Sup. Figure 1. (A) Summary and statistical analysis of experiments in Figure 5E with human normal ovary (n=7) and ovarian tumor tissues (n=9). Scale bar, 25 μ m (magnification, ×40)

Sup. Figure 2. Protein stability experiment between RDM1 and P53 in SKOV3 si*RDM1*/siN cells with Cycloheximide (CHX) treatment.

Sup. Figure 1 Summary and statistical analysis of experiments in IHC

Α

| Protein level | Samples | - | + | ++ | +++ | Total samples |
|------------------|---------------|-----------|-----------|-----------|-----------|------------------|
| RDM1 | Normal ovary | 3 (42.8%) | 3 (42.8%) | 1 (14.3%) | 0 (00.0%) | 7 |
| | Ovarian tumor | 0 (0.0%) | 1 (11.1%) | 5 (55.6%) | 3 (33.3%) | 9 |
| P53 | Normal ovary | 0 (00.0%) | 0 (00.0%) | 4 (57.1%) | 3 (42.8%) | 7 |
| | Ovarian tumor | 1 (11.1%) | 2 (22.2%) | 2 (22.2%) | 4 (66.7%) | 9 |
| RAD51 | Normal ovary | 2 (28.6%) | 3 (42.8%) | 1 (14.3%) | 1 (14.3%) | 7 |
| | Ovarian tumor | 1 (11.1%) | 1 (11.1%) | 4 (66.7%) | 3 (33.3%) | 9 |
| RAD52 | Normal ovary | 3 (42.8%) | 3 (42.8%) | 1 (14.3%) | 0 (00.0%) | 7 |
| | Ovarian tumor | 0 (00.0%) | 2 (22.2%) | 4 (66.7%) | 3 (33.3%) | 9 |

Sup. Figure 2

