

SUPPLEMENTARY MATERIAL:

Tables S1.1 (coding of variables) and S1.2 (multivariate analysis): Additional multivariate analysis of all clinicopathologic features associated to pCR in the overall cohort.

Codificaciones de variables categóricas

		Frecuencia	Codificación de parámetro			
			(1)	(2)	(3)	(4)
Combinada homogeneidadTILs	homogeni <= 20%TILs	190	,000	,000	,000	,000
	Homogeni >20% TILs	94	1,000	,000	,000	,000
	Heterogeni <=20%TILs	78	,000	1,000	,000	,000
	Heterogeni>20%TILs	87	,000	,000	1,000	,000
	No infiltrat	26	,000	,000	,000	1,000
Molecular sutipo	Luminal A	45	1,000	,000	,000	,000
	Luminal B Her-2 neg	142	,000	,000	,000	,000
	Luminal B Her-2 posit	92	,000	1,000	,000	,000
	Her-2 enriched	75	,000	,000	1,000	,000
	Triple negativo	121	,000	,000	,000	1,000
Tipo celularidad infiltrado	Limfocitos	253	,000	,000	,000	
	Plasmaticas	61	1,000	,000	,000	
	Limfocitos+plasmaticas	135	,000	1,000	,000	
	No infiltrado	26	,000	,000	1,000	
Homogeneidad_cat3	Homogeneo	284	,000	,000		
	Heterogeneo	165	1,000	,000		
	No infiltrado	26	,000	1,000		
estromal_intraepitelial-cat3	estromal	395	,000	,000		
	intraepitelial con o sin estromal	54	1,000	,000		
	No infiltrado	26	,000	1,000		
ORHistologia	ductal	455	,000	,000		
	lobellar	14	1,000	,000		
	others	6	,000	1,000		
edatCat50	<=50	245	1,000			
	>50	230	,000			
Graudicotomico	grau 1 o 2	215	,000			
	grau3	260	1,000			
Ki67cat	<30	157	,000			
	=o>30	318	1,000			
TILS20	<=20	294	,000			
	>20	181	1,000			
TNM_cat3	Stage II	326	1,000			
	Stage III	149	,000			

Variables en la ecuación

	B	Error estándar	Wald	gl	Sig.	Exp(B)	95% C.I. para EXP(B)	
							Inferior	Superior
Paso 1 ^a								
TNM_cat3(1)	,329	,280	1,388	1	,239	1,390	,804	2,405
Graudiootómico(1)	,641	,326	3,870	1	,049	1,899	1,002	3,599
ORHistología			,000	2	1,000			
ORHistología(1)	-18,752	10243,430	,000	1	,999	,000	,000	.
ORHistología(2)	-19,105	15460,497	,000	1	,999	,000	,000	.
Molecular_sutipo			28,538	4	,000			
Molecular_sutipo(1)	-,803	1,107	,526	1	,468	,448	,051	3,925
Molecular_sutipo(2)	1,220	,392	9,698	1	,002	3,388	1,572	7,296
Molecular_sutipo(3)	1,919	,403	22,660	1	,000	6,815	3,093	15,020
Molecular_sutipo(4)	,711	,378	3,540	1	,060	2,037	,971	4,273
KI_67_cat(1)	,743	,369	4,061	1	,044	2,102	1,021	4,328
TILs20(1)	-,003	,376	,000	1	,993	,997	,477	2,082
Tipus_infiltrat			1,407	3	,704			
Tipus_infiltrat(1)	-,305	,405	,567	1	,451	,737	,333	1,631
Tipus_infiltrat(2)	,141	,304	,215	1	,643	1,152	,634	2,090
Tipus_infiltrat(3)	-,061	,852	,005	1	,942	,940	,177	4,994
estromal_intraepitelial_cat3			2,064	1	,151			
estromal_intraepitelial_cat3 (1)	,526	,366	2,064	1	,151	1,692	,826	3,465
Homogeneitat_TILs			10,631	2	,005			
Homogeneitat_TILs(1)	1,044	,348	9,024	1	,003	2,840	1,437	5,613
Homogeneitat_TILs(2)	-,508	,411	1,526	1	,217	,602	,269	1,347
Constante	-3,386	,488	48,099	1	,000	,034		

a. Variables especificadas en el paso 1: Homogeneitat_TILs.