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# Study of DYNLL1 in double-strand break repair and synthetic lethality in BRCA1-deficient breast cancer

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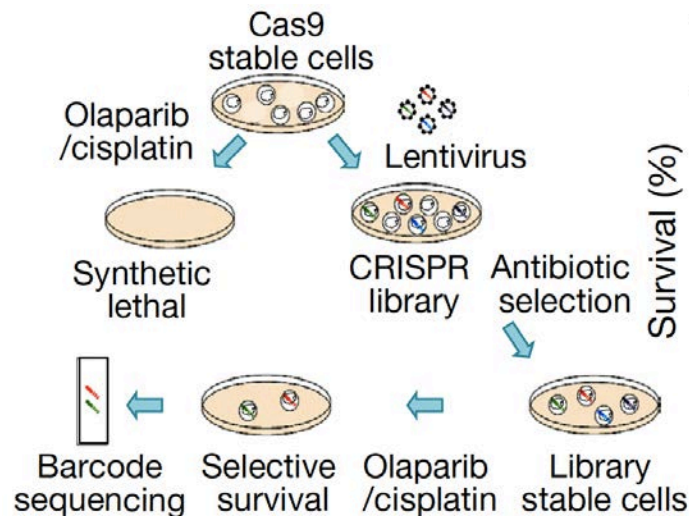
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# A CRISPR-screen to find new genes involved in drug resistance in BRCA1-mutated cells

UWB1.289, COV362 and JHOS-2 cells : BRCA1-mut



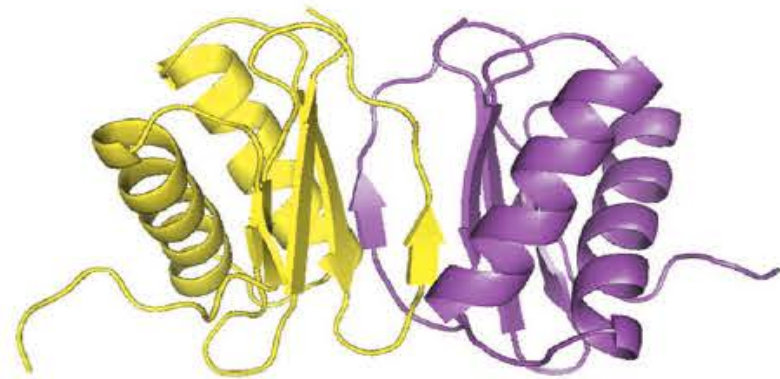
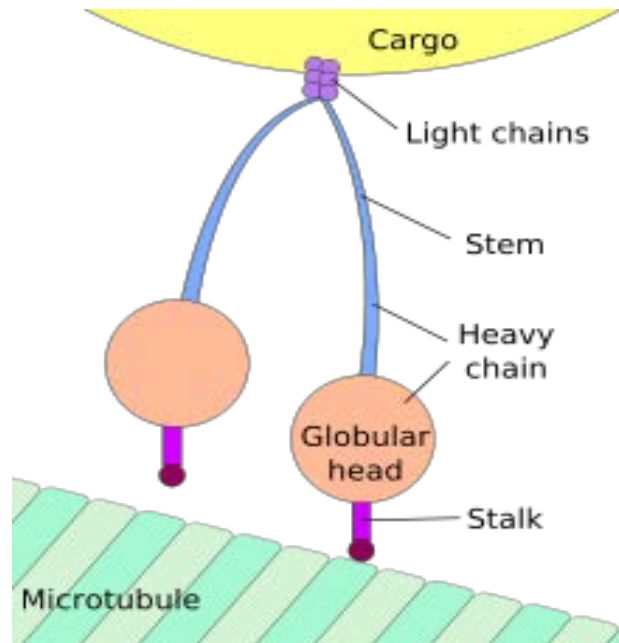
Gene Symbol	STARS Score	p-value
<b>DYNLL1</b>	<b>9,117519799</b>	<b>0</b>
C20orf196	8,947371836	0
TP53BP1	8,895261185	0
<b>ATMIN</b>	<b>7,155399415</b>	<b>6,80E-06</b>
IFNAR1	6,188073532	2,04E-05
IFNAR2	6,159698481	2,04E-05
TEN1	4,886833622	0,000373931
DHX29	4,748660891	0,000482711
NHEJ1	4,431478718	0,000951824

**Olaparib screen**

Gene Symbol	STARS Score	p-value
TAF5L	10,0310281	0
LRR8D	6,94863567	4,5298E-06
EP300	5,6462175	3,6238E-05
TADA2B	4,92430952	0,00020082
<b>DYNLL1</b>	<b>4,81555163</b>	<b>0,00026877</b>
PPFIA4	4,74196526	0,00031256
SUPT20H	4,72601053	0,00032313
DYRK1A	4,43030584	0,00063266
MYBL1	4,41369903	0,00065531
MED12	4,33048023	0,00078366

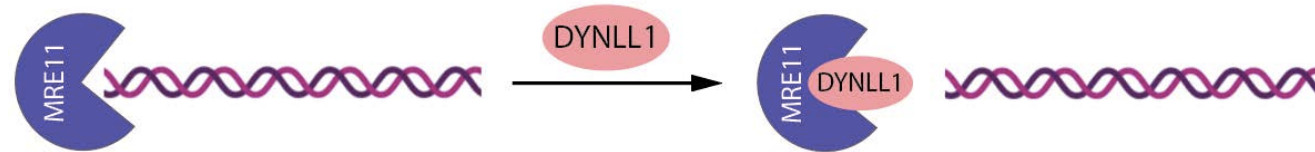
**Cisplatin screen**

# DYNLL1: light chain component of cytoplasmic dynein



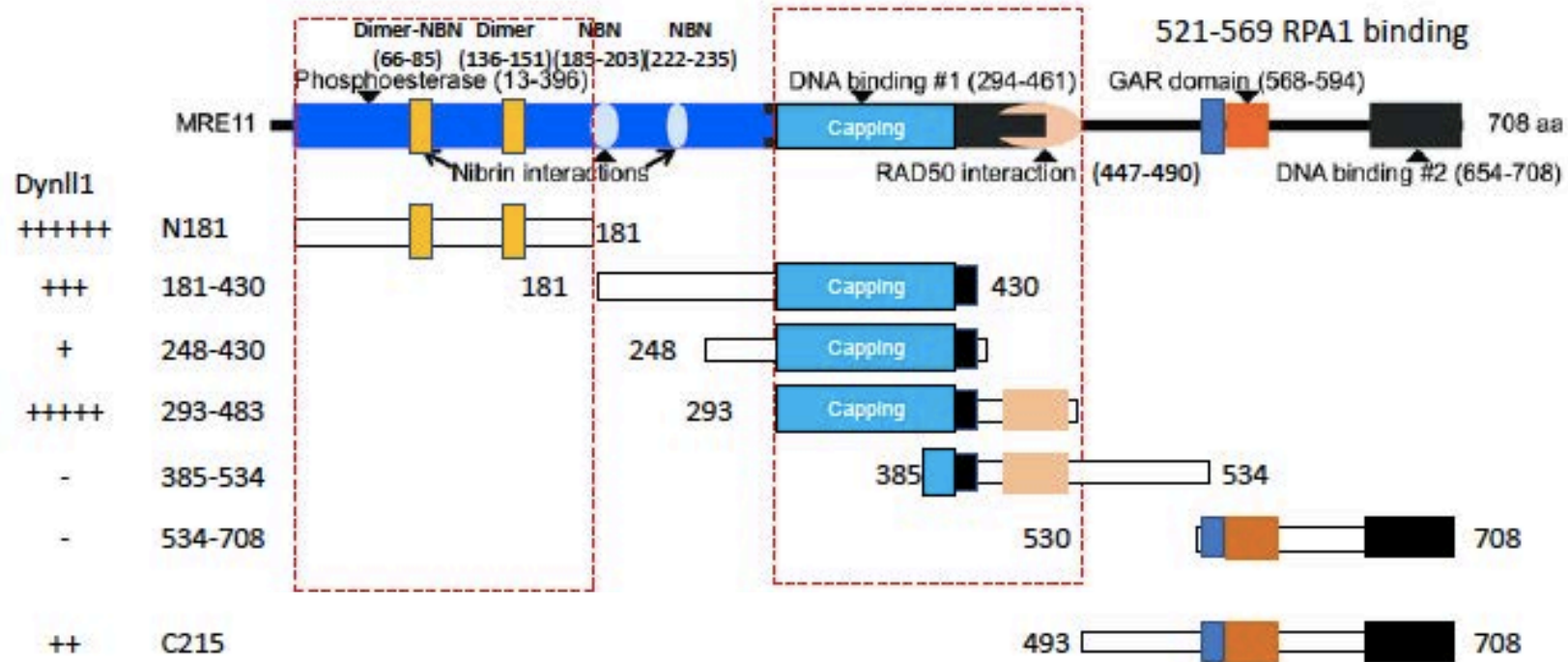
- ❖ Hub protein of **10 kDa**
- ❖ Thought to be involved in linking dynein to cargos
- ❖ May play a role in changing or maintaining the cytoskeletal structures
- ❖ Promotes dimerization of various targets
- ❖ Promotes apoptosis by interacting with BCL2L2
- ❖ **Inhibits resection via its interaction with MRE11**

- To understand the biochemical mechanism by which DYNLL1 prevents DNA-resection, *in vitro*

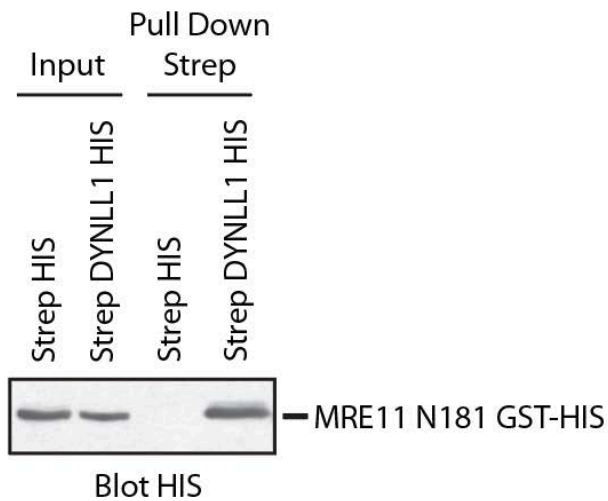


- To determine whether DYNLL1 is a good prognosis marker and a good biomarker for BRCA1-deficient breast cancer therapeutic decision, *in cellulo* and *in vivo*

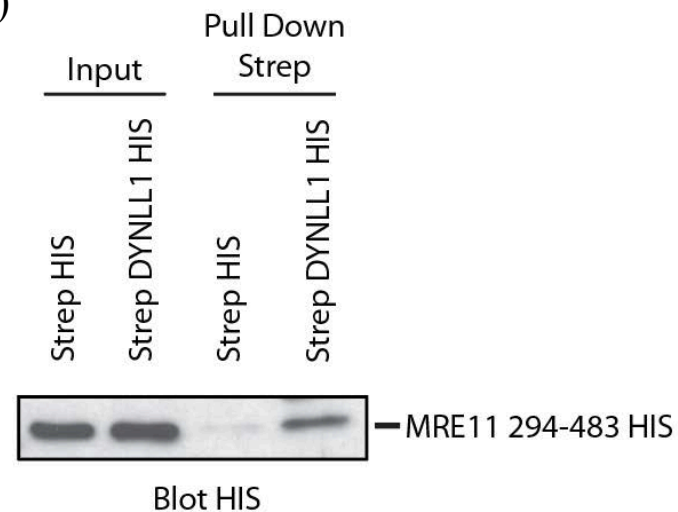
# Fragments of MRE11 bind preferentially DYNLL1 *in cellulo* and *in vitro*



a)



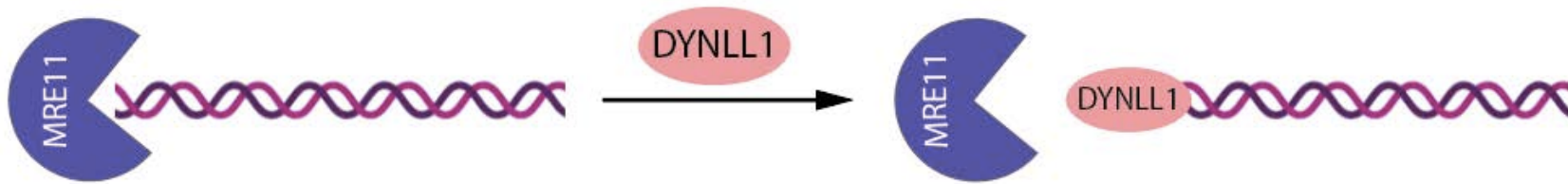
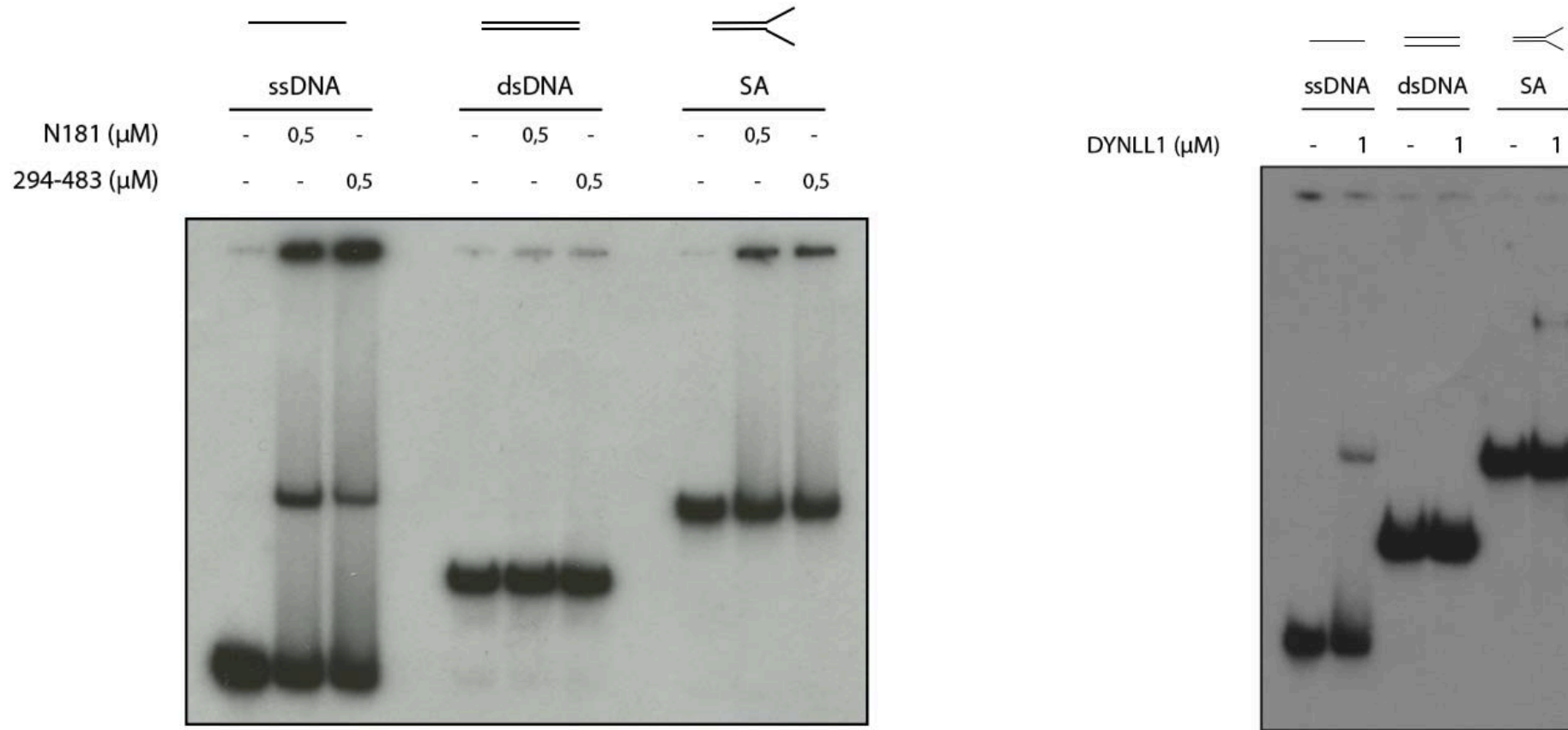
b)



❖ Pulldown *in cellulo* showed that the N181 region and the 294-483 region of MRE11 bind preferentially DYNLL1

❖ Pulldown *in vitro* (co-infection (a) and with purified proteins (b)) show the specific binding of those MRE11 regions to DYNLL1

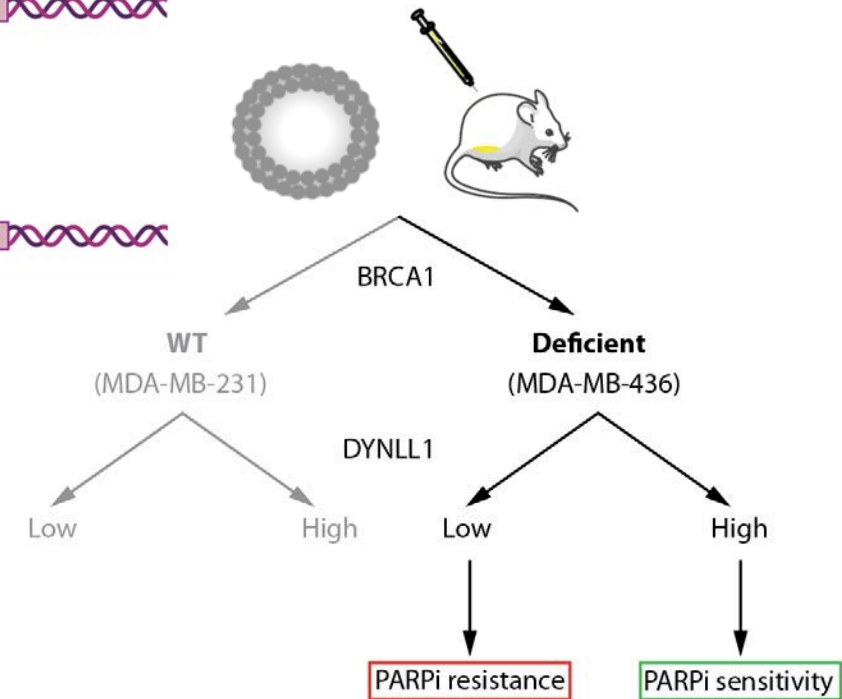
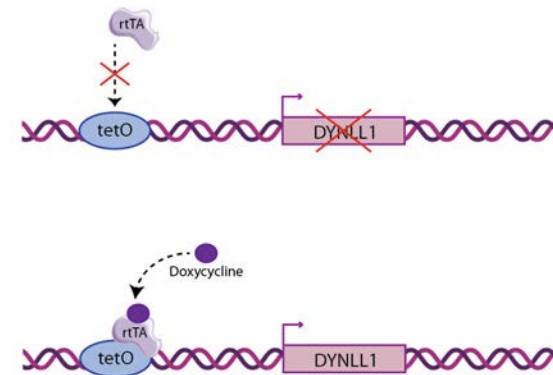
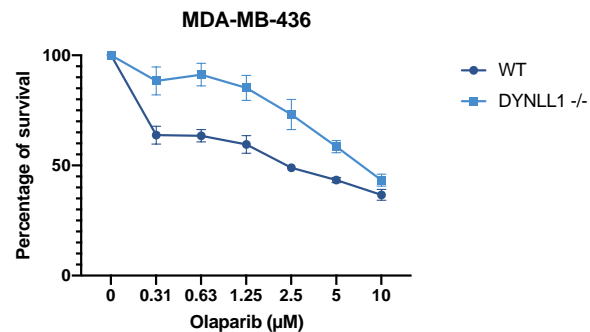
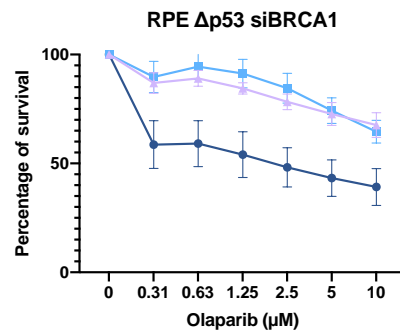
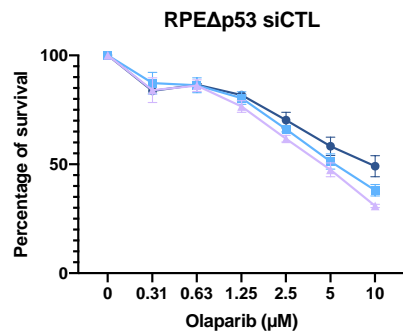
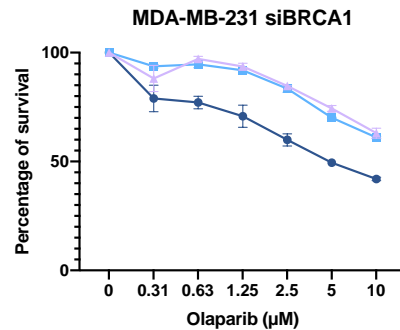
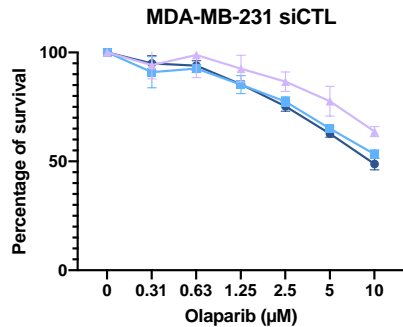
# Fragments of MRE11 and DYNLL1 bind to DNA without fixation



➤ DYNLL1 binds DNA to inhibit resection by MRE11

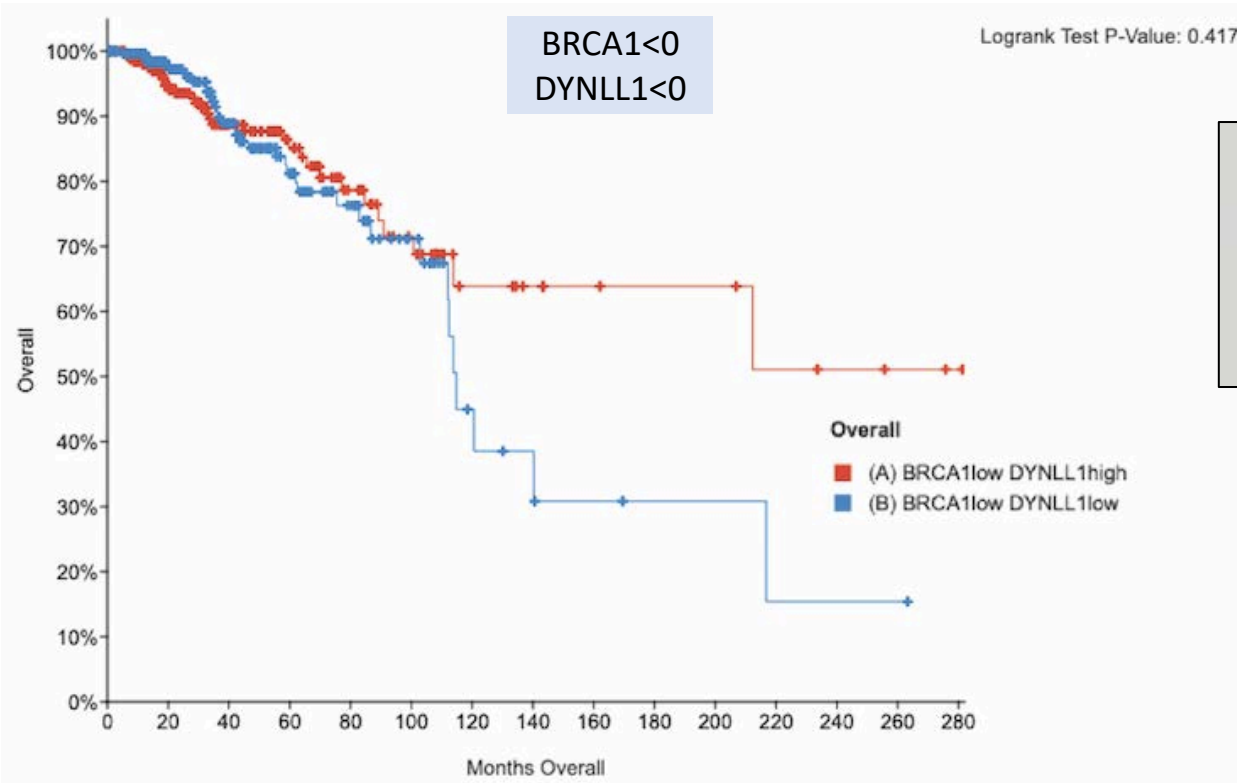
# DYNLL1 knockout leads to PARPi resistance in BRCA1-deficient cells

MDA-MB-231 : BRCA1 WT  
MDA-MB-436 : BRCA1-deficient



- ❖ Cell survival assays show a resistance to the PARPi treatment (Olaparib) only when cells are BRCA1-deficient and DYNLL1-/-
- ❖ We want to induce different levels of DYNLL1 expression in tumours and test the sensitivity to PARPi treatment

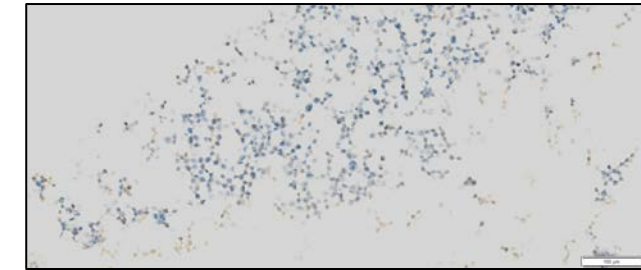
# DYNLL1 is a potential good prognosis marker in BRCA1-deficient cells and tumours



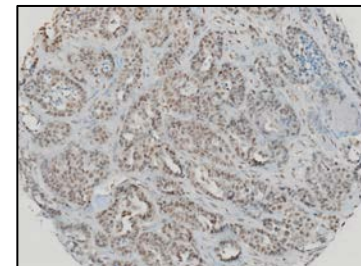
Breast Invasive Carcinoma (TCGA, Firehose Legacy)



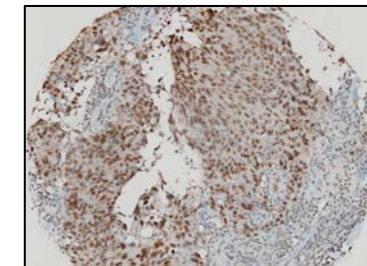
RPE  $\Delta$ p53 WT



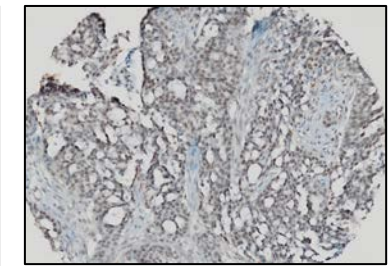
RPE  $\Delta$ p53 DYNLL1 -/-



Ovarian tumour 1



Ovarian tumour 2



Ovarian tumour 3

- ❖ Patient data show a better overall survival when DYNLL1 is more expressed in a BRCA1<sup>low</sup> context
- ❖ DYNLL1 antibody is very specific in immunohistochemistry and can be used further on TMA

➤ BRCA1<sup>low</sup> DYNLL1<sup>low</sup> → HR reactivation → PARPi resistance



- ❖ We aim to determine how DYNLL1 regulates DNA resection via understanding biochemically its interaction with MRE11 ;
- ❖ Understanding the role of DYNLL1 in PARPi resistance using different models (2D, 3D, *in vivo*) will allow us to find better ways to counteract this resistance ;
- ❖ Establishing DYNLL1 as a biomarker for the therapeutic choice of *BRCA1*-deficient cancers could provide a more precise approach for personalized treatment with PARPi

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He YJ, Meghani K, Caron M-C, Yang C, Ronato DA, Bian J, et al. DYNLL1 binds to MRE11 to limit DNA end resection in BRCA1-deficient cells. *Nature*. nov 2018;563(7732):522-6.

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Becker JR, Cuella-Martin R, Barazas M, Liu R, Oliveira C, Oliver AW, et al. The ASCIZ-DYNLL1 axis promotes 53BP1-dependent non-homologous end joining and PARP inhibitor sensitivity. *Nat Commun*. déc 2018;9(1):5406.

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**Thank you for your time!**