## Genetically encoded fluorescent sensor for poly-ADP-ribose

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## **Supplementary Materials**

Figure S1	Protein sequence of Turquoise2-WWE and Venus-WWE.
Figure S2	Nucleus-to-cytoplasm distribution of FP-WWE constructs and FRET efficiency
	change after H <sub>2</sub> O <sub>2</sub> treatment.
Figure S3	sPARroW response to PAR accumulation inducing stimulus in U2OS stable
	cell line.
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Figure S5	Immunofluorescence analysis of PAR accumulation in U2OS cells.

## Turquoise2-WWE

MVSKGEELFTGVVPILVELDGDVNGHKFSVSGEGEGDATYGKLTLKFICTTGKLPVPWPTLVTT LSWGVQCFARYPDHMKQHDFFKSAMPEGYVQERTIFFKDDGNYKTRAEVKFEGDTLVNRIELKG IDFKEDGNILGHKLEYNYFSDNVYITADKQKNGIKANFKIRHNIEDGGVQLADHYQQNTPIGDG PVLLPDNHYLSTQSALSKDPNEKRDHMVLLEFVTAAGITLGMDELYKGGGSSGGGQISYASREF NGEYAWYYEGRNGWWQYDERTSRELEDAFSKGKKNTEMLIAGFPYVADLENMVQYRRNEHGRR<mark>F</mark> KIKRDIIDIPKKGVAGLRLDC

## Venus-WWE

MVSKGEELFTGVVPILVELDGDVNGHKFSVSGEGEGDATYGKLTLKLICTTGKLPVPWPTLVTT LGYGLQCFARYPDHMKQHDFFKSAMPEGYVQERTIFFKDDGNYKTRAEVKFEGDTLVNRIELKG IDFKEDGNILGHKLEYNYNSHNVYITADKQKNGIKANFKIRHNIEDGGVQLADHYQQNTPIGDG PVLLPDNHYLSYQSALSKDPNEKRDHMVLLEFVTAAGITLGMDELYKGGGSSGGGQISYASREF NGEYAWYYEGRNGWWQYDERTSRELEDAFSKGKKNTEMLIAGFLYVADLENMVQYRRNEHGRR<mark>F</mark> KIKRDIIDIPKKGVAGLRLDC

Fig. S1. Protein sequence of Turquoise2-WWE and Venus-WWE. Sequences of Turquoise2, Venus, and WWE are highlighted in cyan, green and pink, respectively. Position of Arg163 mutated in the control constructs to abolish PAR binding (R163A) is highlighted in red.

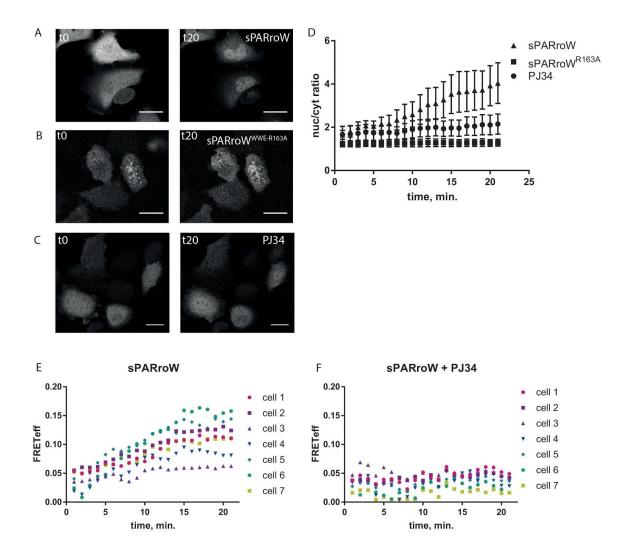


Fig. S2. Nucleus-to-cytoplasm distribution of WWE-FP constructs and FRET efficiency change after  $H_2O_2$  treatment. (A-C) Confocal microscopy images of cells in the donor channel before (left) and 20 min after (right) addition of  $H_2O_2$ . Representative images of cells expressing sPARroW (A), sPARroW-R163A (B) or sPARroW + PJ34 (C) are shown. Scale bars 25  $\mu$ m. (D) Nuclear-to-cytoplasm ratio change after  $H_2O_2$  treatment. (E) Single-cell measurement of FRET efficiency change in PJ34 pretreated cells after  $H_2O_2$  exposure.

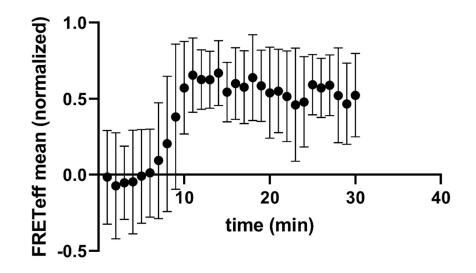


Fig. S3. sPARroW response to PAR accumulation inducing stimulus in U2OS stable cell line. Mean fluorescence intensity signals from donor, acceptor and FRET channels were used to calculate FRET efficiency in selected regions of interest according to the formula in Methods section. Mean values  $\pm$ SD are depicted.

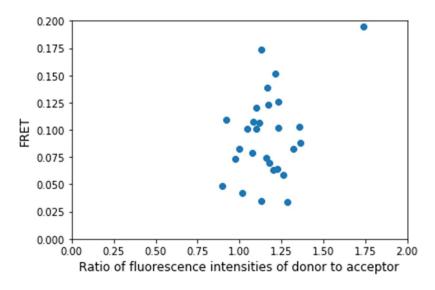


Fig. S4. sPARroW response in transiently transfected U2OS cells after H<sub>2</sub>O<sub>2</sub> treatment. Cell were transiently transfected with Turquoise2-WWE and Venus-WWE encoding plasmids at 1:1 ratio. Graph shows dependence of FRET efficiency on ratio of fluorescence signals of the donor to the acceptor (each dot corresponds to an individual cell). Note strong cell-to-cell variability of FRET efficiency independent on the donor/acceptor ratio within the observed narrow range.

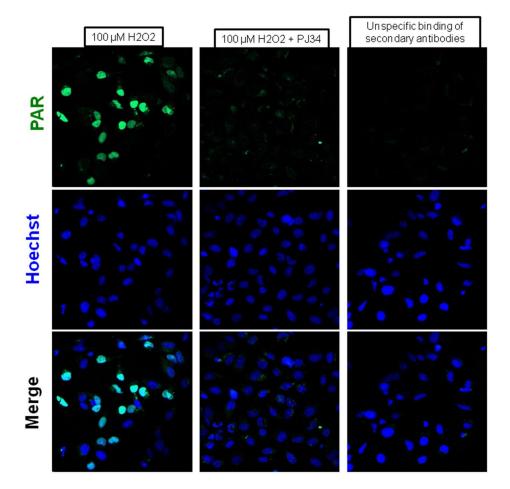


Fig. S5. Immunofluorescence analysis of PAR accumulation in U2OS cells. PAR-specific primary antibodies and Alexa Fluor 488-labeled secondary antibodies were used. Left column – cells treated with  $H_2O_2$  for 20 min. Middle column – cells pretreated with PARP inhibitor PJ34 and treated with  $H_2O_2$  for 20 min. Right column – control cells with no staining by the PAR-specific primary antibodies. Upper row – green fluorescence of the secondary antibodies; middle row – blue fluorescence of nuclei stained with Hoechst; bottom row – overlay of the two channels. Corresponding images were taken under the same conditions. Note strong cell-to-cell variability of PAR accumulation.