## **Supplementary Table 1:** COX-2 expression in Breast TMA comparing COX2 clones CX229 and CX294

TMA Staining Results	COX-2 Clones	COX-2 Positive Breast TMA (n=56)
Dual Positive COX-2	CX229+/CX294+	53 (95%)
Positive for CX229 only	CX229+/CX294-	0
Positive for CX294 only	CX229-/ CX294+	0
Dual negative COX-2	CX229-/ CX294 -	3 (5%)

## **Supplementary Table 2:** COX-2 expression in Breast and Colon TMA comparing clones SP21 and CX229

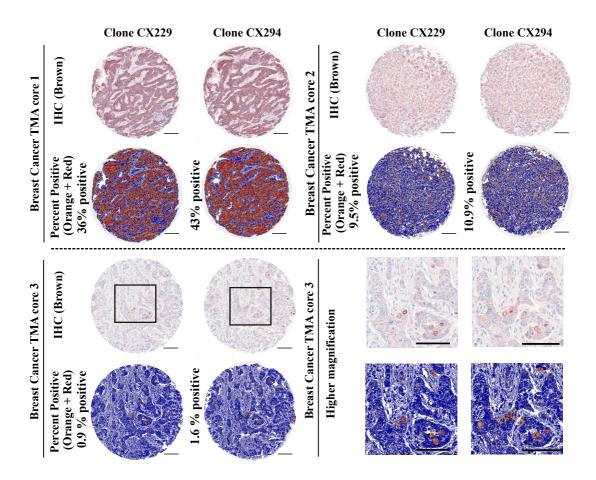
TMA Staining Results	COX-2 Clones	COX-2 Positive Breast TMA (n=52)	COX-2 Positive Colon TMA (n=53)
Non-nitrosylated COX-2 only	CX229+/SP21 <b>-</b>	18 (35%)	23 (43%)
Dual positive COX-2	CX229+/SP21+	29 (55%)	21 (40%)
S-nitrosylated COX-2 only	CX229 <b>-</b> /SP21+	1 (2%)	0 (0%)
Dual negative COX-2	CX229-/SP21-	4 (8%)	9 (17%)

Supplementary Table 3: Clinical characteristics of NHS 1 and YWBC cohorts

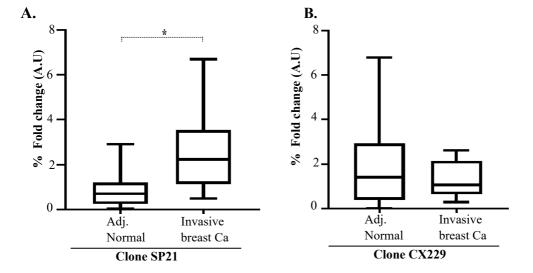
Clinical	<b>Nurses' Health</b>	Young Women Breast	P value
characteristics	Study 1	Cancer	(chi square test)
Overall N	1770 cases	233 cases	
Age at dx, mean	57.8	38.24	
Stage			
1	885 (50%)	67 (28.7%)	
2	600 (33.9%)	97 (41.6%)	< 0.0001
3	250 (14.1%)	52 (22.3%)	
4	35 (2%)	15 (6.52%)	
Grade			
1	303 (17.1%)	22 (9.4%)	
2	935 (52.8%)	86 (36.9%)	< 0.0001
3	416 (23.5%)	109 (46.8%)	
Missing	116 (6.6%)	16 (6.9%)	
Tumor size (cm)			
≤1	439 (24.6%)	30 (12.9%)	
1.1-2.0	682 (38.3%)	73 (31.4%)	< 0.0001
2.1-4.0	481 (27.1%)	65 (27.9%)	
>4.0	164 (9.8%)	53 (22.7%)	
Missing	4 (0.2%)	12 (5.1%)	
ER status			
Positive	1361 (76.9%)	142 (60.9%)	
Negative	404 (22.8%)	87 (37.3%)	< 0.0001
Missing	5 (0.3%)	4 (1.8%)	
PR status	, ,	, ,	
Positive	1137 (64.2%)	135 (57.9%)	
Negative	625 (35.3%)	93 (39.9%)	0.1156
Missing	8 (0.5%)	5 (2.1%)	
HER2 status			
Positive	195 (11.0%)	67 (28.7%)	< 0.0001
Negative	1546 (87.4%)	147 (63%)	
Missing	29 (1.6%)	19 (8.3%)	
Hormone therapy	, ,	,	
Yes	837 (47.4%)	79 (33.9%)	
No	433 (24.6%)	67 (28.8%)	0.0047
Missing	496 (28%)	87 (37.3%)	
Radiation	, ,	, ,	
Yes	545 (30.7%)	90 (38.6%)	
No	735 (41.5%)	55 (43.5%)	< 0.0001
Missing	490 (27.8%)	88 (37.8%)	
Chemotherapy	` '	` '	
Yes	492 (27.8%)	137 (58.9%)	
No	772 (43.6%)	19 (8.1%)	< 0.0001
Missing	506 (28.6%)	77 (33%)	

## Supplementary Table 4: The methodological details for IHC and IF staining

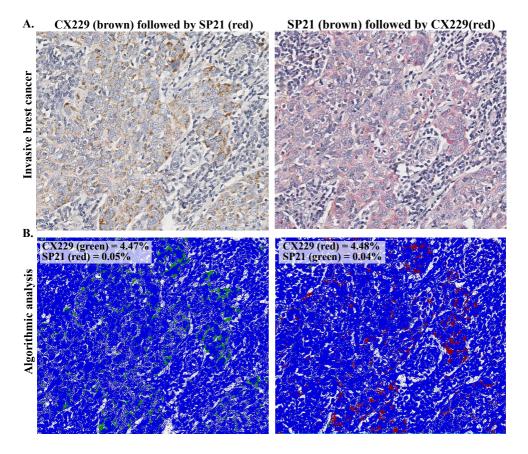
Stain/single	Antigen retrieval	Peroxidase and Protein Blocks	Primary Antibody	Secondary Antibody	Chromogen
or dual			•	· ·	
IHC –	125°C under	Peroxidase block	SP21 clone	Envision +	DAB (Dako
SP21 stain	pressure for	(1ml 3% H2O2 +	(Thermoscienti	HRP	# K3468)
	5 mins with	9 ml methanol) for	fic # RM-	antibody	for 10 mins
	TRS (Dako	10 mins	9121) 1:200	(Dako #	
	#S1699)	Protein block (Biocare	for 60 mins	K4003	
		# BS966L) for 10 mins		rabbit) for	
				30 mins	
IHC-	125°C under	Peroxidase block	CX229 clone	Envision +	DAB (Dako
CX229	pressure for	(1ml 3% H2O2 +	(Cayman, #	HRP	# K3468)
stain	5 mins with	9 ml methanol) for	160112)	antibody	for 10 mins
	TRS (Dako	10 mins	1:250	(Dako #	
	#S1699)	Protein block (Biocare	for 60 mins	K4001	
		# BS966L) for 10 mins		mouse) for	
				30 mins	
IHC-	125°C under	Peroxidase block	CX294	Envision +	DAB (Dako
CX294	pressure for	(1ml 3% H2O2 +	clone	HRP	# K3468)
stain	5 mins with	9 ml methanol) for	(Agilent#	antibody	for 10 mins
	TRS (Dako	10 mins	M3617)	(Dako #	
	#S1699)	Protein block (Biocare	1:50	K4001	
		# BS966L) for 10 mins	for 60 mins	mouse) for	
				30 mins	
IHC- Dual	125℃ under	1st antibody stain:	CX229 clone	Envision +	DAB (Dako
SP21 and	pressure for	Peroxidase	(Cayman,	HRP antibody	# K3468)
CX229	5 mins with	(1ml 3% H2O2 + 9 ml	#160112)	(Dako #	for 10 mins
stain	TRS (Dako	methanol) for 10 mins	1:250for 60	K4001mouse)	
	#S1699)		mins	for 30 mins	
		Protein block (Biocare			
		# BS966L) for 10 mins			
		2nd antibody			
		stain: Protein			
		block (Biocare			
		# BS966L) for			
		10 mins			



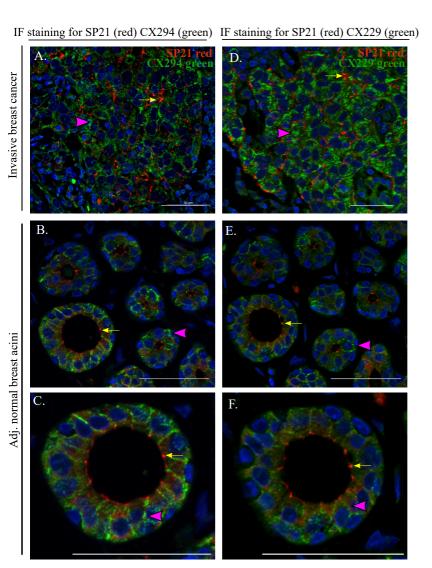
Supplementary Figure 1: TMA cores from breast cancer cases stained for CX229 and CX294 have similar staining intensity, ranging from high (core #1), medium (core #2) or low (core #3). Higher magnification of core #3 (black box) shows that CX229 and CX294 have similar specificity in staining the same cells on sequential sections. Algorithmic analysis for each TMA core shows COX2 positive signal (orange and red) compared to negatively stained tissue (blue). Scale bar for all images is 100μm.



**Supplementary Figure 2.** SP21 and CX229 clones show inverted staining patterns between adjacent normal and invasive cancer in Kaiser Pacific North West (KPNW) cases. A. SP21 staining intensity was low in adjacent normal breast tissue and increased in invasive breast cancer (n = 10, P value:  $\leq 0.05$ ). B. CX229 (n = 10) shows highest COX2 expression in normal adjacent breast epithelium with a trend towards decreased COX2 expression in invasive breast cancer.

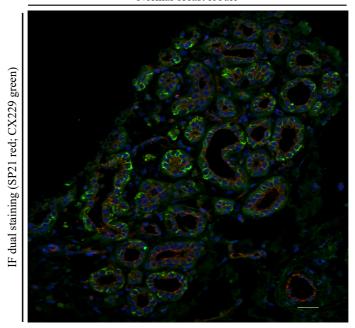


**Supplementary Figure 3.** A. Dual IHC stained images of SP21 and CX229 show similar staining patterns when the order of antibodies and chromogens were switched during staining process. B. Algorithmic analysis of the images in part A of this figure (upper panel) revealed 4.47% CX229 (green) and 0.05% SP21 (red) expression (left, lower panel). With reversed antibody order, there was 4.48% CX229 (red) and 0.04% SP21 (green) expression (right, lower panel).



**Supplementary Figure 4.** IF staining of serial sections of an invasive cancer case and adjacent normal breast acini show distinct sub-cellular localization of SP21, CX294 and CX229. The left column shows dual stains of SP21 and CX294 with minimal co-localization (SP21 = red, CX294 = green, co-localization = yellow). The right column shows dual stains of SP21 and CX229 with minimal co-localization (SP21 = red, CX294 = green, co-localization = yellow). Pink arrow heads show intense localized staining for CX294/CX229. Yellow arrows show intense localized staining for SP21. Scale bar for all images is 50µm.

## Normal breast lobule



Supplementary Figure 5. IF staining of a true normal lobule with breast acini show distinct subcellular localization of SP21 and CX229. The dual stains of SP21 and CX229 have minimal colocalization (SP21 = red, CX294 = green, co-localization = yellow) but have heterogeneous expression in each acini. Scale bar for all images is  $30\mu m$ .