

SUPPLEMENTAL DATA

Fig S1 : Control analysis of reciprocal bleed-through fluorescence from the red and green channels.

COS-7 cells were transiently transfected with the expression vector encoding HA-SRC-1. Cells were fixed after 40 h and immunolabeled overnight at 4°C with either the primary rat anti-HA (upper panel) or the rabbit anti-S7/Rpt1 antibodies (lower panel). First antibody was followed by a 30 min incubation with a mix of the two corresponding secondary antibodies Dylight 549 anti-rat (red) and alexa 488 anti-rabbit (green). Cells were analyzed by confocal microscopy as described in Methods section. Note the lack of overlap of both fluorescences. DIC: Differential Interferential Contrast.

Fig S2 : Colocalization analyses of SRC-1 with antigens and organelles.

A and B, COS-7 cells were transiently transfected with vectors encoding HA-SRC-1. Cells were fixed after 40 h and immunolabeled with the anti-PML (A) or anti-SC-35/SRp30 spliceosome component (B) antibodies. Colocalization analysis was performed by confocal microscopy. Note the partial colocalization of SRC-1 with PML nuclear bodies and the absence of colocalisation with SC-35.

C and D, COS-7 cells were transiently transfected with vector encoding GFP-SRC-1. After 40 h, cells were incubated with a mitochondrial (C) or lysosomal tracker (D) (Invitrogen), followed by fixation and observation by confocal microscopy. No significant colocalization was observed.

E, COS-7 cells were cotransfected with the vector encoding HA-SRC-1 and the pEGFP-Peroxy vector (Clontech). Immunodetection was performed with the anti-HA antibody and cells were analyzed by confocal microscopy. No significant colocalization was observed.

Fig S3: Epoxomicin increases SRC-1 expression level.

COS-7 cells were transfected with the vector encoding SRC-1 and incubated during 15 h in the presence of epoxomicin (500 nM) or vehicle (DMSO, -). Expression of SRC-1 was analyzed by Western blot using anti-HA monoclonal antibody. The bands were quantified as described in “Materials and Methods”.

Fig S4: The Nuclear Export Signal deletion of SRC-1 decreases its turnover rate.

COS-7 cells were transfected with the wt SRC-1 or the Δ (NES) encoding vector (deleted of the Nuclear Export Signal). Seventy-two hours after transfection, cells were treated with cycloheximide (100 μ g/ml) during 1, 4 or 6 h. Whole cell extracts were analyzed by electrophoresis on 7.5% SDS-PAGE and immunoblotted with the indicated antibodies. Band intensities (right panel) were quantified as described in “Material and Methods”. Note the decrease in kinetics degradation rate of the nuclear mutant Δ (NES) compared to wt SRC-1.

Fig S5 : PRB is degraded in the presence of R5020 but not RU486.

Ishi PR-B cells were treated during 24 h with vehicle (control, -), the agonist R5020 (10 nM) or the antagonist RU486 (10 nM) and western blot analysis was performed with the corresponding whole cell extracts as described in “Materials and Methods”. PR-B was detected using monoclonal antibody Let126 and loading control protein α -tubulin was detected using anti- α -tubulin antibody.

Fig S6 : R5020 or RU486 does not modify endogenous SRC-1 mRNA levels.

Ishi PR-B cells were treated during 24 h with vehicle (EtOH, -), R5020 (10 nM) or RU486 (10 nM). Total RNA was extracted and relative expression of SRC-1 mRNA levels were quantified by quantitative RT-PCR as described in “Materials and Methods”. Results were normalized by the 18S RNA and are expressed as mean \pm SEM of six independent determinations.

Fig S7 : Coactivator specificity in the ligand-induced down-regulation.

Ishi PR-B cells were transfected as indicated, with either the SRC-1, SRC-2 or SRC-3 encoding vector. Forty-eight h after transfection, cells were cultured for 24 h either in the absence of ligand (control vehicle, -H) or in the presence of R5020 (10 nM). Whole cell extracts were analyzed by electrophoresis on 7.5% SDS-PAGE and immunoblotted with the indicated antibodies.

Fig S8 : Epoxomicin and MG132 do not increase expression levels of the two non degradable Δ (PEST) and Δ (bHLH) mutants.

COS-7 cells were transfected with Δ (PEST) and Δ (bHLH) encoding vectors and incubated in the absence (vehicule, -) or presence of either epoxomicin (500 nM) or MG132 (5 μ M) for 15 h. Expression level of SRC-1 mutants was analyzed by Western blot using the indicated antibodies. Note the absence of accumulation of both mutants in the presence of both proteasome inhibitors.