

SUPPLEMENTARY INFORMATION LEGENDS

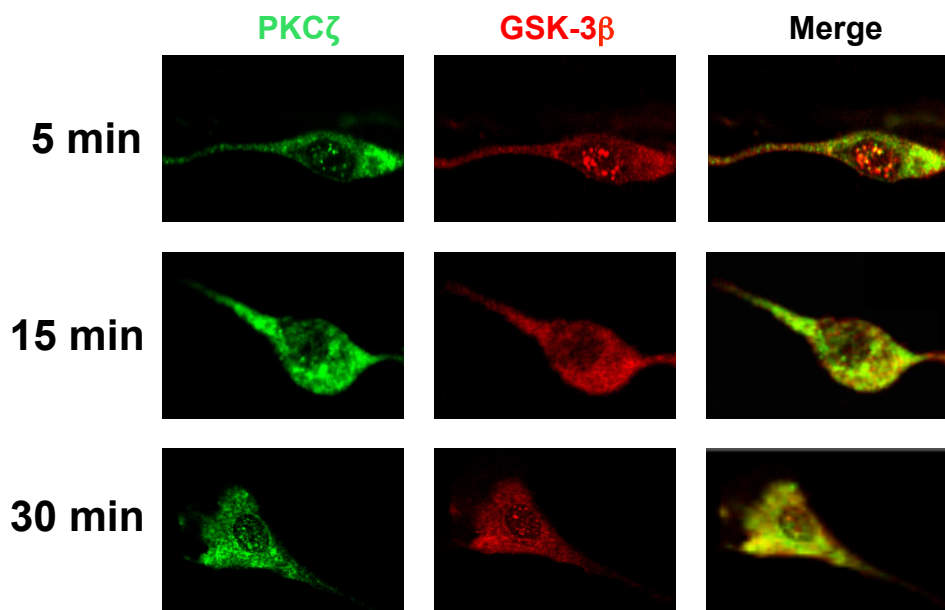
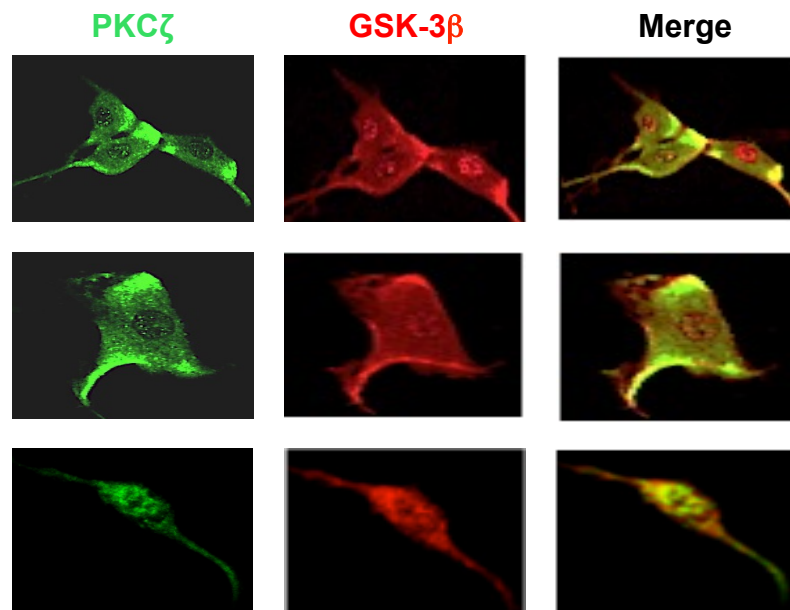
Figure S1. Time course of GSK-3 β stimulation by Wnt ligands (100 ng/ml). Changes in the intracellular localization of PKC ζ and GSK-3 β occur in a short time (5 min) while after 15 min, this returns to baseline in both non-malignant 112CoN cells (panel A) and malignant RKO and SW480 cells (panels B and C, respectively). Cells were fixed, permeabilized and co-immunostained with antibodies against GSK-3 β and against PKC ζ . Fluorescence was analyzed by laser confocal microscopy as described in the Experimental Procedures section. PKC ζ was visualized with FITC-conjugated goat anti-rabbit antibody and GSK-3 β with rhodamine goat anti-mouse antibody. Data are representative of three independent experiments.

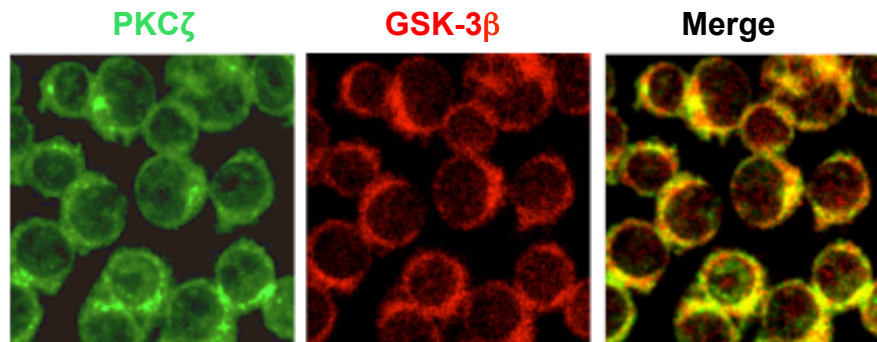
Figure S2. Efficiency of the knockdown of PKC ζ . RKO or SW480 cells were transfected with 2 μ g of pSuperPKC ζ -RNAi plasmid or with control pSuper plasmid using Lipofectamine 2000. The efficiency of the silencing of PKC ζ was determined 36 h post-transfection of cells by Western blot.

Figure S3. *In silico* analysis of the GSK-3 β aminoacid sequence showing the putative PKC phosphorylation sites in GSK-3 β . The sites obtained from the analysis appear underlined in black; three sites displaying the classical PKC consensus sequence with high probability score are underlined in red, and the Serine 147, that exactly matched with the

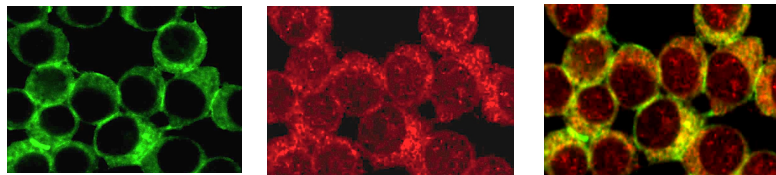
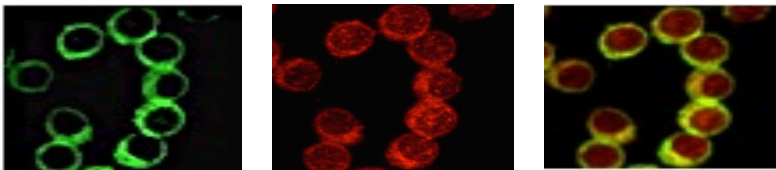
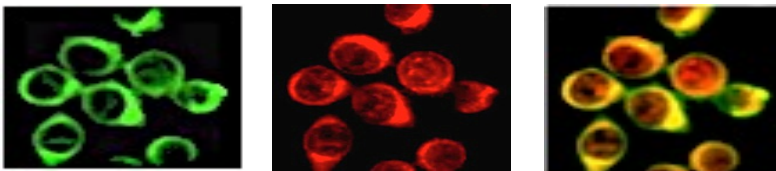
consensus sequence recognized by the P-Ser-PKC-substrate antibody used in the experiments (R/K₍₀₋₂₎ Ser X_(hydrophobic)R/K) is shown underlined with a curly red line.

Figure S4. Efficiency of transfection of GSK-3 β wild-type, GSK-3 β /S147A and GSK-3 β /S147E. RKO cells were transfected with the void plasmid (vector) or with plasmid-encoding wild type GSK-3 β or plasmid-encoding GSK-3 β mutants. GSK-3 β was immunoprecipitated 24 h post-transfection from cell extracts and analyzed by Western blot. Actin antibody was used to control for equal loading. The results are representative of three independent experiments using different cell preparations.

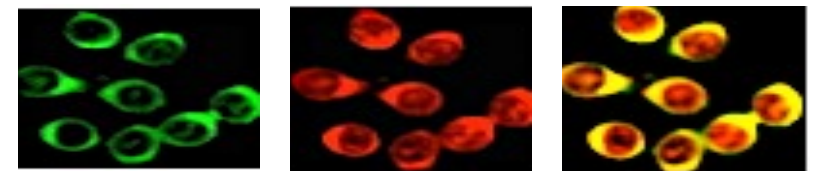
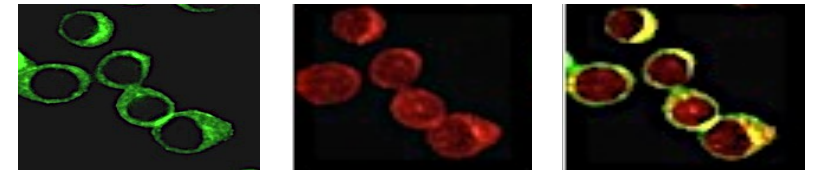
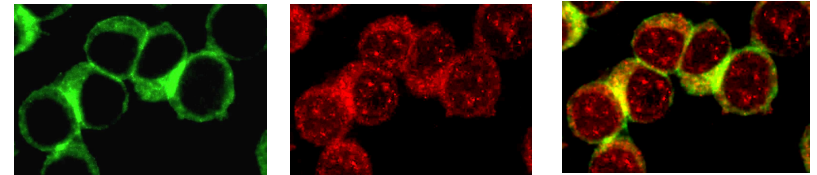
A**Control****Wnt3a****Wnt5a****Figure S1A**

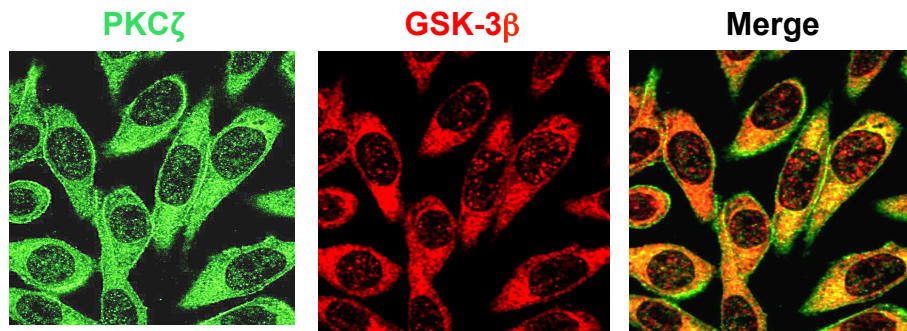
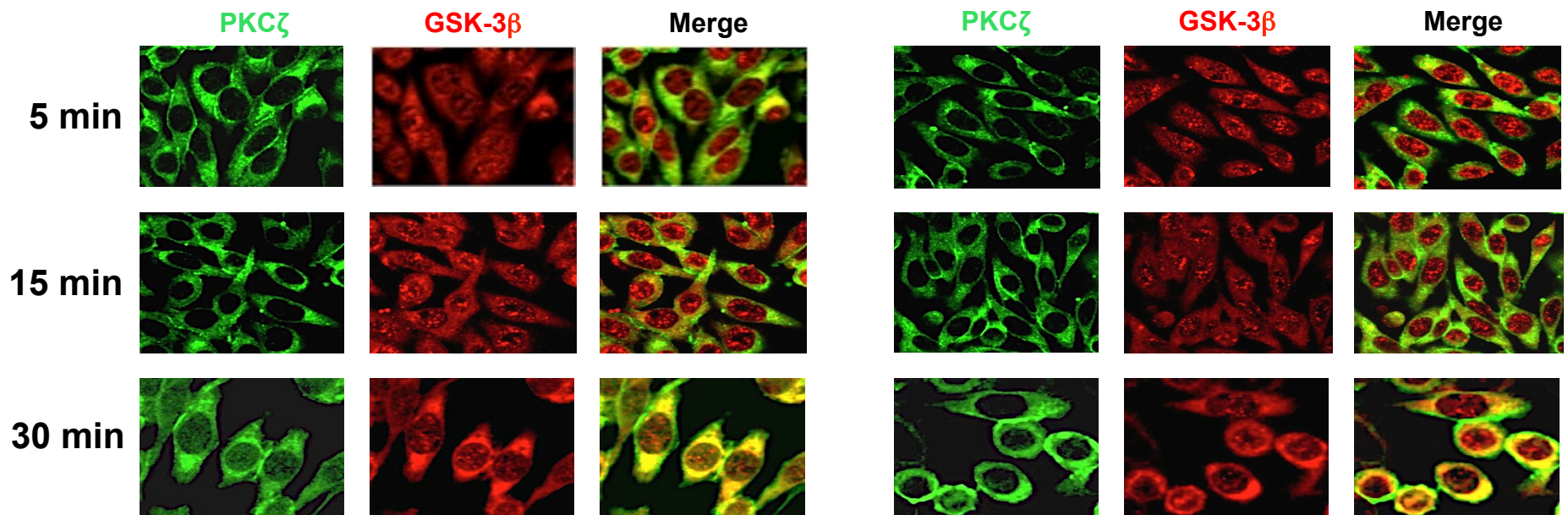
B**Control****Figure 1S B****Wnt3a**

PKC ζ GSK-3 β Merge

5 min**15 min****30 min****Wnt5a**

PKC ζ GSK-3 β Merge



C**Control****Figure 1S C****Wnt3a****Wnt5a**

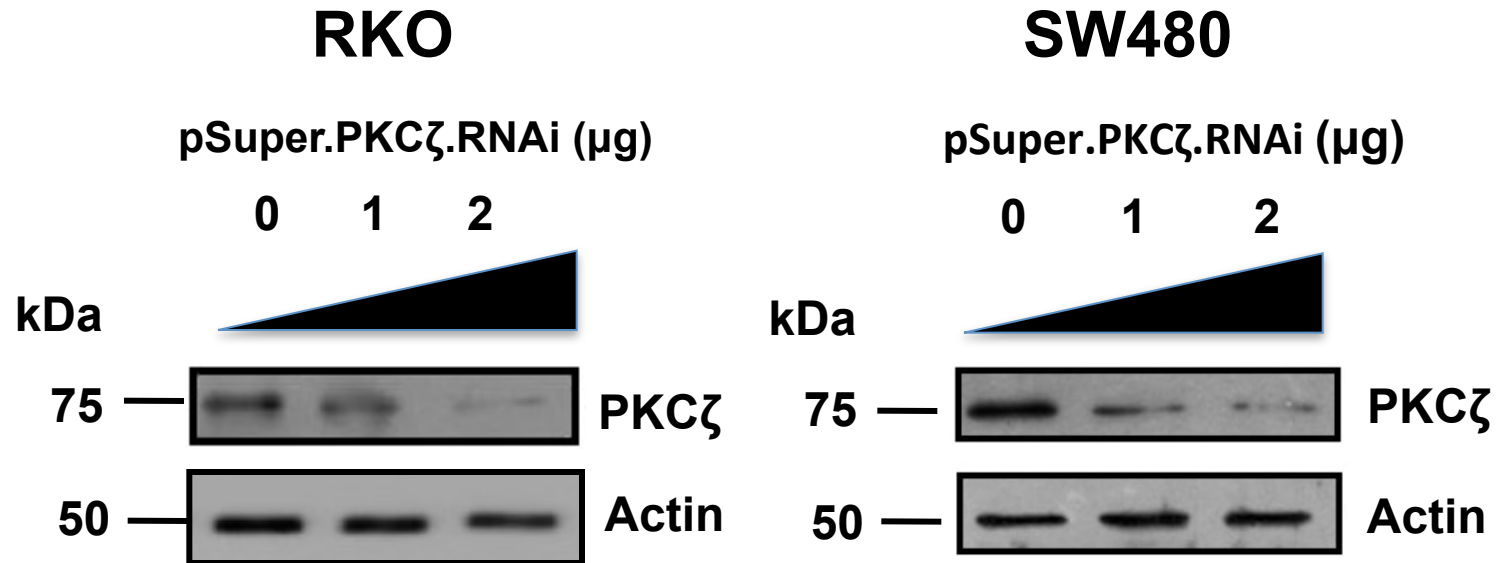
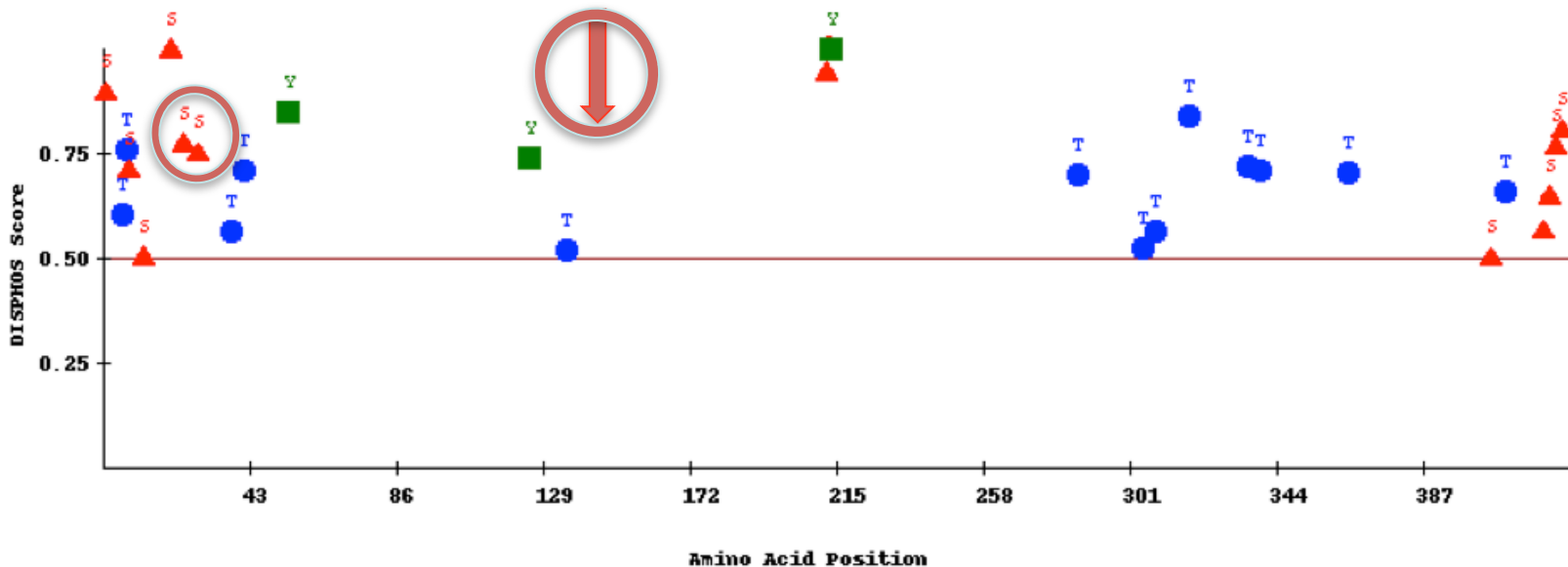


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>gi|21361340|ref|NP_002084.2| glycogen synthase kinase-3 beta isoform 1 [Homo sapiens]

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 ASTPTNATAASDANTGDRGQTNNAASASASNST



<http://www.dabi.temple.edu/disphos/pred/predict>

Figure S3. Putative PKC phosphorylation sites in GSK-3 β (NetphosK 2.0) The sites obtained from the analysis appear underlined in black; three sites displaying the classical PKC consensus sequence with high probability score are underlined in red, and the Serine 147, that exactly matched with the consensus sequence recognized by the P-Ser-PKC-substrate antibody used in the experiments ($R/K_{(0-2)}$ Ser X_(hydrophobic) R/K) is shown underlined with a curly red line.

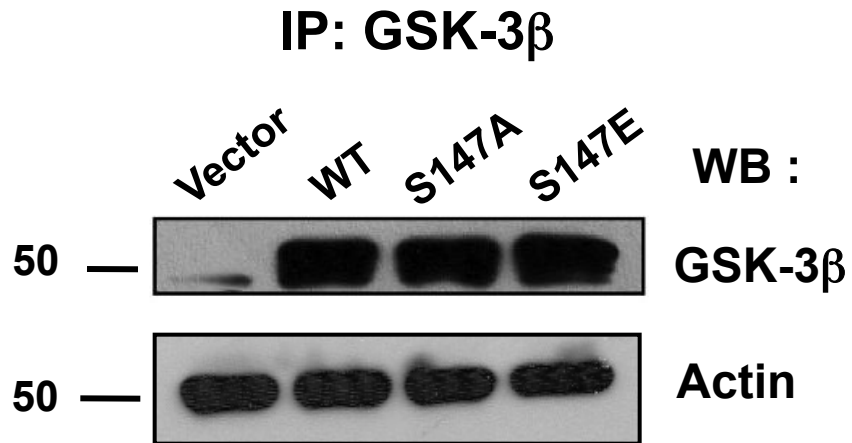


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