CD74 interacts with CD44 and enhances tumorigenesis and metastasis via RHOA-mediated cofilin phosphorylation in human breast cancer cells

Supplementary Materials



Supplementary Figure S1: Detection of CD74 expression level. (A) CD74 was knocked down using CD74 shRNA contained vector in MDA-MB-231 and T47D cells, CD74 level was assayed by western blot. (B) CD74 expression level in MDA-MB-231 pLT-shN. C., MDA-MB-231 pLT-shCD74 #1 and MDA-MB-231 pLT-shCD74 #2 cell lines were detected by western blot.



Supplementary Figure S2: CD74 binds to CD44 in human breast cancer cells. (A) For immunofluorescence, CD74 was stained with AlexaFluor 568 dye (red), CD44 was stained with AlexaFluor 488 dye (green), yellow staining showed that CD74 colocalized with CD44. (B) MDA-MB-231 and T47D cells were transfected with pcDNA3.1 or pcDNA3.1-Flag-CD74, cell lysate was immunoprecipitated with anti-Flag antibody, CD74 and CD44 were detected by western blot. (C) For immunoprecipitation, MDA-MB-231 cell lysate was seperated into two group and incubated with IgG antibody and CD44 antibody at 4°C for 1 h, respectively. Then, cell lysate incubated with protein G at 4°C overnight, CD74 and CD44 protein were detected by western blot.



Supplementary Figure S3: Knockdown of CD74 or/and CD44 decreased cell protrusion formation. (A) Knockdown of CD74, CD44 or double knockdown of CD74 and CD44 were performed in MDA-MB-231 cells, the cells were resuspended and reseeded onto adhesion microscope slides after 24 h of transfection, F-actin was stained with TRITC-conjugated phalloidin (red), nuclei was visualized with DAPI (blue). Cell morphology was detected by confocal laser scanning microscopy. (B) T47D cells were treated similar to MDA-MB-231 cells, and representative images are shown as indicated. Arrows stand for the typical protrusions.



Supplementary Figure S4: Inhibition of CDC42 or RAC minimally down-regulated CFL1 phosphorylation. (A) MCF-7 cells were co-transfected with CD74 and CDC42 N17 (Dominant Negative), western blot analysis showed the phosphorylation of CFL1. (B) MCF-7 cells were co-transfected with CD74 and RAC N17 (Dominant Negative), CFL1 phosphorylation was assessed by western blot assay.



Supplementary Figure S5: RHOA is required for CD74 and p-CFL1-dependent cell migration. (A) MDA-MB-231 cells were co-transfected with CD74 shRNA contained plasmid and RHOA L63 (Constitutively Active) vector, RHOA-dependented CFL1 phosphorylation was analyed by western blotting. (B) MDA-MB-231 cells were subjected to wound-healing scratch assay after tranfecting as described above, cell were allowed to recover for 72 h. Representative images of cell recover are shown on the right. (C) Differences between groups were analyzed by *t*-test. The error bars represent the SD, *P < 0.05; **P < 0.01; ***P < 0.001. (D) MCF-7 cells were co-transfected with pcDNA3.1-CD74 and RHOA N19 (Dominant Negative) plasmids, RHOA regulated CFL1 phosphorylation was detected by immunoblot assay. (E) Wound-healing scratch assay was conducted after MCF-7 cotransfected with CD74 and RHOA N19. Cells were allowed to recover for 48 h. Representative images of cell recover are shown. (F) Differences between pcDNA3.1 plasmid transfected group were analyzed by *t*-test. The error bars represent the SD, *p-value < 0.001. (G) MCF-7 cells were co-transfected with CD74 and RHOA N19 plasmid transfected group were analyzed by *t*-test. The error bars represent the SD, *p-value < 0.05 and ***p-value < 0.001. (G) MCF-7 cells were co-transfected with CD74 and RHOA N19 for 24 h, then, re-suspended cells were re-seeded onto adhesion microscope slides, F-actin was stained with TRITC-conjugated phalloidin (red), nuclei was visualized with DAPI (blue).

Patients characteristics	No. of patients (%)
Age (years)	
\leq 45	91 (48.1)
> 45	98 (51.9)
Clinical stages	
Stage I	9 (4.8)
Stage II	109 (57.7)
Stage III	50 (26.5)
Stage IV	21 (11.1)
Lymph node status	
N0	49 (26.0)
N1/N2/N3	140 (74.0)
Pathological grades	
Well	11 (5.8)
Moderate	106 (56.1)
Poor	72 (38.1)
Survival status	
Alive	144 (76.2)
Death	45 (23.8)

Supplementary Table S1: 189 cases of BIDC patients' features

Supplementary Table S2: Comparison of CD74 protein expression in BIDC and non-cancerous control breast tissues

	CI			
Features (n)	Low expression (%)	High expression (%)	<i>r</i> -value	
Cancer $(n = 189)$	46 (24.3)	143 (75.7)		
NCBT(n = 40)	22 (55.0)	18 (45.0)	0.000*	

*Chi-square test, statistically significant difference (P < 0.05). NCBT: non-cancerous control breast tissues.

Variables	D	SE	Wald	Sia	Even (D)	95.0% CI for Exp (B)	
variables	D	5.E .	walu	51g.	стр (р)	Lower	Upper
Clinical stages	1.863	.529	12.418	.000*	6.446	2.286	18.172
Pathologic grades	.067	.420	.025	.873	1.069	.469	2.438
Survival status	.255	.557	.210	.647	1.291	.433	3.848
ER	117	.880	.018	.894	.889	.158	4.989
PR	.372	.875	.181	.670	1.451	.261	8.064
CerbB-2	.175	.398	.193	.660	1.191	.546	2.596
Age(years)	.118	.360	.108	.742	1.126	.556	2.278
CD74	1.205	.412	8.573	.003*	3.337	1.490	7.478

Supplementary Table S3: Multivariate logistic regression analysis of lymph node metastasis factors in BIDC patients (n = 189)

95% CI: 95% confidence interval.*P < 0.05.

Supplementary	Table S4:	Analysis	of the	association	between	expression	of CD74	protein	and
clinicopathologi	cal feature	es of BIDC	C (n = 1	189)					

	CI	Develope			
reatures	Low expression (%)	High expression (%)	<i>P</i> -value		
Age (years)					
\leq 45 (<i>n</i> = 91)	21 (23.1)	70 (76.9)			
> 45 (<i>n</i> = 98)	25 (25.5)	73 (74.5)	0.697		
Pathological grades					
Well/Moderate ($n = 117$)	30 (25.6)	87 (74.4)			
Poor $(n = 72)$	16 (22.2)	56 (77.8)	0.595		
Clinical stages					
Stage I–II ($n = 118$)	23 (19.5)	95 (80.5)			
Stage III–IV ($n = 71$)	23 (32.4)	48 (67.6)	0.045*		
Lymph node status					
LNM (<i>n</i> = 140)	28 (20.0)	112 (80.0)			
No LNM (<i>n</i> = 49)	18 (36.7)	31 (63.3)	0.019*		
ER					
Negative $(n = 60)$	17 (28.3)	43 (71.7)			
Positive $(n = 129)$	29 (22.5)	100 (77.5)	0.383		
PR					
Negative $(n = 65)$	19 (29.2)	46 (70.8)			
Positive $(n = 124)$	27 (21.8)	97 (78.2)	0.256		
HER-2					
Negative $(n = 124)$	31 (25.0)	93 (75.0)			
Positive $(n = 65)$	15 (23.1)	50 (76.9)	0.770		
Survival status					
Alive $(n = 144)$	33 (22.9)	111 (77.1)			
Death $(n = 45)$	13 (28.9)	32 (71.1)	0.415		

*Chi-square test, statistically significant difference (p < 0.05). LNM: lymph node metastasis. *P* is used to test the association between expression of CD74 and the clinicopathological features of BIDC.