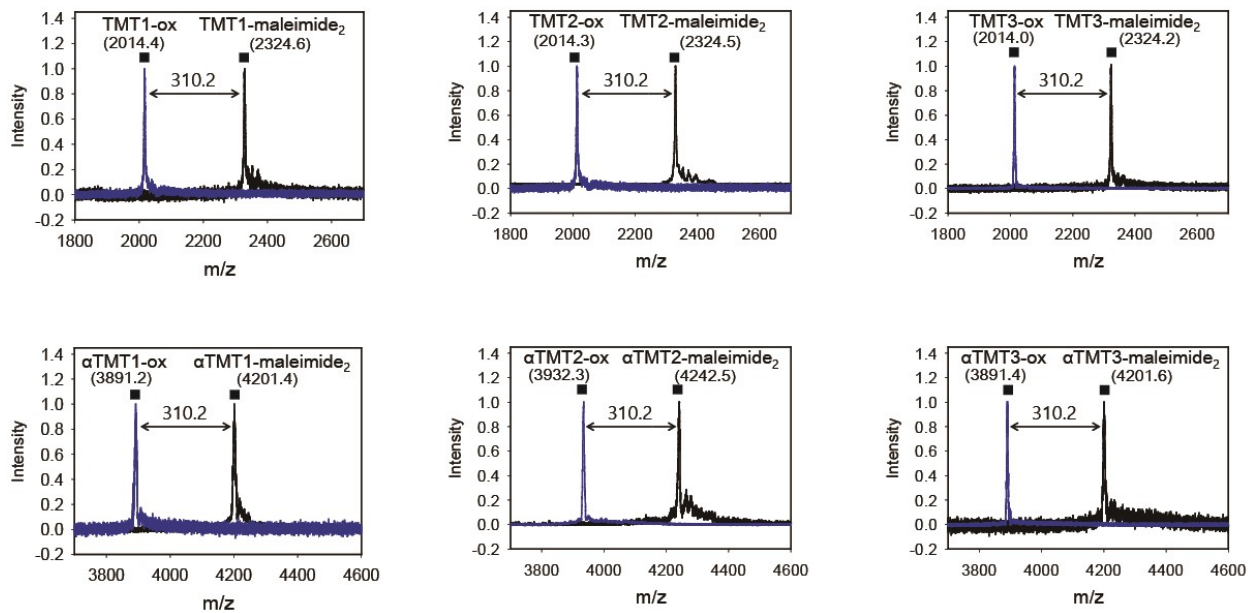


Fig. S7 MALDI-TOF MS spectra of peptides and their conjugates with n-methoxycarbonyl maleimide (MW: 155.11 Da).

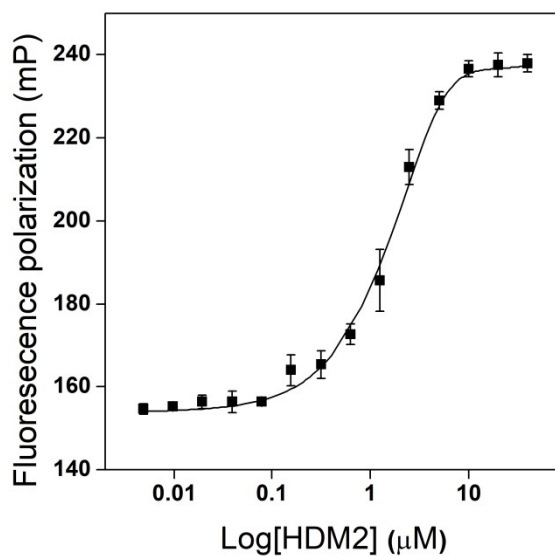


Fig. S8 Changes in the FP intensity of p53f (100 nM) upon titration with hDM2 in Tris-Cl buffer solution (pH 8.0, 20 mM, 10 mM imidazole, 150 mM NaCl, 10% glycerol) at room temperature. The filter sets were 485/15 excitation and 535/25 emission.

Peptides	α -Helix Sequence	EC50 (μ M)	K _d (μ M)
Nutlin-3a	-	0.154	0.145
α TMT1	Ac-SQK(ϵ)TFSDLWKLLC(α)EN-NH ₂	0.547	0.515
α TMT2	Ac-SQETFK(ϵ)DLWKLLC(α)EN-NH ₂	0.149	0.140
α TMT3	Ac-SQETFSDLWK(ϵ)LLC(α)EN-NH ₂	0.655	0.616
α TMT2A	Ac-SQETAK(ϵ)DLWKLLC(α)EN-NH ₂	4.584	4.314
α TMT2-ox	Ac-SQETFK(ϵ)DLWKLLC(α)EN-NH ₂	0.569	0.536
p53 α -stapled	Ac-SQETFCDLWKLLCEN-NH ₂	0.343	0.323

Fig. S9 Calculated EC50 and dissociation constant (K_d) values for the p53f/hDM2 complexes with the investigated peptides.

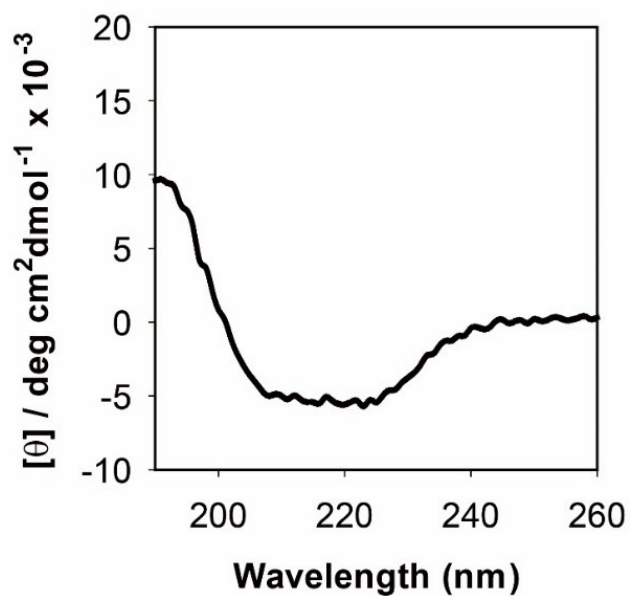


Fig. S10 CD spectrum of the stapled p53 α peptides (20 μ M) in PBS (pH 7.0, 20 mM, 150 mM NaCl).

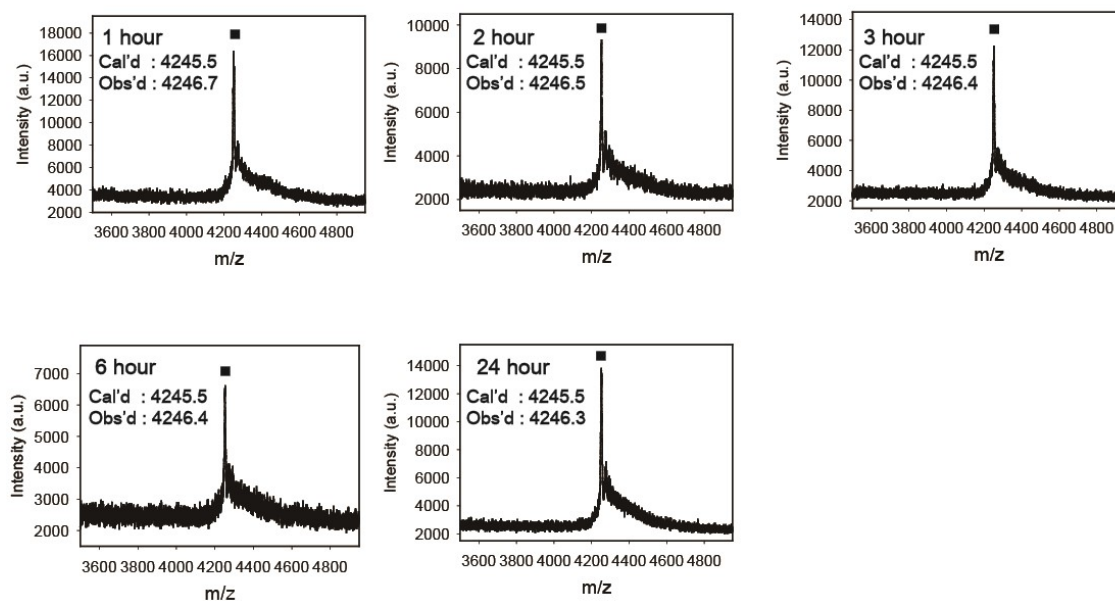


Fig. S11 MALDI-TOF analysis of the α TMT2 peptides after incubation in PBS (pH 7.0, 20 mM, 150 mM NaCl) for different times at room temperature and n-methoxycarbonyl maleimide (2 mM) coupling.

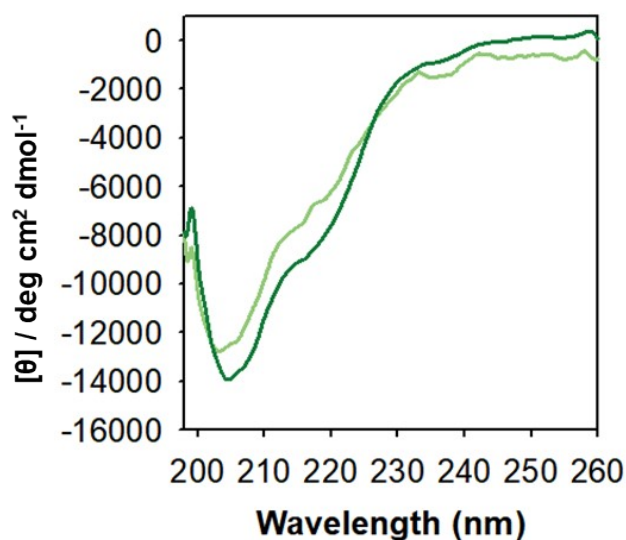
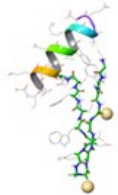


Fig. S12 CD spectra of α TMT2 (green) and α TMT2-ox (light green) in PBS (pH 7.0, 20 mM, 150 mM NaCl) at room temperature. $[\text{peptide}] = 20 \mu\text{M}$.

α TMT2

10 ns



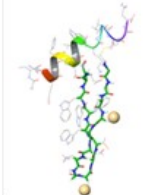
20 ns



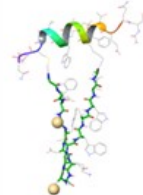
30 ns



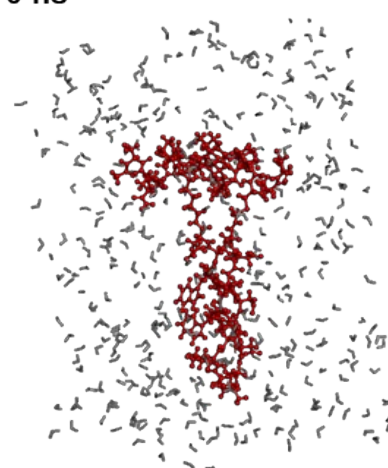
40 ns



50 ns



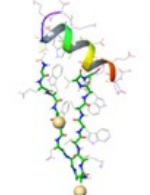
0 ns



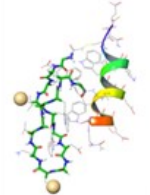
60 ns



70 ns



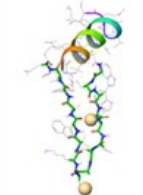
80 ns



90 ns



100 ns

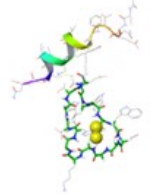


α TMT2-ox

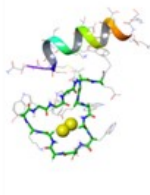
10 ns



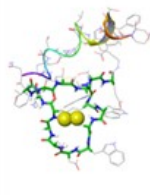
20 ns



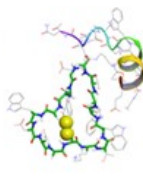
30 ns



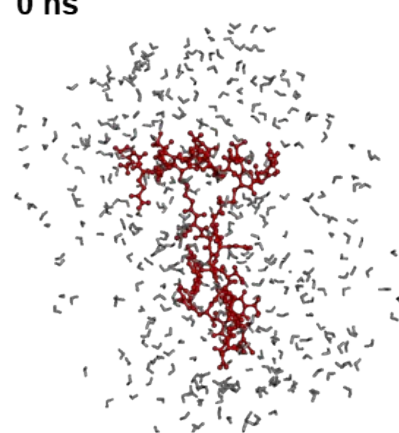
40 ns



50 ns



0 ns



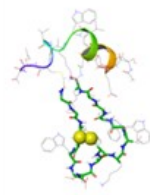
60 ns



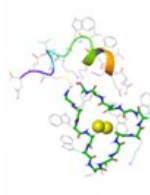
70 ns



80 ns



90 ns



100 ns

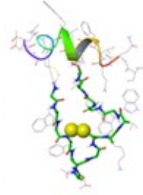


Fig. S13 MD simulation results of folding behaviors of α TMT2 and α TMT2-ox and snapshots of the input models with their geometry.

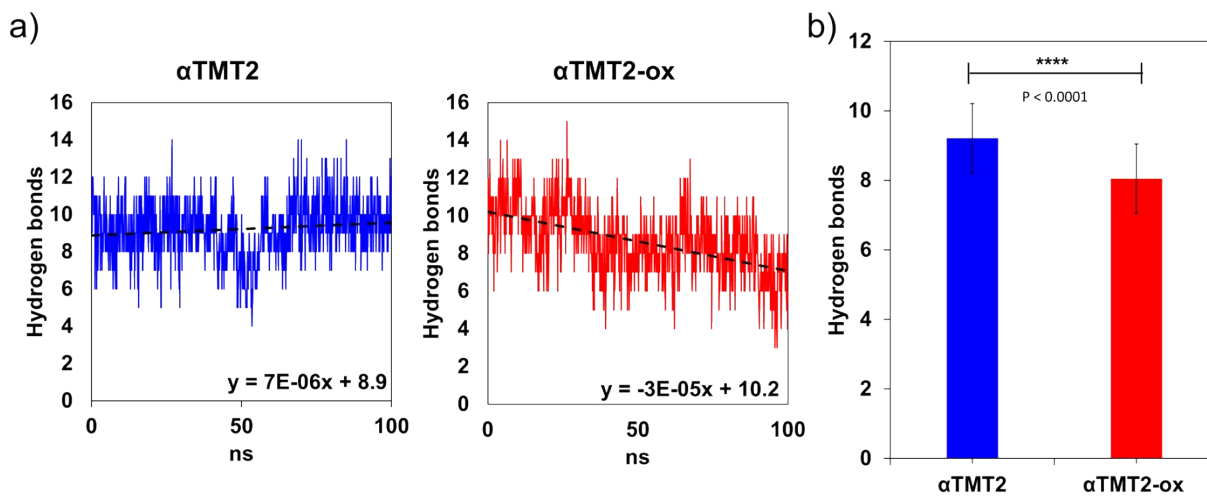


Fig. S14 (a) The numbers of hydrogen bonds in the helical backbone of α TMT2 and α TMT2-ox and (b) the average numbers of these hydrogen bonds over the simulation time (100 ns).

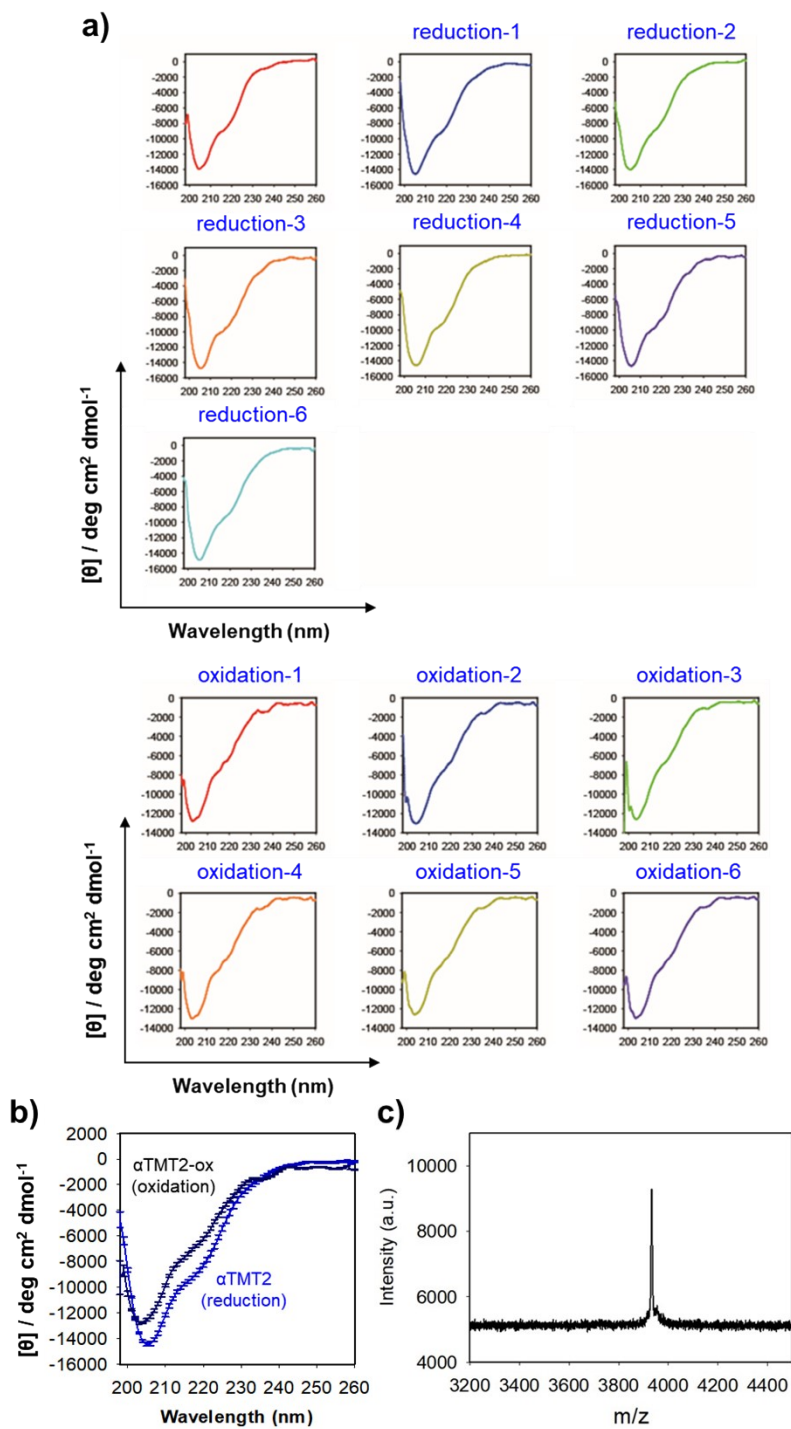


Fig. S15 (a) CD spectra of αTMT2 and $\alpha\text{TMT2-ox}$ during the six cycles of oxidation-reduction processes. (b) Averaged CD spectra of the oxidized and reduced αTMT2 peptides from the repeated oxidation-reduction experiment. (c) MALDI-TOF spectrum of αTMT2 after the repeated oxidation-reduction experiment (calculated MW: 3934.3, observed MW: 3935.6).

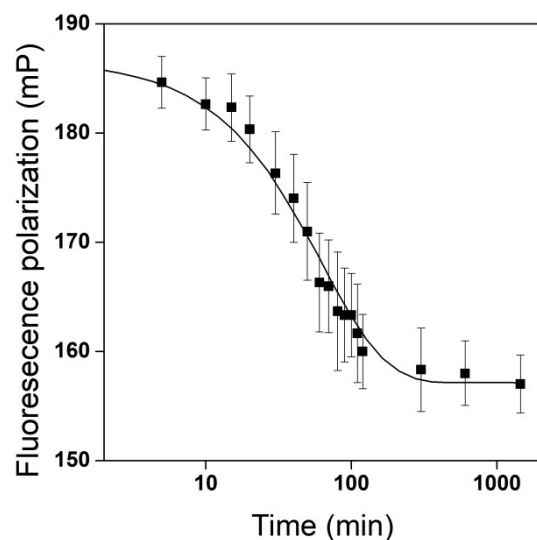


Fig. S16 Time-dependent FP competition assay on the α TMT2-ox peptides against p53f/hDM2 (100 nM/4 μ M) complexes in Tris-Cl buffer solution (pH 8.0, 20 mM, 10 mM imidazole, 150 mM NaCl, 10% glycerol) at room temperature, with the reducing agents (1 mM TCEP). The filter sets were 485/15 excitation and 535/25 emission. The experiment was performed in triplicate.

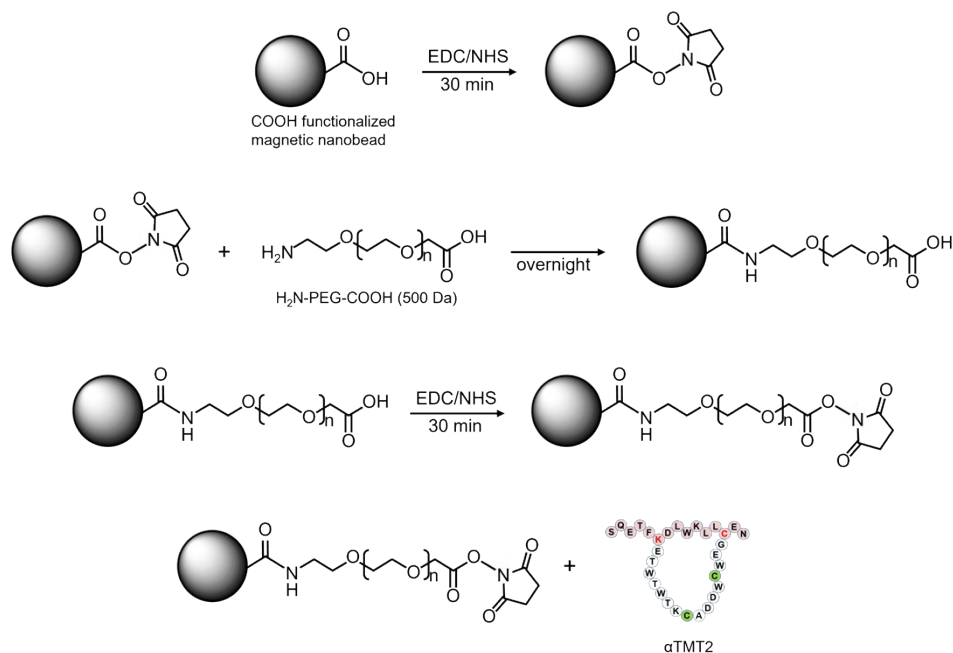


Fig. S17 Synthetic scheme of α TMT2-bead.

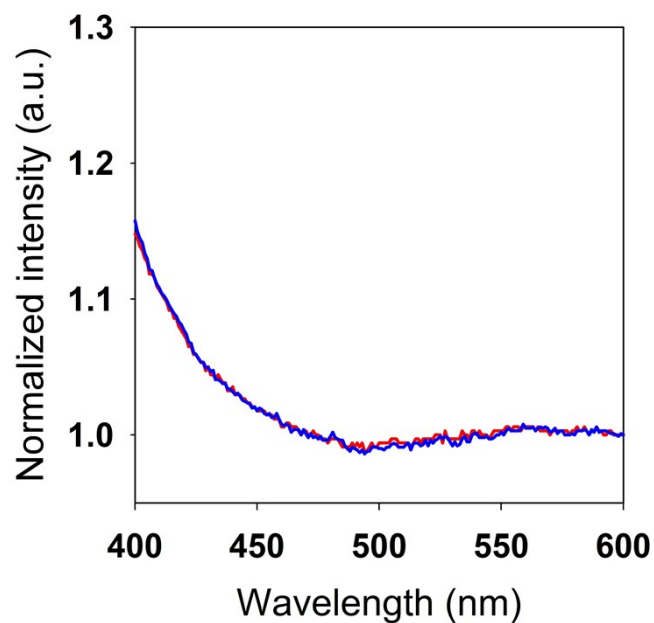


Fig. S18 The capture-and-release (CAR) experiment using PEG-magnetic bead conjugates without peptides. UV-Vis absorption spectra of released hGCs before (red) and after (blue) 5 h incubation in PBS (pH 7.0, 20 mM, 150 mM NaCl) containing the oxidizing agents (35 μ M) at room temperature.