

Supplementary Table 5. Multivariable Cox regression survival analyses of *CAV1* haplotypes in relation to breast cancer events, distant metastases, and death due to any cause for the entire follow-up period.

Breast cancer events								
Number of Haplotypes	Total	Events	Crude		Adjusted model 1		Adjusted model 2	
TTGTA	<i>n</i>	<i>n</i>	HR	(95% CI)	HR	(95% CI)	HR	(95% CI)
0	252	51	Ref.		Ref.		Ref.	
1	511	96	0.90	0.64 – 1.27	0.89	0.64 – 1.26	0.90	0.64 – 1.26
2	254	48	0.89	0.60 – 1.32	0.91	0.61 – 1.35	0.89	0.59 – 1.32
TCATA	<i>n</i>	<i>n</i>	HR		HR		HR	
0	710	135	Ref.		Ref.		Ref.	
1	283	57	1.08	0.79 – 1.47	1.09	0.80 – 1.49	1.13	0.83 – 1.56
2	24	3	0.65	0.21 – 2.05	0.72	0.23 – 2.26	0.73	0.23 – 2.30
CTACC	<i>n</i>	<i>n</i>	HR		HR		HR	
0	711	135	Ref.		Ref.		Ref.	
1	287	56	1.08	0.79 – 1.48	1.06	0.77 – 1.44	1.03	0.75 – 1.41
2	19	4	1.14	0.42 – 3.07	1.22	0.45 – 3.31	1.15	0.42 – 3.11
CTGTA	<i>n</i>	<i>n</i>	HR		HR		HR	
None (0)	858	172	Ref.		Ref.		Ref.	
Any (1+)	159	23	0.67	0.43 – 1.03	0.66	0.43 – 1.03	0.67	0.43 – 1.04
Distant metastasis								
Number of Haplotypes	Total	Events	Crude		Adjusted model 1		Adjusted model 2	
TTGTA	<i>n</i>	<i>n</i>	HR	(95% CI)	HR	(95% CI)	HR	(95% CI)
0	252	32	Ref.		Ref.		Ref.	
1	511	58	0.88	0.57 – 1.35	0.84	0.54 – 1.30	0.85	0.55 – 1.32
2	254	32	0.96	0.58 – 1.58	0.99	0.60 – 1.64	0.99	0.60 – 1.63
TCATA	<i>n</i>	<i>n</i>	HR		HR		HR	
0	710	83	Ref.		Ref.		Ref.	
1	283	38	1.16	0.79 – 1.70	1.15	0.78 – 1.70	1.19	0.80 – 1.76
2	24	1	0.36	0.05 – 2.59	0.47	0.06 – 3.37	0.45	0.06 – 3.28
CTACC	<i>n</i>	<i>n</i>	HR		HR		HR	
0	711	89	Ref.		Ref.		Ref.	
1	287	32	0.94	0.63 – 1.41	0.89	0.59 – 1.34	0.88	0.59 – 1.32
2	19	1	0.38	0.05 – 2.74	0.47	0.07 – 3.38	0.43	0.43 – 3.10
CTGTA	<i>n</i>	<i>n</i>	HR		HR		HR	
None (0)	858	106	Ref.		Ref.		Ref.	0.50 – 1.38
Any (1+)	159	16	0.77	0.46 – 1.31	0.78	0.46 – 1.33	0.79	0.46 – 1.33
Death								
Number of Haplotypes	Total	Events	Crude		Adjusted model 1		Adjusted model 2	
TTGTA	<i>n</i>	<i>n</i>	HR	(95% CI)	HR	(95% CI)	HR	(95% CI)
0	252	40	Ref.		Ref.		Ref.	
1	511	98	1.17	0.81 – 1.69	1.11	0.77 – 1.61	1.13	0.79 – 1.64
2	254	50	1.21	0.79 – 1.83	1.22	0.80 – 1.86	1.21	0.80 – 1.85
TCATA	<i>n</i>	<i>n</i>	HR		HR		HR	
0	710	126	Ref.		Ref.		Ref.	
1	283	59	1.17	0.86 – 1.61	1.18	0.86 – 1.61	1.18	0.86 – 1.61
2	24	3	0.78	0.25 – 2.45	0.85	0.27 – 2.68	0.91	0.29 – 2.87
CTACC	<i>n</i>	<i>n</i>	HR		HR		HR	
0	711	135	Ref.		Ref.		Ref.	
1	287	52	1.01	0.73 – 1.39	0.99	0.71 – 1.36	0.99	0.71 – 1.37
2	19	1	0.25	0.04 – 1.81	0.29	0.04 – 2.05	0.29	0.04 – 2.09
CTGTA	<i>n</i>	<i>n</i>	HR		HR		HR	
None (0)	858	163	Ref.		Ref.		Ref.	
Any (1+)	159	25	0.76	0.50 – 1.17	0.71	0.46 – 1.09	0.70	0.46 – 1.08

Adjusted model 1: Age at inclusion, tumor size, nodal status, grade III, and ER status. Missing data for four patients for at least one variable.

Adjusted model 2: Model 1+ chemotherapy, radiotherapy, trastuzumab, tamoxifen, and aromatase inhibitors. Missing data for four patients for at least one variable.

Caveolin-1 genotypes as predictor for locoregional recurrence and contralateral disease in breast cancer

Breast Cancer Research and Treatment

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