

**Supplementary Table S1:  
validation of Sequencing results by ddPCR.**

Sample ID	sequencing		validation: ddPCR	
	position	% reads	position	% fractional abundance
ESR 3	D538G	6.86	D538G	3.36
ESR 5	D538G	39.30	D538G	37.21
ESR 9	D538G	7.93	D538G	6.71
ESR 13	D538G	1.97	D538G	4.96
ESR 14	D538G	45.57	D538G	44.09
ESR 20b	D538G	37.93	D538G	37.97
ESR 40	D538G	42.74	D538G	41.94
ESR 73	D538G	1.01	D538G	10.93
ESR 6	D538G	not detected	D538G	not detected
ESR 20a	D538G	not detected	D538G	not detected
ESR 50	D538G	not detected	D538G	not detected
ESR 71	D538G	not detected	D538G	not detected

Figure S1:  
DFS and OS for the metastatic cohort

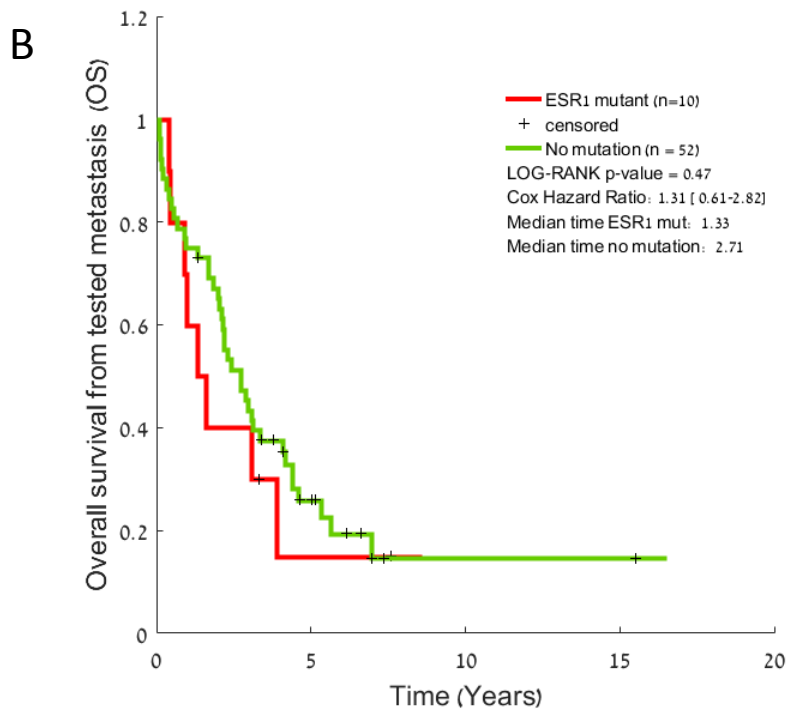
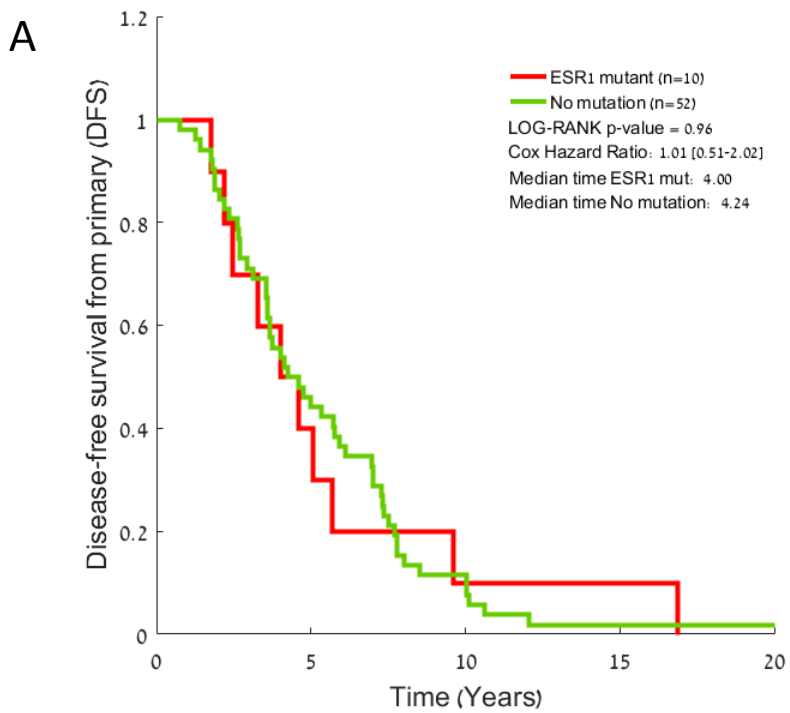
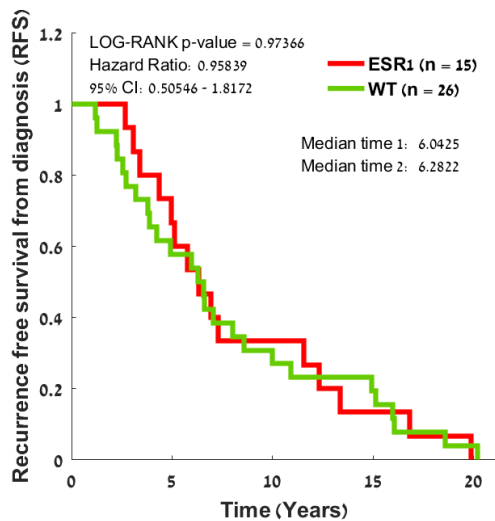


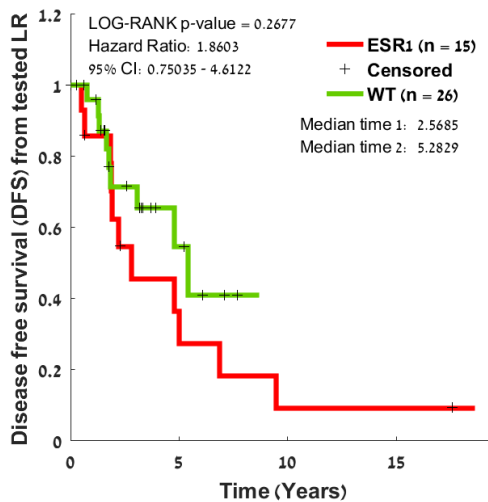
Figure S2:

RFS, DFS and DRFS for the loco-regional cohort: mutant vs. WT

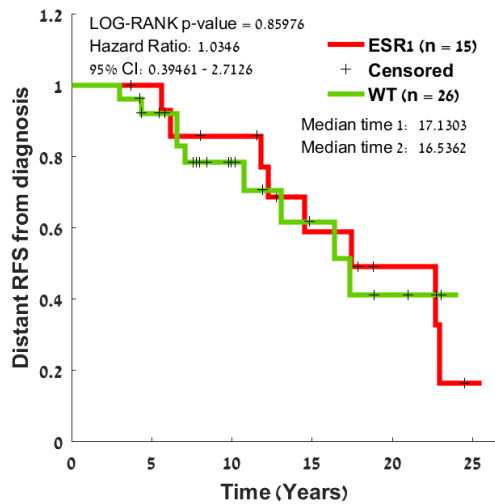
A



B



C



D

