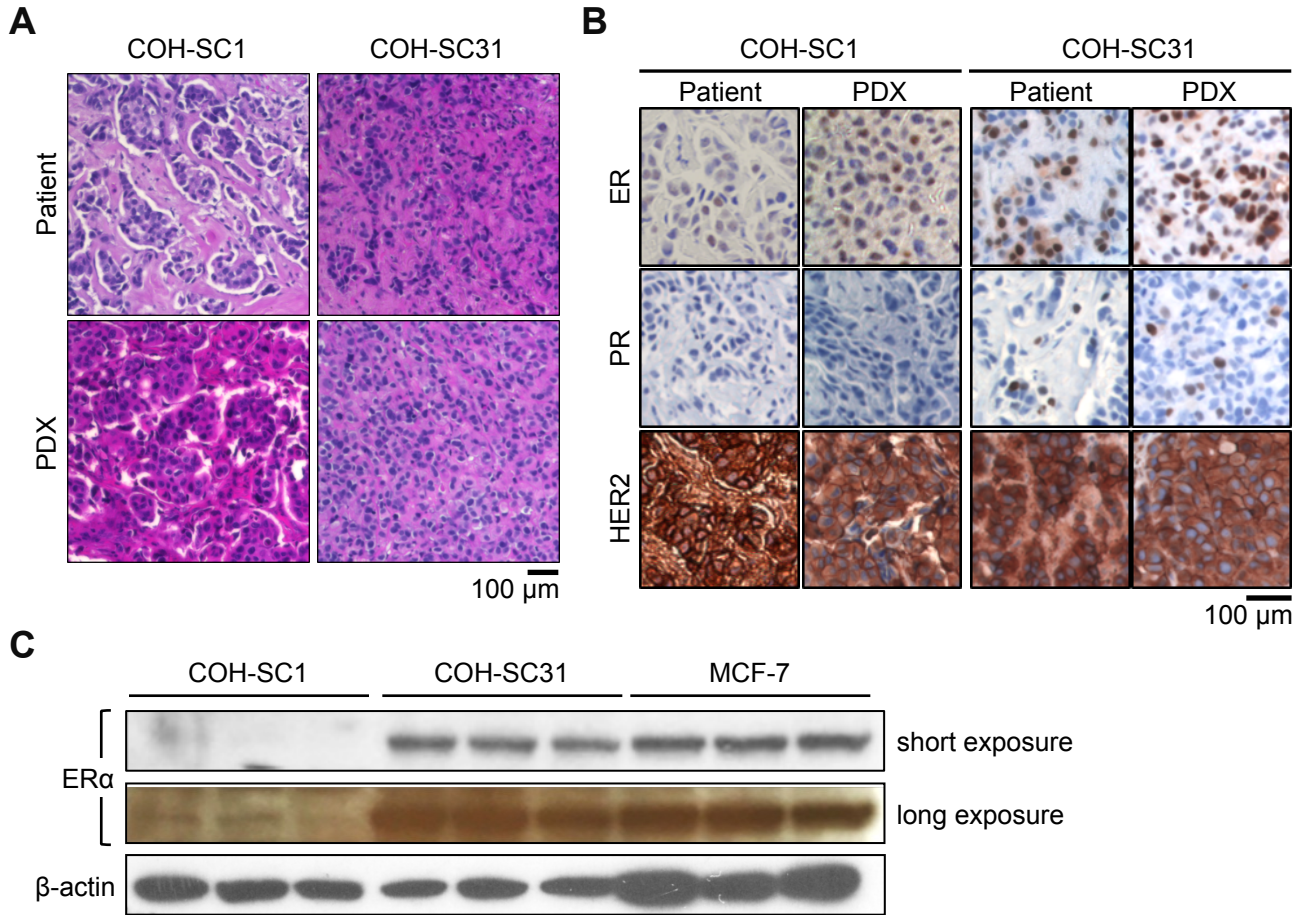


Figure S1, Hsu & Wu *et al.*



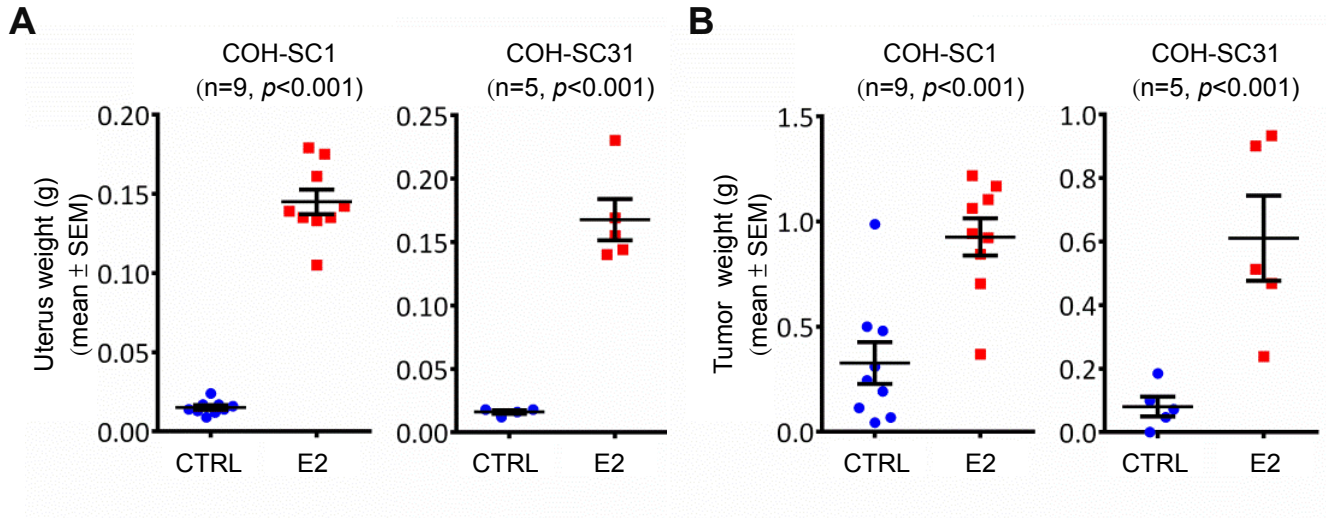
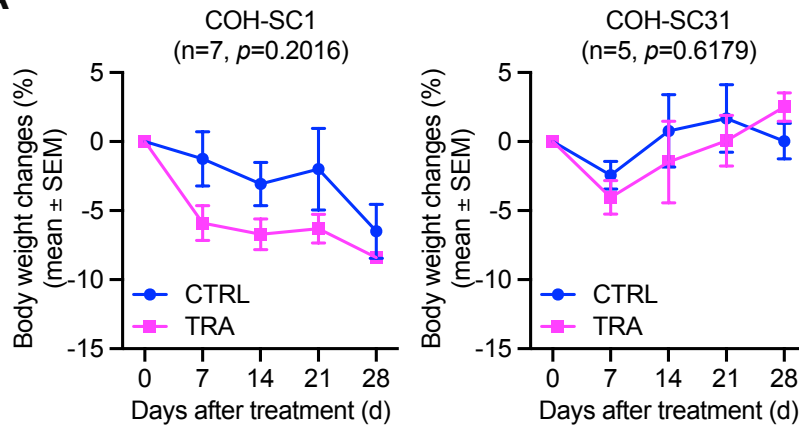
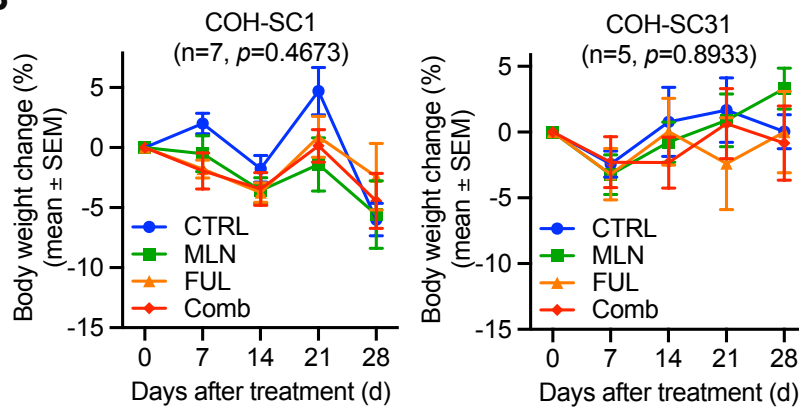


Figure S3, Hsu & Wu *et al.*

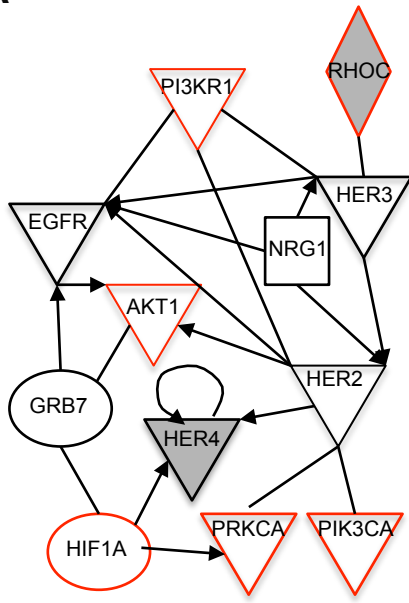
A



B



A



B

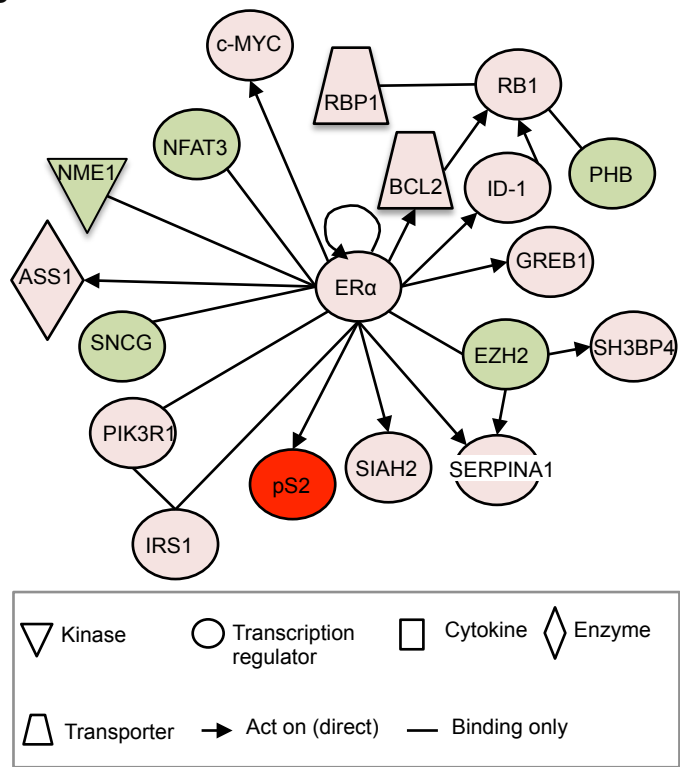


Figure S5, Hsu & Wu *et al.*

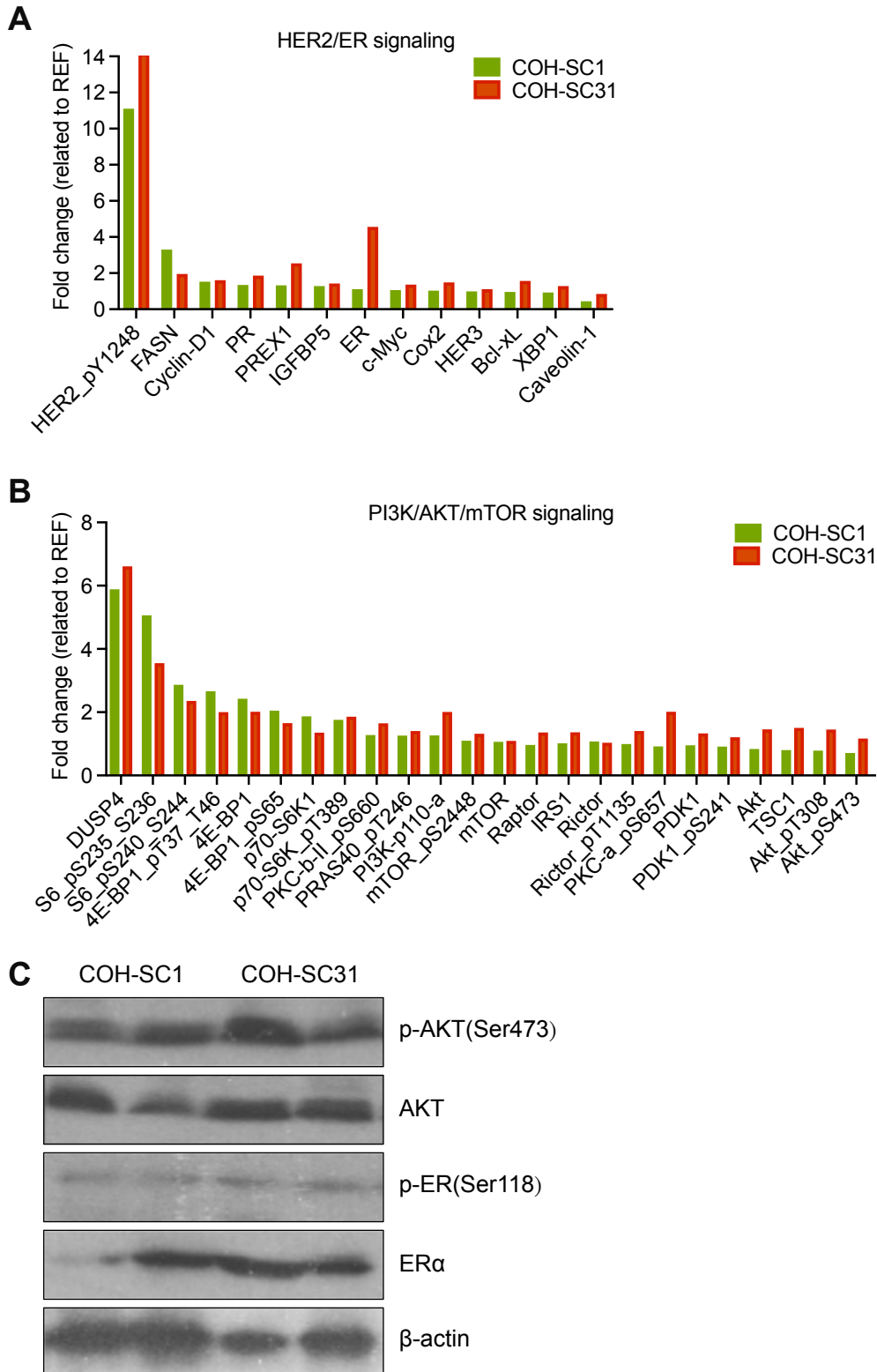
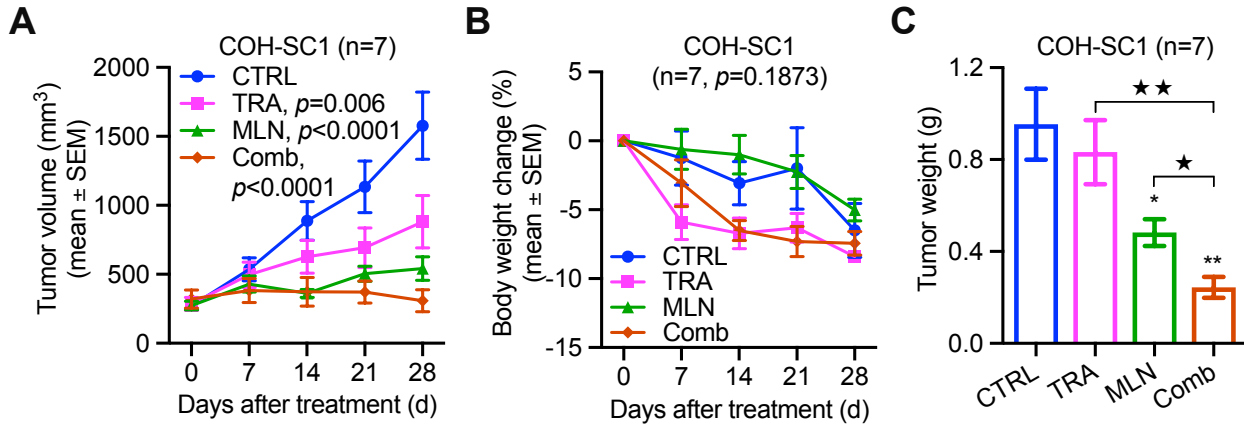


Figure S6, Hsu & Wu *et al.*



D

| Sample | Tumor ER expression (IHC) | | Mitotic Counts | |
|--------|---------------------------|-----|--------------------------------|---------|
| | negative/positive | % | # of mitoses per 15 40X-fields | |
| CTRL-1 | borderline positive | 1-5 | 35 | 46 ± 12 |
| CTRL-2 | positive | 20 | 44 | |
| CTRL-3 | borderline positive | 1-5 | 59 | |
| TRA-1 | positive | 20 | 30 | 20 ± 9 |
| TRA-2 | positive | 30 | 17 | |
| TRA-3 | positive | 20 | 14 | |
| MLN-1 | negative | < 1 | 23 | 25 ± 9 |
| MLN-2 | borderline positive | 1-5 | 35 | |
| MLN-3 | negative | < 1 | 18 | |
| Comb-1 | positive | 20 | 2 | 2 ± 1 |
| Comb-2 | positive | 10 | 3 | |
| Comb-3 | positive | 10 | 1 | |