

# Supplementary Figure 11

<b>A.</b>	<b>Cell Line</b>	<b>p53 Mutation</b>	<b>p53 Aggregation Status</b>
	Kuramochi	D281Y	Unknown
	OVCAR8	Y126_K132del	Unknown
	OVCAR4	L130V	Unknown
	OVCAR3	R248Q	Aggregating <sup>†</sup>
	OAW28	P152fs	Unknown
	SKOV3	P89fs	Unknown
	SNU-119	P151A	Unknown
	CaOV3	Q136*	Unknown

<sup>†</sup> Xu J, Reumers J, Couceiro JR, De Smet F, Gallardo R, Rudyak S, et al. Gain of function of mutant p53 by coaggregation with multiple tumor suppressors. Nat Chem Biol. 2011 May;7(5):285–95.

<b>B.</b>	<b>Patient Tumor Sample</b>	<b>p53 Mutation</b>	<b>p53 Aggregation Status</b>
	HGSOC1	R273H	Aggregating <sup>†</sup>
	HGSOC2	W53*	Unknown
	HGSOC3	C238Y	Unknown
	HGSOC4	E204*	Unknown
	HGSOC5	R273H	Aggregating <sup>†</sup>
	HGSOC6	V157G	Unknown
	HGSOC7	R273C	Suggested <sup>§</sup>
	HGSOC8	S127F	Unknown
	HGSOC9	R342fs	Unknown
	HGSOC10	R248Q	Aggregating <sup>†</sup>

<sup>†</sup> Xu J, Reumers J, Couceiro JR, De Smet F, Gallardo R, Rudyak S, et al. Gain of function of mutant p53 by coaggregation with multiple tumor suppressors. Nat Chem Biol. 2011 May;7(5):285–95.

<sup>‡</sup> Levy CB, Stumbo AC, Ano Bom AP, Portari EA, Cordeiro Y, Silva JL, et al. Co-localization of mutant p53 and amyloid-like protein aggregates in breast tumors. Int J Biochem Cell Biol. 2011 Jan;43(1):60–4.

<sup>§</sup> Palanikumar L, Karpauskaite L, Al-Sayegh M, Chehade I, Alam M, Hassan S, et al. Protein mimetic amyloid inhibitor potently abrogates cancer-associated mutant p53 aggregation and restores tumor suppressor function. Nat Commun. 2021 Jun 25;12(1):3962.

**Supplementary Figure 11. TP53 mutations and predicted aggregation statuses of all ovarian cancer cell lines and patient HGSOC specimens tested using the in vitro organoid drug assay.** The p53 mutation status and aggregation potential for **(A)** ovarian cancer cell lines and **(B)** primary patient HGSOC tumor samples tested using the in vitro organoid drug assay. TP53 mutation statuses for cell lines were obtained from the IARC TP53 database. Patient TP53 mutations were determined by clinical sequencing or by whole exome sequencing. The aggregating status for some TP53 mutations has previously been suggested or demonstrated experimentally other investigators. The aggregation potential for vast majority of mutations in p53 are unknown.