Supplemental Data

Primers

The sequence of primer sets used in this study are:

For ChIP analysis

Primer sets used for p107, E2F1, $cyclin\ A$ and cdc25A promoters and U_{2C} have been described (Takahashi et al., 2000 and Tyagi et al., 2007).

1. Apafl

Forward GGGTGTGTTTATTTGCATAAGCGGGC

Reverse TCTGGACAGCGGAGCAGTCAAAT

2. *p73*

Forward TGCGACGGCTGCAGGTAGGA

Reverse TGTGTCGCCTTGTCCACTAGCTC

3. *p14*^{ARF}

Forward CGCTGAGGGTGGGAAGATGGT

Reverse GTTCCTCTCCCTCCCGCCTA

4. *p21*

Forward AGCTGCCGAAGTCAGTTCCTTGT

Reverse TCTCTCACCTCCTGAGTGCCT

For RT-PCR analysis

Primer sets used for *E2F1* and *GAPDH* mRNA have been described (Tyagi et al., 2007).

1. Apaf1 mRNA

Forward TCCAGTCCAGGTTTCAGCACAAGA

Reverse TTCACTGTTTCCTGATGGCCTCGT

2. *p73* mRNA

Forward CGGGCCATGCCTGTTTACAAGAAA

Reverse TGGAGCAGACTGTCCTTCGTTGAA

3. *p14*^{ARF} mRNA

Forward AGGTTCTTGGTGACCCTCCGGATT

Reverse CCCATCATCATGACCTGGTCTTCT

4. *p21* mRNA

Forward TTAGCAGCGGAACAAGGAGTCAGA

Reverse ACACTAAGCACTTCAGTGCCTCCA

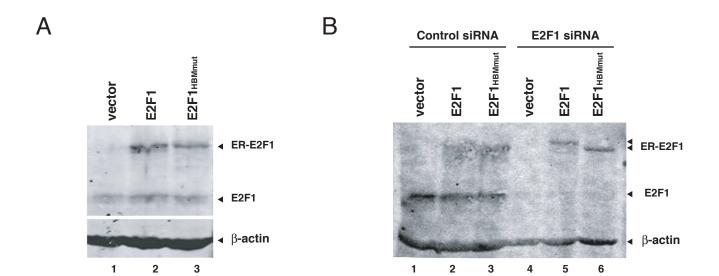
5. hcf-1 mRNA

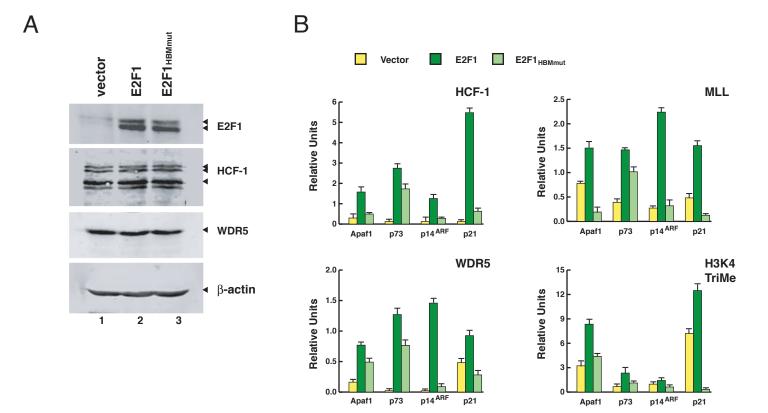
Forward AAGACAGCTCTGGCACCAA

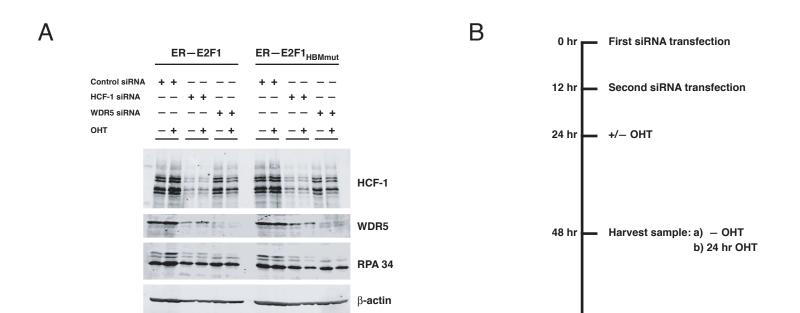
Reverse GTCTGGAGAAGAATCCAGG

Antibodies

Antisera used in these experiments were: polyclonal anti-HCF-1_N subunit, anti-WDR5, anti-β actin, anti-E2F1 for immunoblots (KH95), anti-E2F1 for ChIP and immunoprecipitation (sc-193), anti-MLLc (clone 9-12); anti-trimethyl-Histone H3K4; ALEXA FLUOR 680 goat anti-rabbit and ALEXA FLUOR 680 goat anti-mouse have been described before (Tyagi et al., 2007); p53 (sc-6243, Santa Cruz biotechnology); anti-PCNA (Transduction Laboratories); ALEXA FLUOR 488 goat anti-rabbit, ALEXA FLUOR 546 goat anti-mouse (Invitrogen); anti 53BP1 (kind gift of T. Halazonetis, University of Geneva); anti RPA 70 and RPA 34 antibody (kind gift of B. Stillman, Cold Spring Harbor Laboratory)







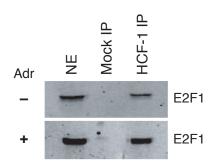
72 hr

Harvest sample: c) 48 hr OHT

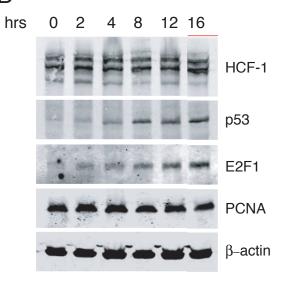
C

| siRNA | ОНТ | hrs with OHT | Exp 1 | | Exp 2 | | Av of Exp 1 & 2 | SD |
|---------|-----|-----------------|----------|----------|----------|----------|--------------------|-------|
| | | | % sub-G1 | Fold Inc | % sub-G1 | Fold Inc | | |
| Control | - | 0 | 5.6 | 1 | 6.25 | 1 | 1 | |
| Control | + | 24 | 14.6 | 2.6 | 15.3 | 2.44 | 2.52 | 0.08 |
| Control | + | 48 | 19 | 3.39 | 22.8 | 3.64 | 3.51 | 0.12 |
| HCF-1 | - | 0 | 12.9 | 1 | 15 | 1 | 1 | |
| HCF-1 | + | 24 | 15.2 | 1.17 | 18.02 | 1.2 | 1.18 | 0.015 |
| HCF-1 | + | 48 | 16.4 | 1.27 | 22.3 | 1.48 | 1.37 | 0.1 |
| WDR5 | - | 0 | 16.6 | 1 | 17.9 | 1 | 1 | |
| WDR5 | + | 24 | 17.2 | 1.03 | 21.7 | 1.21 | 1.12 | 0.09 |
| WDR5 | + | 48 | 18.4 | 1.1 | 22 | 1.22 | 1.16 | 0.06 |





В



Adriamycin

C

| siRNA | Adr | hrs with Adr | Exp 1 | | Exp 2 | | Av of Exp 1 & 2 | SD |
|---------|-----|-----------------|----------|----------|----------|----------|--------------------|------|
| | | | % sub-G1 | Fold Inc | % sub-G1 | Fold Inc | | |
| | | | | | | | | |
| Control | - | 0 | 4.3 | 1 | 3.9 | 1 | 1 1 | |
| Control | + | 24 | 14.3 | 3.32 | 11.4 | 2.92 | 3.12 | 0.2 |
| Control | + | 48 | 23.8 | 5.53 | 24.6 | 6.31 | 5.92 | 0.38 |
| | | | | | | | | |
| HCF-1 | - | 0 | 11.5 | 1 | 11.9 | 1 | 1 | |
| HCF-1 | + | 24 | 12.3 | 1.07 | 13.2 | 1.11 | 1.08 | 0.01 |
| HCF-1 | + | 48 | 18.7 | 1.62 | 21.4 | 1.79 | 1.71 | 0.08 |
| | | | | | | | | |
| WDR5 | - | 0 | 10.3 | 1 | 9.3 | 1 | 1 | |
| WDR5 | + | 24 | 11.9 | 1.15 | 10.3 | 1.11 | 1.13 | 0.02 |
| WDR5 | + | 48 | 23 | 2.23 | 20.6 | 2.21 | 2.22 | 0.01 |

Figure Legends

Figure S1

- A Immunoblot analysis of U2OS cell extracts stably expressing empty vector, ER-E2F1, or ER-E2F1_{HBMmut} are shown. Endogenous (E2F1) and recombinant (ER-E2F1) E2F1 proteins were visualized using anti-E2F1 antisera.
- B Immunoblot analysis of control or E2F1 siRNA treated U2OS cell extracts stably expressing ER-E2F1 or ER-E2F1_{HBMmut} are shown. (The difference in ER-E2F1 and ER-E2F1_{HBMmut} mobility clearly seen in lanes 5 and 6 is a phosphatase-sensitive effect, which varies with electrophoresis conditions.)

Figure S2

- A Immunoblot analysis of E2F1, HCF-1, WDR5, and β-actin proteins (as indicated) from U2OS cell extracts transfected with plasmid encoding empty vector, E2F1, or E2F1_{HBMmut} mutant and harvested 24-hrs post-transfection, and used for the ChIP experiment in Fig. **5B** are shown.
- B Quantitation of HCF-1, MLL, WDR5, and H3K4 trimethylation ChIP analyses of U2OS cells transfected with plasmids encoding empty vector, E2F1 or E2F1_{HBMmut}, by triplicate real-time PCR, of indicated promoters is shown. The cells samples are the same as in Fig. **5B** and are similarly normalized to E2F1 levels.

Figure S3.

- A Immunoblot analysis of HCF-1, WDR5, RPA34, and β-actin proteins from U2OS cells stably expressing ER-E2F1 or ER-E2F1_{HBMmut} and treated with control, HCF-1 or WDR5 siRNA for 72 hrs are shown. The cells were induced with OHT as indicated.
- **B** Schematic outline of experiment performed in Fig. 6C.

C Raw data (% sub-G1) and calculated relative-fold increase (Fold Inc) obtained in Fig 6C.

Figure **S4**.

- A Association of endogenous HCF-1 and E2F1, in nuclear extracts from HeLa cell treated with adriamycin (Adr) for 16 h. HCF-1 or mock immunoprecipitates (IP) of adriamycin treated (+) or untreated (-) samples were probed for E2F1. NE, input nuclear extract (3%).
- B Immunoblot analysis of endogenous levels of HCF-1, p53, E2F1, PCNA and β -actin proteins from U2OS cells treated with 2μ g/ml Adriamycin and harvested after indicated times are shown.
- C Raw data (% sub-G1) and calculated relative-fold increase (Fold Inc) obtained in Fig 7C.