

Supplementary Materials

A Molecular Stratification of Chilean Gastric Cancer Patients with Potential Clinical Applicability

Mauricio P. Pinto, Miguel Córdova-Delgado, Ignacio N. Retamal, Matías Muñoz-Medel, M. Loreto Bravo, Doris Durán, Francisco Villanueva, César Sanchez, Francisco Acevedo, Sebastián Mondaca, Érica Koch, Carolina Ibañez, Héctor Galindo, Jorge Madrid, Bruno Nervi, José Peña, Javiera Torres and Marcelo Garrido

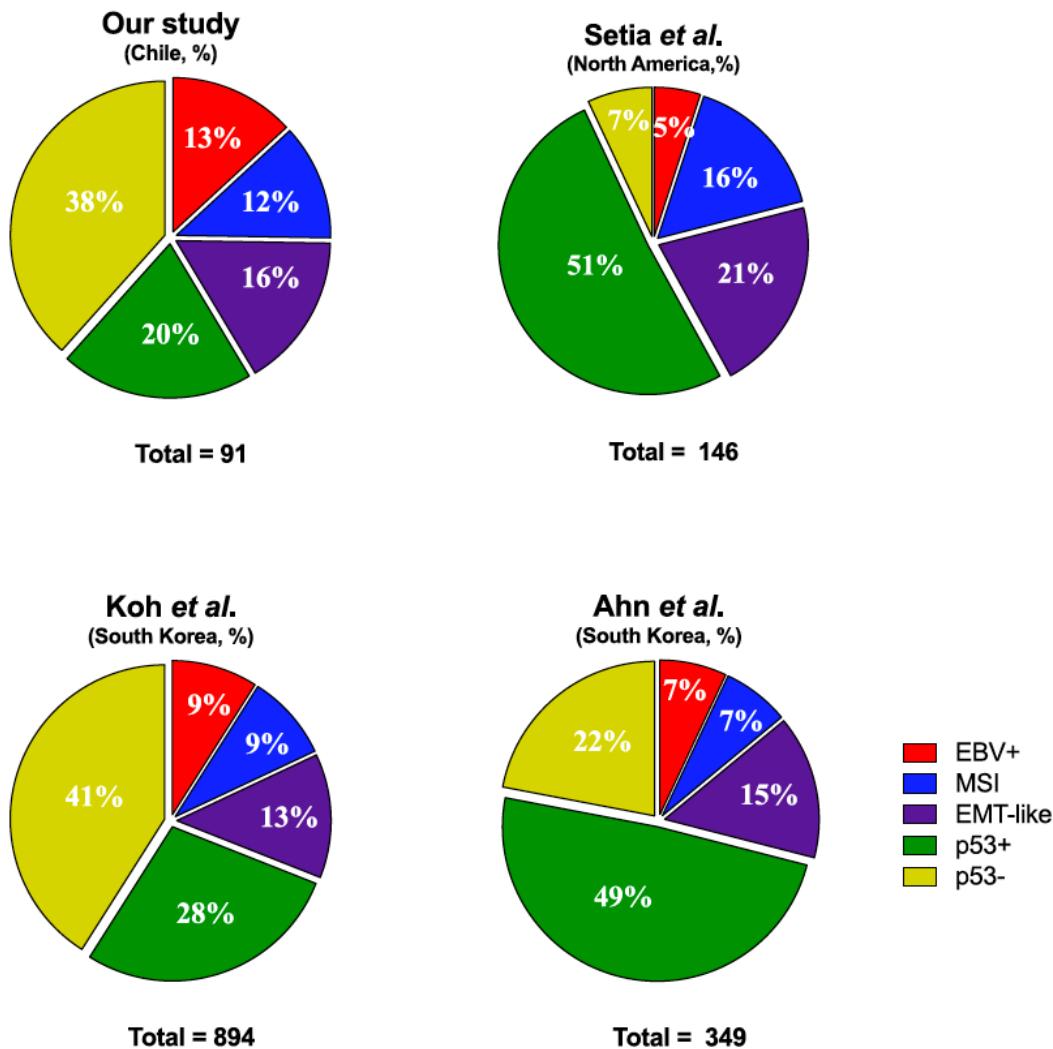
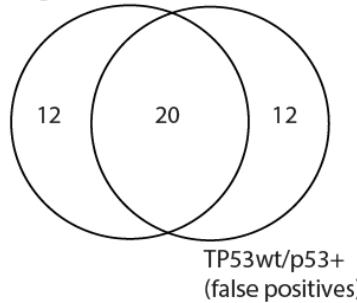


Figure S1. Distribution of gastric cancer molecular subtypes among Chilean, North American or Asian studies.

≥10% and 2+ or 3+

TP53mut/p53-
(false negatives)

ID	P53_NGS		P53_IHC	
	Kind of mutation	AA change	% cel	Intensity (1-3)
GCTF00003T	nonsense	p.Arg342*	0	0
GCTF00005T	nonsense	p.Arg306*	0	0
GCTF00016T	nonsense	p.Arg196*	0	0
GCTF00020T	frameshift/insertion	p.Ser240fs	0	0
GCTF00044T	nonsense	p.Arg213*	0	0
GCTF00053T	missense	p.Ser183Leu	10	1
GCTF00068T	nonsense	p.Arg213*	0	0
GCTF00072T	missense	p.Tyr220His	20	1
GCTF00082T	missense	p.Gly154Ser	0	0
GCTF00096T	frameshift/insertion	p.Tyr327fs	10	1
GCTF00108T	frameshift/insertion	p.Pro223fs	0	0
GCTF00119	frameshift/insertion	p.Arg158fs	0	0

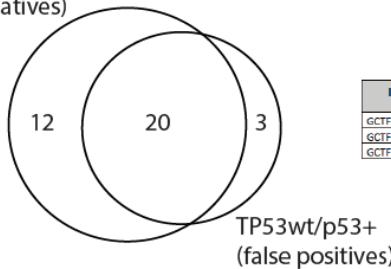


ID	P53_IHC		P53_NGS	
	% cel	Intensity (1-3)	% cel	Intensity (1-3)
GCTF00027	40	2	Wild type	
GCTF00031	10	2	Wild type	
GCTF00038	10	3	Wild type	
GCTF00055	10	3	Wild type	
GCTF00090	90	3	Wild type	
GCTF00103	10	2	Wild type	
GCTF00111	10	3	Wild type	
GCTF00114	60	2	Wild type	
GCTF00115	10	3	Wild type	
GCTF00116	10	2	Wild type	
GCTF00118	10	2	Wild type	
GCTF00125	10	2	Wild type	

≥20% and 2+ or 3+

TP53mut/p53-
(false negatives)

ID	P53_NGS		P53_IHC	
	Kind of mutation	AA change	% cel	Intensity (1-3)
GCTF00003T	nonsense	p.Arg342*	0	0
GCTF00005T	nonsense	p.Arg306*	0	0
GCTF00016T	nonsense	p.Arg196*	0	0
GCTF00020T	frameshift/insertion	p.Ser240fs	0	0
GCTF00044T	nonsense	p.Arg213*	0	0
GCTF00053T	missense	p.Ser183Leu	10	1
GCTF00088T	nonsense	p.Arg213*	0	0
GCTF00072T	missense	p.Tyr220His	20	1
GCTF00082T	missense	p.Gly154Ser	0	0
GCTF00096T	frameshift/insertion	p.Tyr327fs	10	1
GCTF00108T	frameshift/insertion	p.Pro223fs	0	0
GCTF00119	frameshift/insertion	p.Arg158fs	0	0

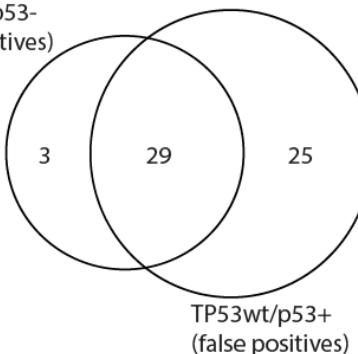


ID	P53_IHC		P53_NGS	
	% cel	Intensity (1-3)	% cel	Intensity (1-3)
GCTF00027	40	2	Wild type	
GCTF00090	90	3	Wild type	
GCTF00114	60	2	Wild type	

**≥20% and 2+ or 3+ or 0% and 0+ (complete loss)
like Setia et al.**

TP53mut/p53-
(false negatives)

ID	P53_NGS		P53_IHC	
	Kind of mutation	AA change	% cel	Intensity (1-3)
GCTF00053T	missense	p.Ser183Leu	10	1
GCTF00072T	missense	p.Tyr220His	20	1
GCTF00096T	frameshift/insertion	p.Tyr327fs	10	1



ID	P53_IHC		P53_NGS	
	% cel	Intensity (1-3)	% cel	Intensity (1-3)
GCTF00001	9	0	Wild type	
GCTF00005	0	0	Wild type	
GCTF00007	9	0	Wild type	
GCTF00010	0	0	Wild type	
GCTF00016	0	0	Wild type	
GCTF00023	0	0	Wild type	
GCTF00024	0	0	Wild type	
GCTF00027	40	2	Wild type	
GCTF00028	9	0	Wild type	
GCTF00033	0	0	Wild type	
GCTF00040	0	0	Wild type	
GCTF00043	0	0	Wild type	
GCTF00053	0	0	Wild type	
GCTF00042	0	0	Wild type	
GCTF00046	0	0	Wild type	
GCTF00047	0	0	Wild type	
GCTF00075	0	0	Wild type	
GCTF00086	0	0	Wild type	
GCTF00090	90	3	Wild type	
GCTF00096	0	0	Wild type	
GCTF00112	0	0	Wild type	
GCTF00114	0	0	Wild type	
GCTF00121	0	0	Wild type	
GCTF00127	0	0	Wild type	
GCTF00130	0	0	Wild type	

Figure S2. Comparison of p53 false positive/negatives using different criteria.

Table S1. clinico-pathological characteristics in *TP53* Wild type or mutant patients.

Characteristic	<i>TP53</i> -WT	<i>TP53</i> -Mut
Median OS: months (range)	40 (26–52)	23 (13–53)
Males; n (%)	12 (57)	13 (65)
Age (median, IQT)	64 (58–77)	63 (54–67)
Stage; n (%)		
I/II	4 (19)	8 (40)
III	14 (67)	9 (45)
IV	3 (14)	3 (20)
Primary tumor; n (%)		
Proximal	3 (15)	7 (39)
Medial	9 (45)	4 (22)
Distal	7 (35)	6 (33)
Multiple	1 (5)	1 (6)
IHC status; n (%)		
HER2+	1 (5)	8 (40)
PDL1+	3 (14)	3 (15)
P53+	8 (38)	14 (70)
Signet ring cell; n (%)		
No	12 (57)	14 (70)
<50%	6 (29)	5 (25)
≥50%	3 (14)	1 (5)
Lauren histotype; n (%)		
Diffuse	9 (50)	4 (22)
Intestinal	7 (39)	11 (61)
Mixed	2 (11)	3 (17)

IQT interquartile range, IHC immunohistochemistry.

Table S2. Response assessment to first line peri-operative/palliative chemotherapy by RECIST v1.1 (n=32).

RECIST v1.1	MMR-D n = 2	EBV+ n = 5	<i>TP53</i> -WT n = 8	<i>TP53</i> -Mut n = 8	EMT-like n = 9	Total n = 32	p- Value
CR/PR	0	4 (80)	3 (37.5)	5 (62.5)	6 (66.7)	20 (56)	0.31
SD/PD	2 (100)	1 (20)	5 (62.5)	3 (37.5)	3 (33.3)	16 (44)	

CR complete response, PR partial response, SD stable disease, PD progressive disease.



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