

Discovery of ERD-3111 as a Potent and Orally Efficacious Estrogen Receptor PROTAC Degrader with Strong Antitumor Activity



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Introduction

Estrogen receptor α (ER α) is a prime target for the treatment of ER+ breast cancer. Despite the development of several effective therapies targeting ER α signaling, clinical resistance remains a major challenge. In addition to oral selective estrogen receptor degraders (SERDs), another promising therapeutic strategy to overcome endocrine resistance in ER+ breast cancer is the development of ER degraders using the proteolysis targeting chimera (PROTAC) technology.

However, identifying highly potent and orally efficacious ER α PROTACs is difficult due to their relatively larger molecular weights and poorer physicochemical properties. Extensive research has led to the discovery of ARV-471 as an orally bioavailable ER PROTAC degrader. These preclinical and initial clinical data for ARV-471 suggested that oral ER α PROTACs may have a promising therapeutic potential for the treatment of ER+ human breast cancer.

Herein, we describe the design, synthesis, and biological evaluation of new classes of ER α PROTACs based on a new cereblon ligand and three classes of ER ligands. This work led to the discovery of **ERD-3111** as a potent, orally bioavailable, and highly efficacious ER α PROTAC that effectively inhibits the *in vivo* growth of breast cancers with either wild-type or mutated ER in mice.

Result

Table 1. Profiling of new CCRN ligand TX-16 and its analogue 9 (TX-16-Me)

CCRN Binding affinity IC ₅₀ (μ M)		Caco-2 Permeability of Compound 9		Plasma Protein Binding of Compound 9 (%)	
TX16 (9a)	Lenalidomide	Thalidomide	P _{app} (10^{-6} cm/s)	ER	human/dog/monkey/rat/mouse
2.6	3.6	2.9	7.4	1.1	12.5/17.0/14.4/36.5/26.3
Rat Pharmacokinetic Profile of Compound 9					
IV/PO (mg/kg)	V _{ss} (L/kg)	Cl (mL/min/kg)	T _{1/2} (h)	C _{max} (ng/mL)	AUC (h·ng/mL)
1/3	5.0	29.2	1.8	228.4	1147
					F (%)
					70

Figure 1. Design of new classes of oral ER PROTACs based on CCRN ligand TX-16 and different classes of ER ligands

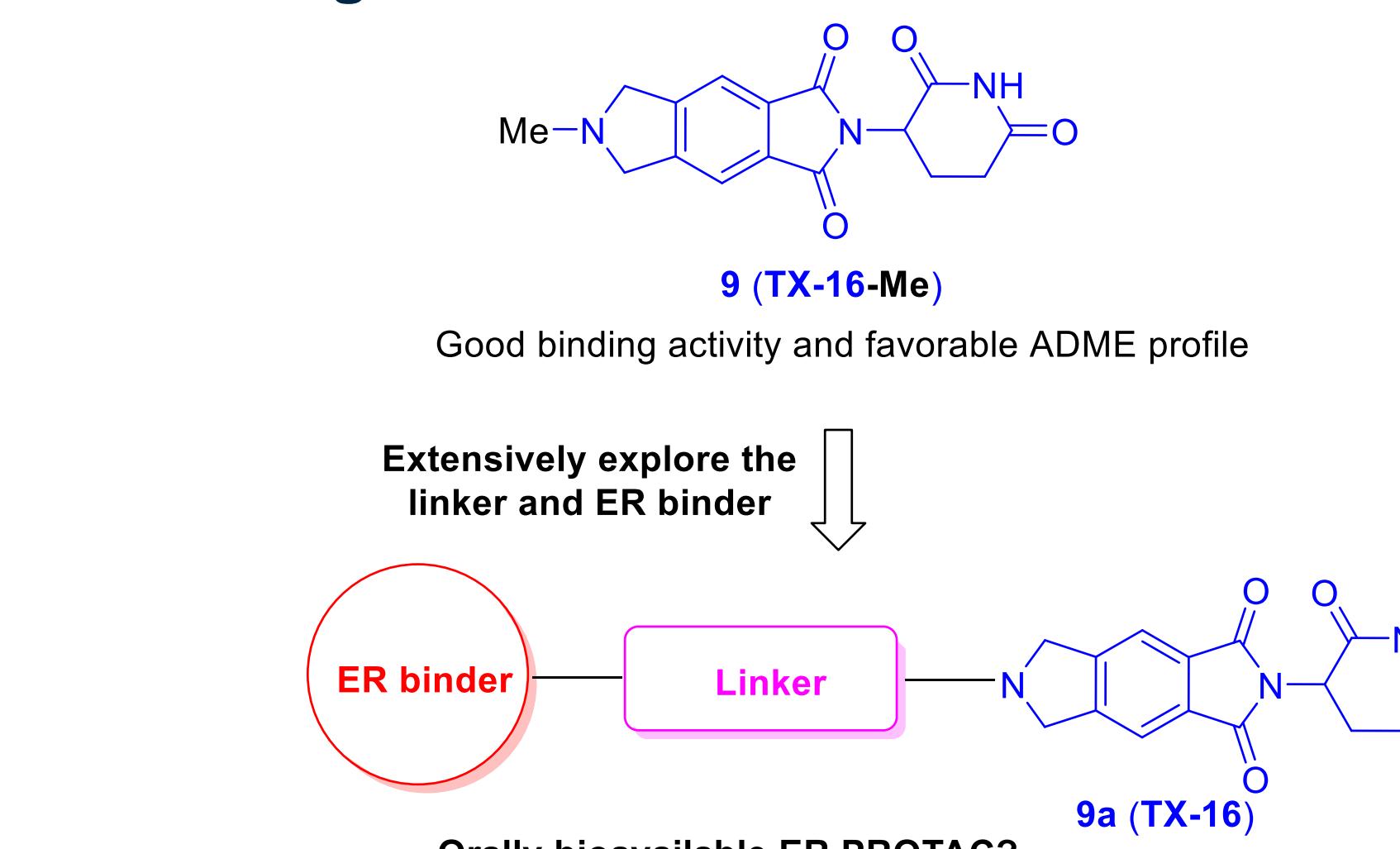


Table 2. Determination of optimal linker length based on ER binder in ARV-471 and TX16

Compound	Linker length (n)	ER α degradation	
		DC ₅₀ (nM)	D _{max} (%)
Fulvestrant	N.A.	0.9 ± 0.2	100 ± 3
ARV-471	N.A.	0.4 ± 0.04	89 ± 1
10	0	>1000	32 ± 5
11	1	415 ± 75	65 ± 8
12	2	>1000	50 ± 4
13	3	171 ± 48	79 ± 12
14	4	236 ± 40	67 ± 6
15	5	31 ± 4	106 ± 4
16	6	129 ± 15	101 ± 6

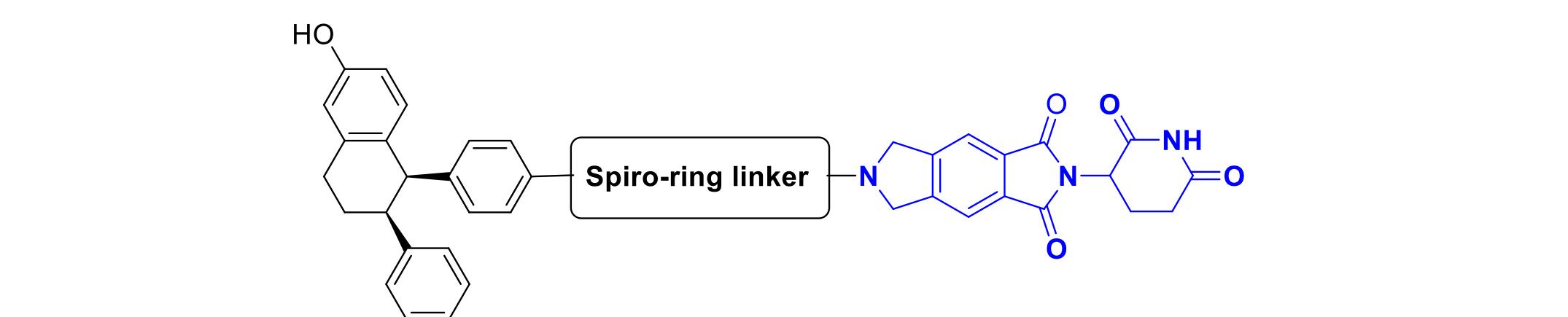
Table 3. Rigidification of linker to enhance the potency

Compound	Linker	ER α degradation ^a	
		DC ₅₀ (nM) ^b	D _{max} (%) ^c
17		989 ± 237	50 ± 2
18		21 ± 7	59 ± 3
19		2.3 ± 0.7	84 ± 5
20		4.3 ± 1.0	79 ± 3
21		295 ± 94	57 ± 2
22		19 ± 7	64 ± 5
23		6.0 ± 1.4	84 ± 4
24		1.6 ± 0.3	95 ± 4
25		13 ± 3.2	76 ± 4
26		9.7 ± 3.4	63 ± 4
27		32 ± 6.3	87 ± 5
28		>1000	40 ± 8

Table 4. The PK profiles for compounds 19, 20, 24, 36 in rats and/or mice

Cpd	Species	IV/PO (mg/kg)	V _{ss} (L/kg)	Cl (mL/min/kg)	T _{1/2} (h)	C _{max} (ng/mL)	AUC (h·ng/mL)	F (%)
19	Rat	1/3	8.8	25.7	4.0	54.6	552	30
	mouse	1/3	16.2	14.2	9.8	93.3	1482	58
20	Rat	1/3	55.4	17.0	16.0	24.9	399	36
24	Rat	1/3	2.0	23.1	2.7	40.6	206	10
36	Rat	1/3	1.6	20.0	2.5	69.5	328	13
	Mouse	1/3	2.2	31	3.1	146	550	35

Table 5. Improvement of the PK profiles by employing more rigid spiro-ring-containing linkers



Cpd	Spiro-ring Linker	ER α degradation	
		DC ₅₀ (nM)	D _{max} (%)
29		11 ± 3.4	65 ± 3
30		>1000	23 ± 21
31		4.0 ± 1.5	63 ± 3
32		1.8 ± 0.6	68 ± 3
33		2.8 ± 0.9	71 ± 3
34		18 ± 6.3	58 ± 2
35		4.6 ± 1.2	75 ± 3
36		5.5 ± 1.2	90 ± 4
37		393 ± 165	58 ± 8
38		>1000	16 ± 8

Table 6. Replacing the ER core of compound 36 with tricyclic indole cores

Cpd	R	ER α degradation		Oral plasma exposure in rat ^a (Drug concentration, ng/mL)			
		DC ₅₀ (nM)	D _{max} (%)	1 h	3 h	6 h	24 h
39		0.1 ± 0.02	97 ± 3	9.3 ± 4.1	6.6 ± 3.2	3.0 ± 1.0	N/A
40		0.8 ± 0.18	83 ± 3	7.9 ± 4.3	8.2 ± 3.4	8.3 ± 3.5	4.0 ± 0.2
41		0.8 ± 0.11	106 ± 2	8.3 ± 0.4	14.0 ± 1.5	10.3 ± 4.9	

^aDose: compounds 39-40 (3 mg/kg); compound 41 (5 mg/kg).

Table 7. Replacing the ER core of compound 36 with tricyclic indazole cores

Cpd	X	R ₁	R ₂	ER α degradation	
				DC ₅₀ (nM)	D _{max} (%)
42	N	CF ₃	H	3.8 ± 0.6	107 ± 4
43	C-F	CF ₃	F	2.0 ± 0.3	88 ± 2
44 ^a (ERD-3111)	C-F	CHF ₂	F	0.5 ± 0.04	91 ± 1
45	N	CHF ₂	H	14 ± 3.0	112 ± 6
46	C-H	CHF ₂ </td			