



Influence of *TONSL* on tumor suppressor function of *RAD51* and resistance to CDK4/6 inhibitors in ER+ breast cancer.

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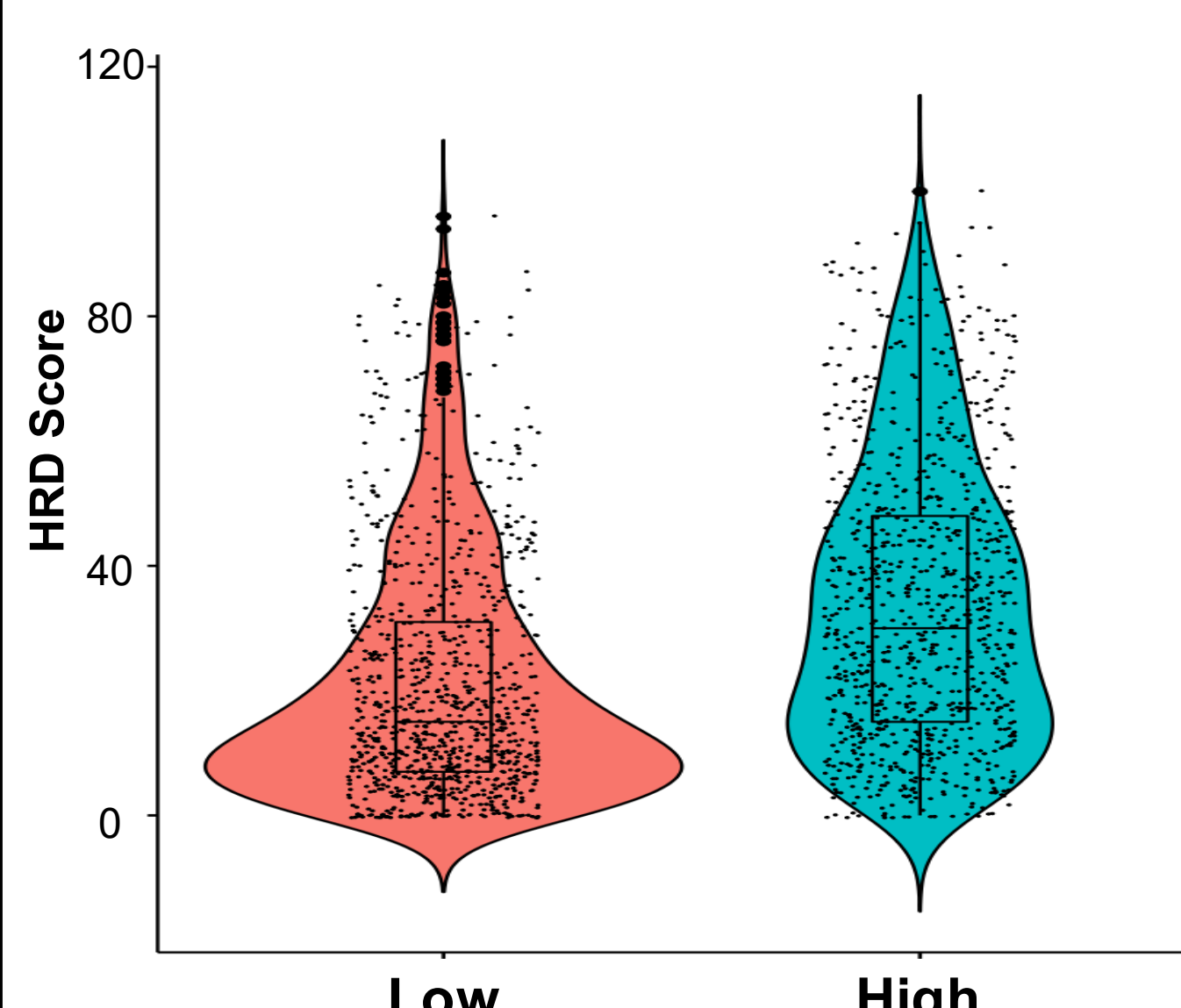
BACKGROUND

- Homologous Recombination (HR) and Replication Stress (RS) resolution mechanisms help to maintain genomic stability. *TONSL*, *MMS22L*, *RAD51*, *BRCA1/2* are the few proteins that cooperatively coordinate HR and RS resolution.
- Therefore, genes in these pathways are expected to function as tumor suppressors.
- However, only *BRCA1/2* function as tumor suppressors as their germline mutations increase cancer susceptibility.
- *TONSL* and *RAD51* are frequently overexpressed in tumors.
- Here, we investigated the relationship between *TONSL-RAD51* axis in breast cancer (BC) outcome and response to CDK4/6 inhibitors (CDK4/6i) and endocrine therapy (ET).

METHODS

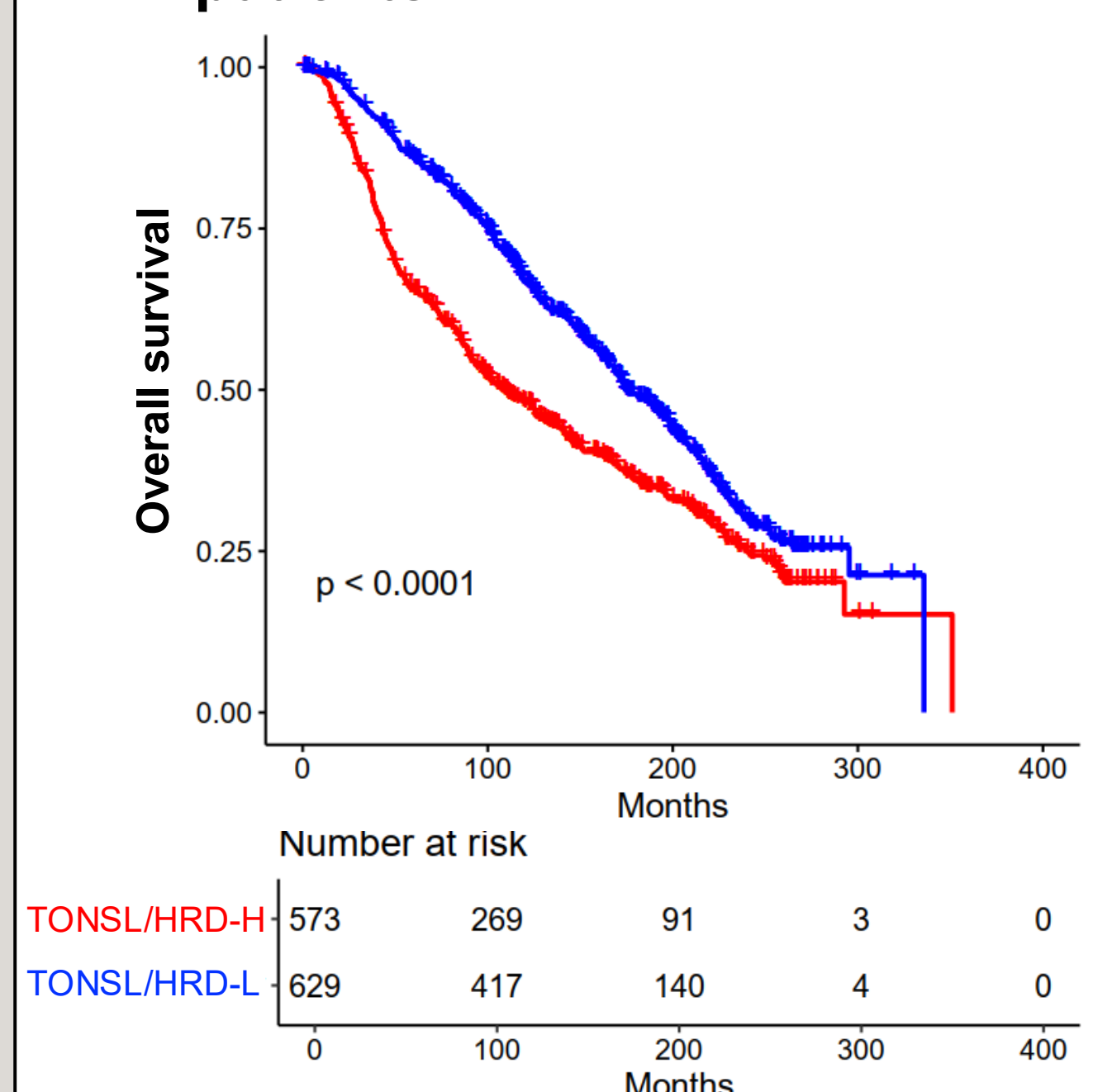
- 10,980 ER+ BC samples tested by NGS (592, NextSeq; WES/WTS, NovaSeq, Caris Life Sciences).
- Dual expression group *TONSL/RAD51*-high (H), *TONSL-H/RAD51*-low (L), *TONSL-L/RAD51*-H, and *TONSL/RAD51*-L were classified by RNA expression above or below median.
- HRD score and *TONSL/HRD*-H/L survival was analyzed using METABRIC (ER+ BC, n = 1904) dataset.
- *TONSL-RAD51* role in sensitivity to CDK4/6i was determined using *TONSL*-amplified ER+ MCF-7 cells.
- Real-world median overall survival (mOS) of *TONSL/RAD51* groups was derived from insurance claims and calculated from biopsy, start of ET or CDK4/6i to last contact using Kaplan-Meier. Statistical significance was assessed using chi-square and Mann-Whitney U with multiple comparison adjustments (q < 0.05).

Figure 1. HRD score in *TONSL*-high (H) compared to *TONSL*-low (L)



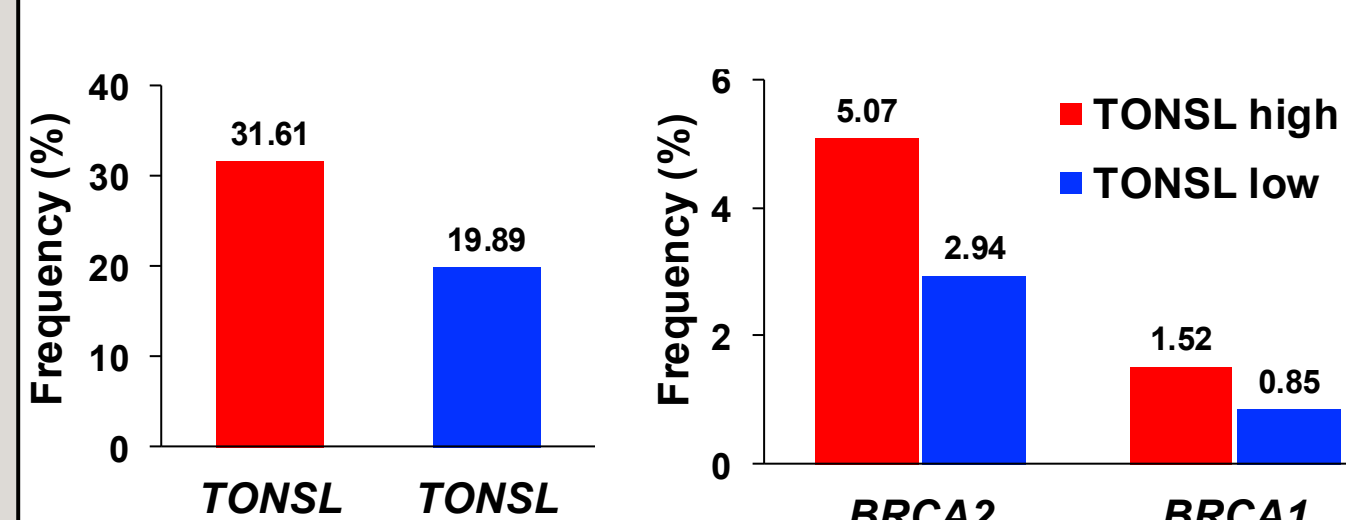
TONSL-H had higher HRD score (33.7 vs 21.1) compared to *TONSL*-L (n = 952 each), p<0.05.

Figure 2. *TONSL/HRD*-H is associated with poor outcome in ER+ patients



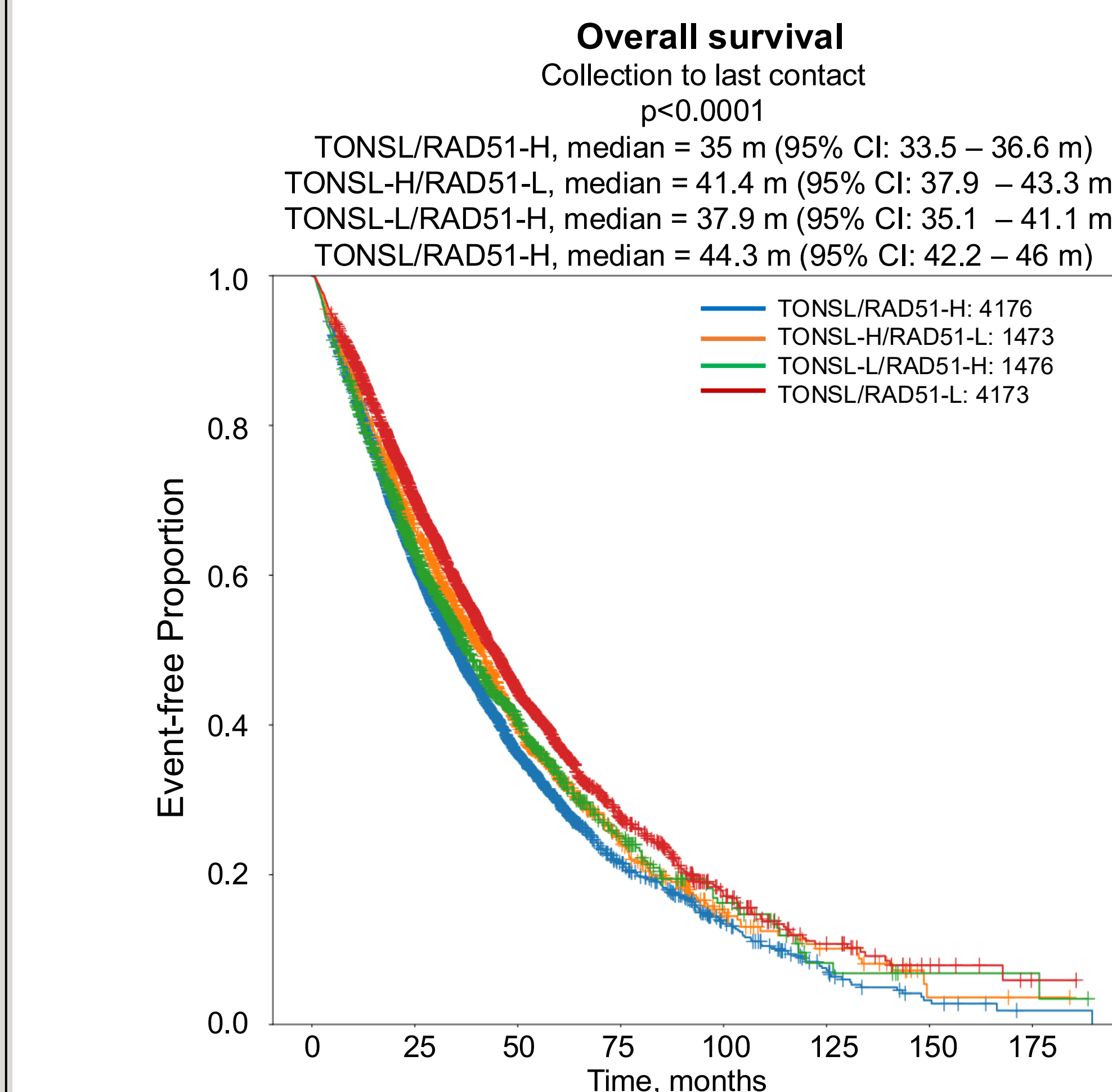
TONSL/HRD-H had worse mOS (109 m vs 138 m, p<0.05) compared to *TONSL/HRD*-L

Figure 3. Frequency of LOH and *BRCA2/1* mutation

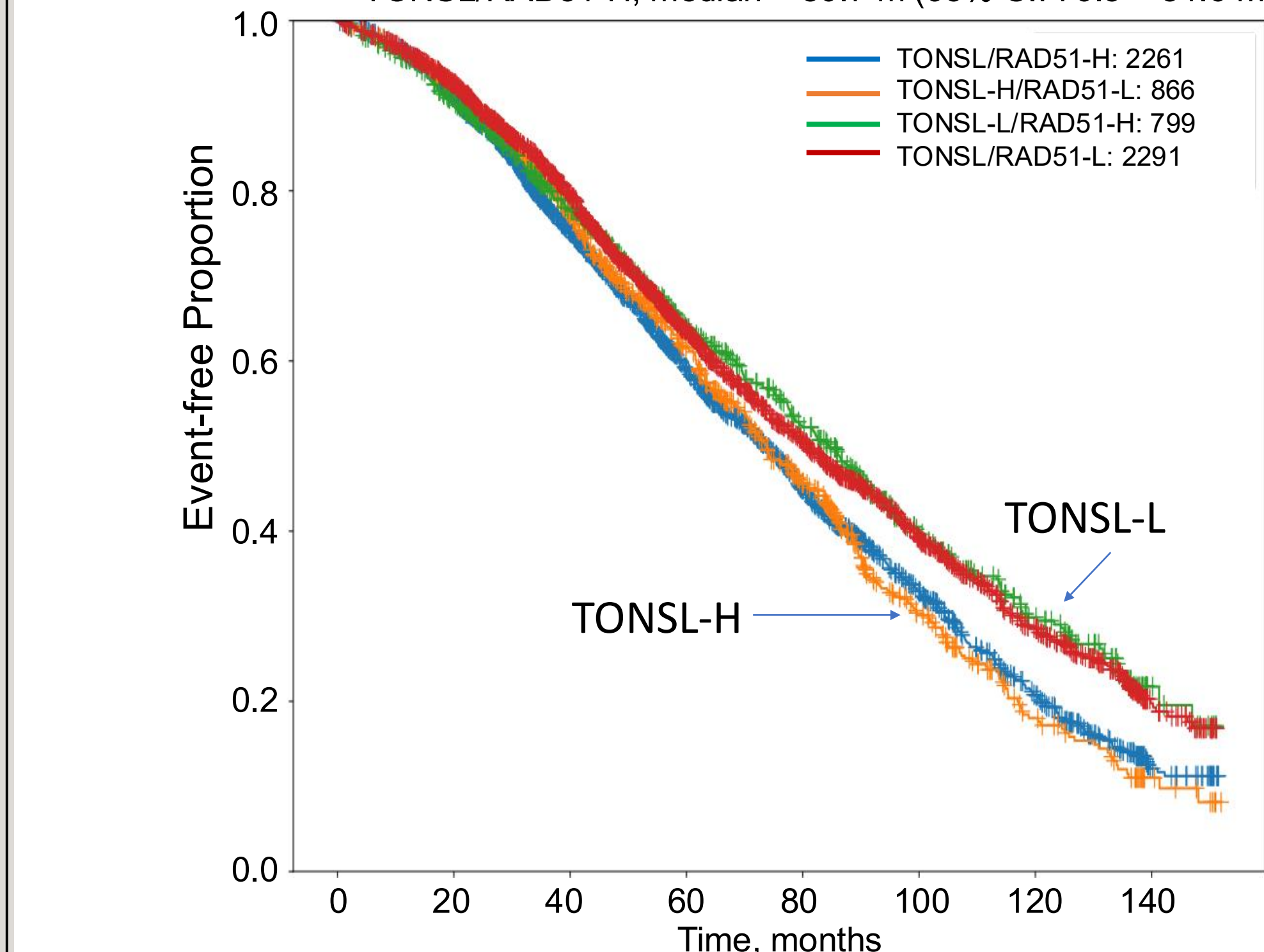
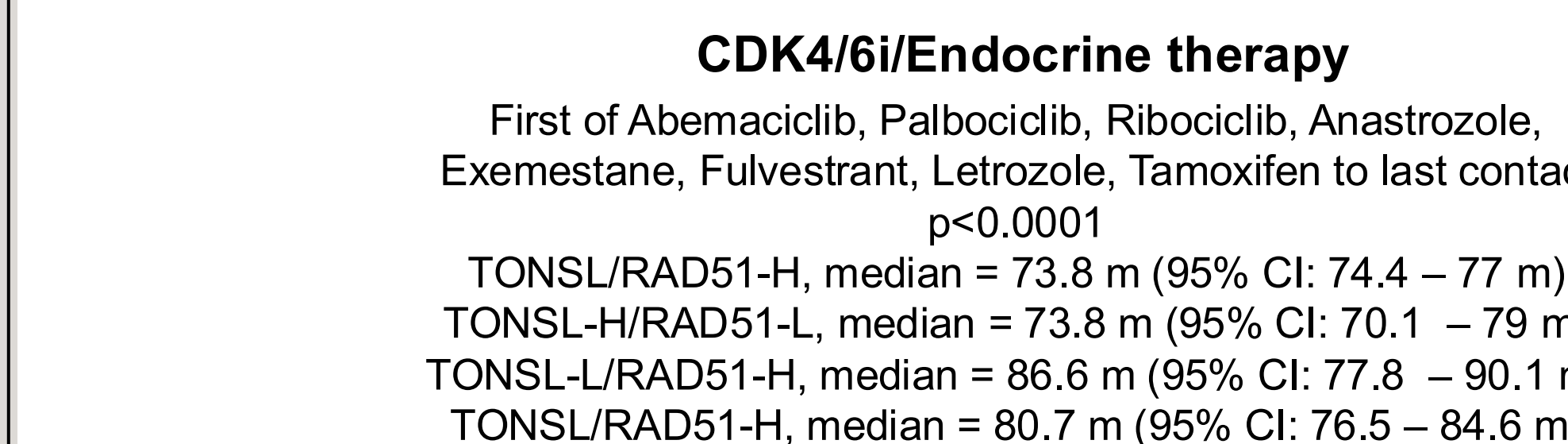


TONSL-H had higher frequency of LOH (31.6% vs 19.8%), *BRCA2* (5.1% vs 2.9%) and *BRCA1* (1.5% vs 0.8%) mutation, all q < 0.05.

Figure 4. ER+ breast cancer patients' survival

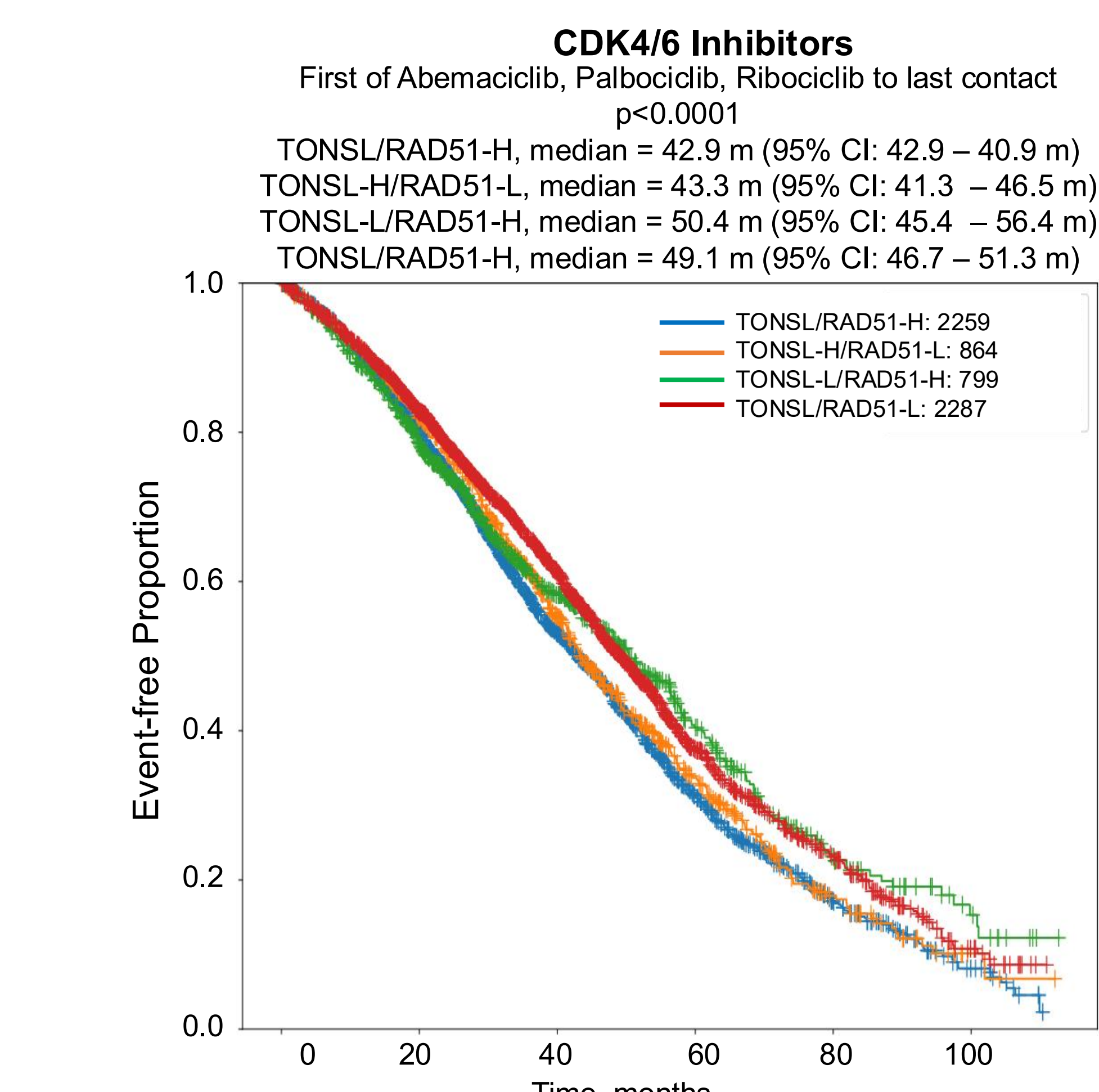


TONSL/RAD51-H	4176	2134	671	206	78	26	7	1
TONSL-H/RAD51-L	1473	886	328	99	35	16	2	1
TONSL-L/RAD51-H	1476	740	227	68	24	6	2	2
TONSL/RAD51-L	4173	2450	818	199	70	26	8	1



TONSL/RAD51-H	2261	1971	1415	947	602	339	164	29
TONSL-H/RAD51-L	866	781	584	388	243	109	45	10
TONSL-L/RAD51-H	799	689	510	348	246	138	74	11
TONSL/RAD51-L	2291	2016	1504	985	659	384	211	40

RESULTS



TONSL/RAD51-H	2259	1624	762	303	93	17
TONSL-H/RAD51-L	864	644	332	133	37	3
TONSL-L/RAD51-H	799	550	282	122	40	11
TONSL/RAD51-L	2287	1667	904	307	117	19

Table 1. Survival by *TONSL* and *RAD51* expression

	TONSL/ RAD51-H	TONSL-H/ RAD51-L	TONSL-L/ RAD51-H	TONSL/ RAD51-L	p-value
Overall	35 m (n=4176)	41.4 m (n=1473)	37.9 m (n=1476)	44.3 m (n=4173)	<0.01
ET	63.4 m (n=1361)	58 m (n=461)	65.2 m (n=508)	76.1 m (n=1430)	<0.01
ET+ CDK4/6i	73.8 m (n=2261)	73.8 m (n=866)	84.6 m (n=799)	80.7 m (n=2291)	<0.01

TONSL/RAD51-H had worse mOS compared to *TONSL-H/RAD51*-L, *TONSL-L/RAD51*-H or *TONSL/RAD51*-L. *TONSL-H/RAD51*-L had worse survival with ET compared to other groups. With ET+CDK4/6i, *TONSL/RAD51*-H and *TONSL-H/RAD51*-L had similar but worse mOS compared to *TONSL-L/RAD51*-H and *TONSL/RAD51*-L.

Figure 5. TMCF7-*RAD51* siRNA effects confirmation

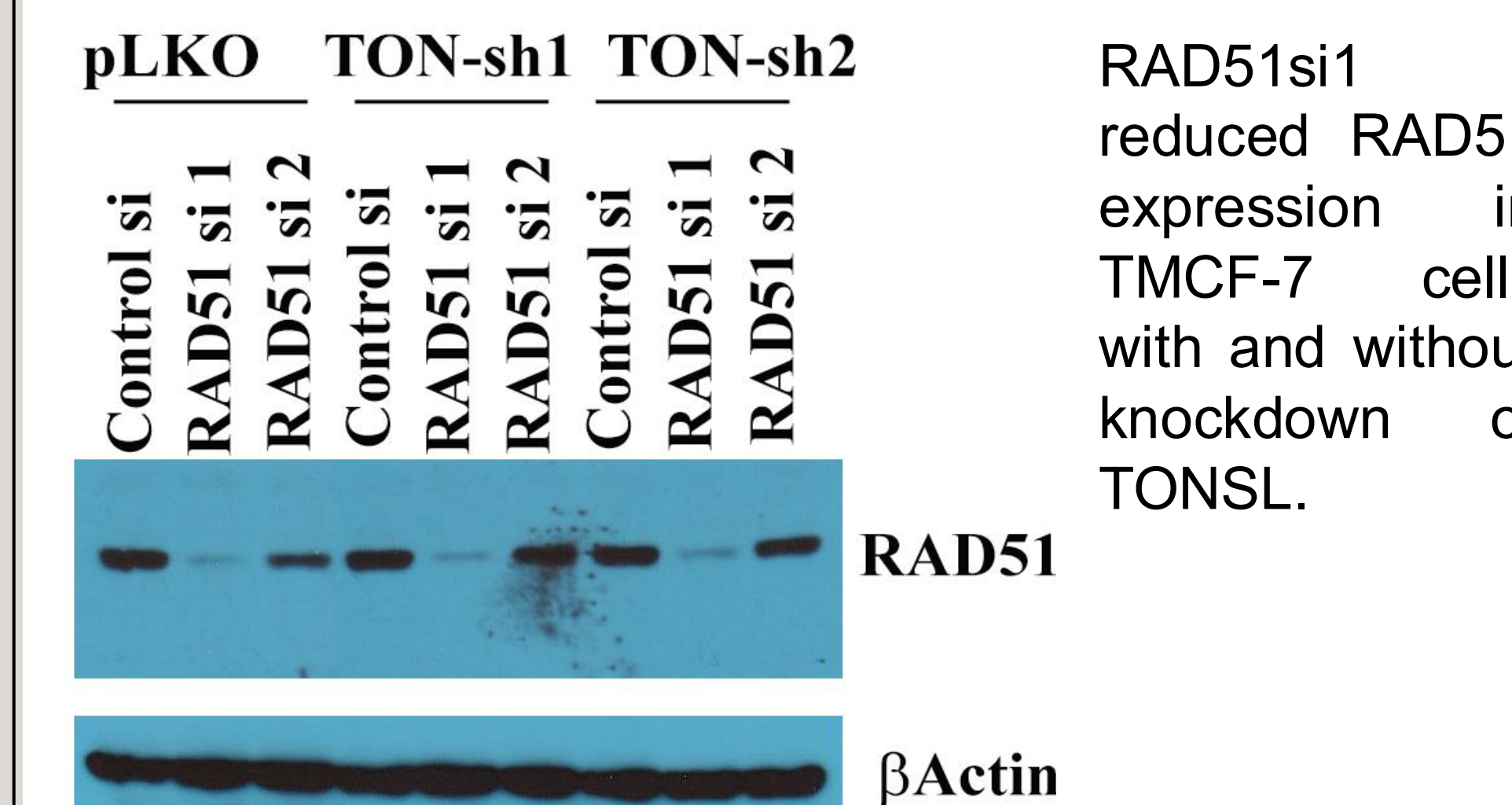
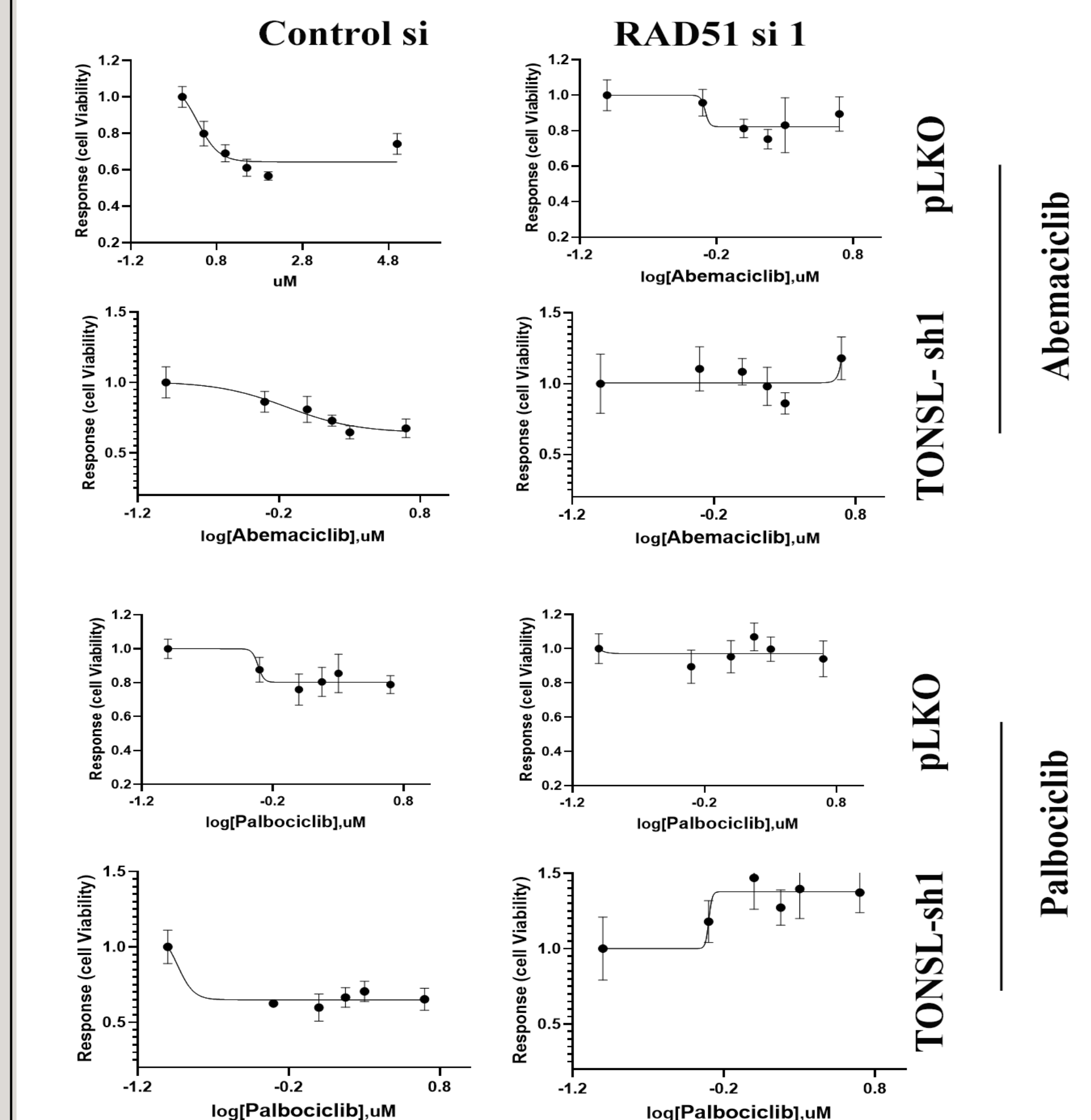


Figure 6. *RAD51* is required for the activity of Abemaciclib and Palbociclib



TONSL knockdown sensitizes TMCF-7 cells to Palbociclib. *RAD51* is required for Palbociclib and Abemaciclib-mediated growth inhibition of TMCF-7 pLKO and TMCF-7 *TONSL*-sh cells. These results are consistent with better response of *TONSL*-L/*RAD51*-H tumors to ET+/CDK4/6i treatment (Table 1).

CONCLUSIONS

- *TONSL*-H independent of *RAD51* expression is associated with poor response to ET or ET+CDK4/6i.
- Therapeutic targeting of *TONSL* to create *TONSL*-L/*RAD51*-H status may improve response to ET+CDK4/6i. Activity of *TONSL* is regulated by various protein complexes, destabilizers of *TONSL*-protein complexes could potentially help in sensitizing ER+ BC to ET or CDK4/6i.