

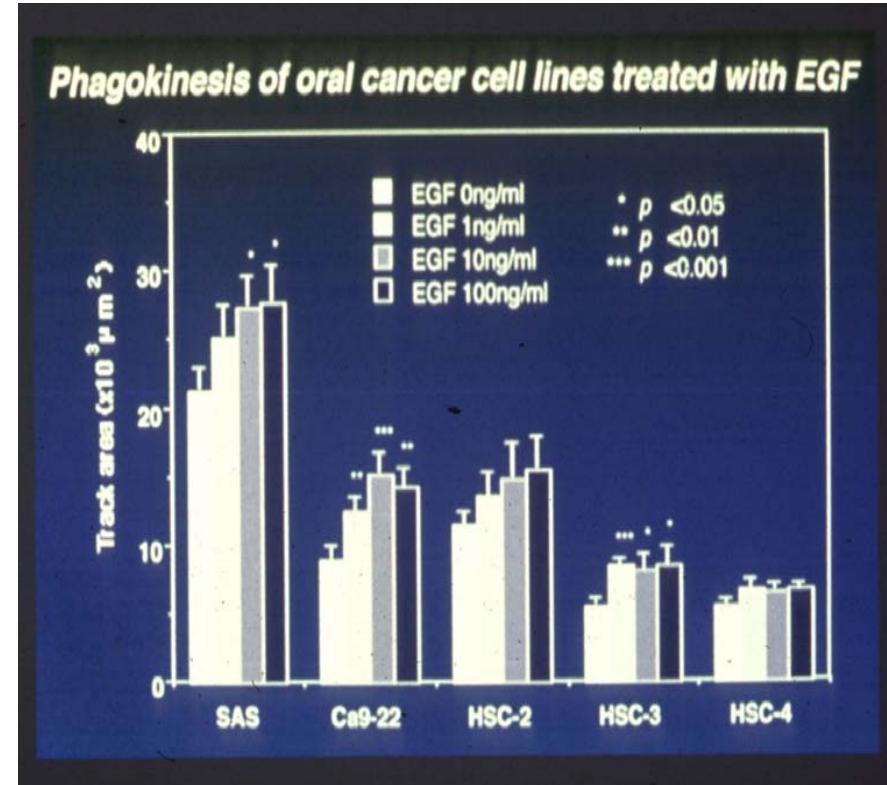
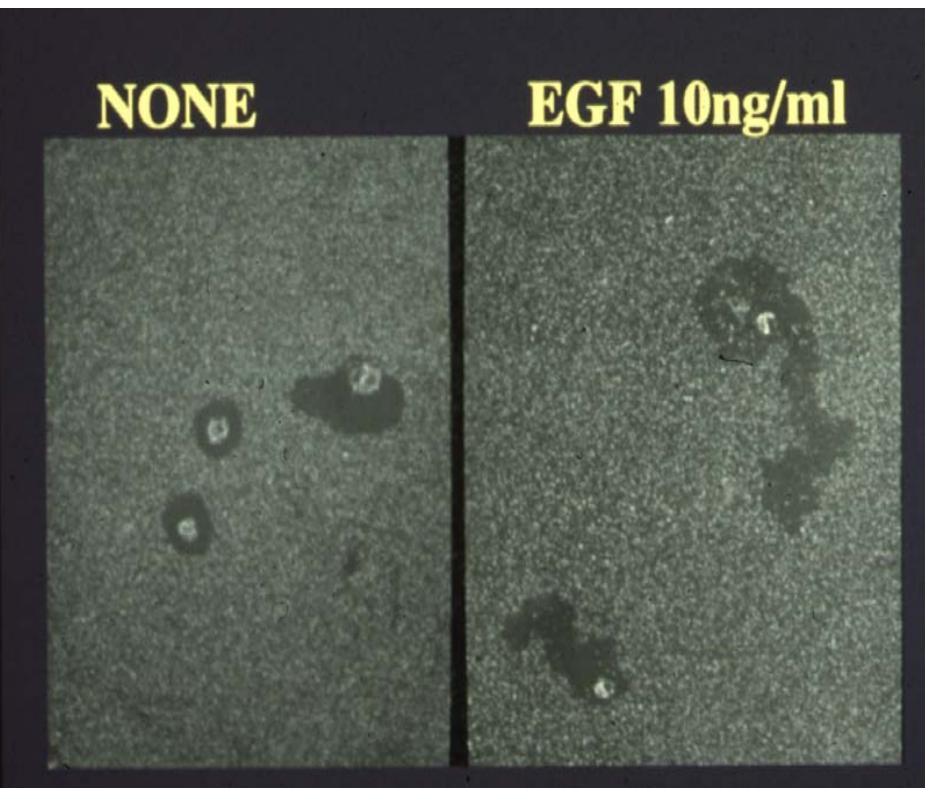


## EGF受容体・シグナル伝達を分子標的とする口腔がん治療の基礎的検討

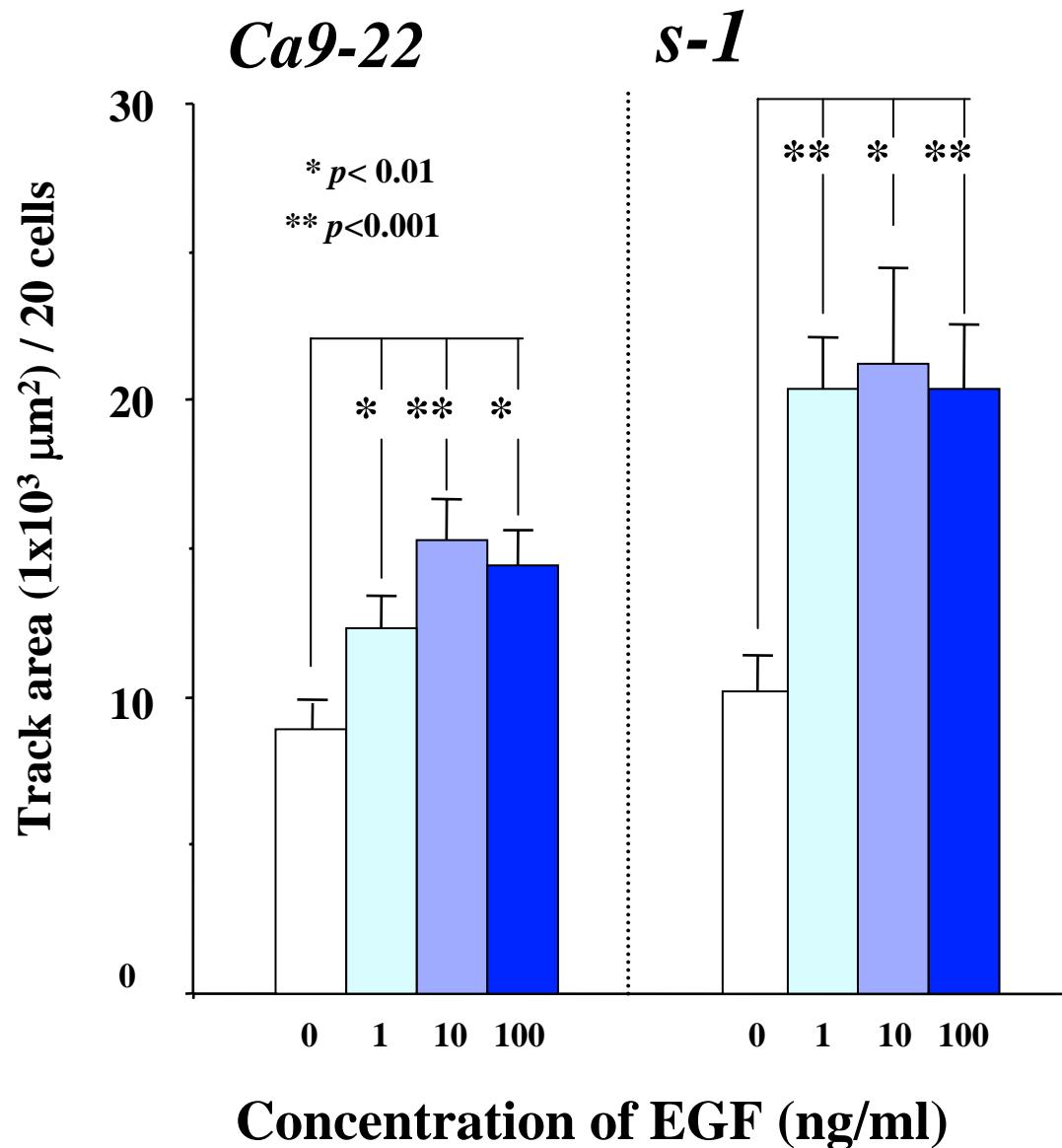
メタデータ	言語: Japanese 出版者: 公開日: 2008-03-12 キーワード (Ja): キーワード (En): SCC, HUMAN, EGF, INVASION, EGF RECEPTOR, PKC 作成者: 柴田, 敏之, 浜田, 淳一, 土井田, 誠, 山下, 知巳, 牧田, 浩樹 メールアドレス: 所属:
URL	<a href="http://hdl.handle.net/20.500.12099/2816">http://hdl.handle.net/20.500.12099/2816</a>

# ヒト口腔がん細胞の浸潤シグナルの伝達

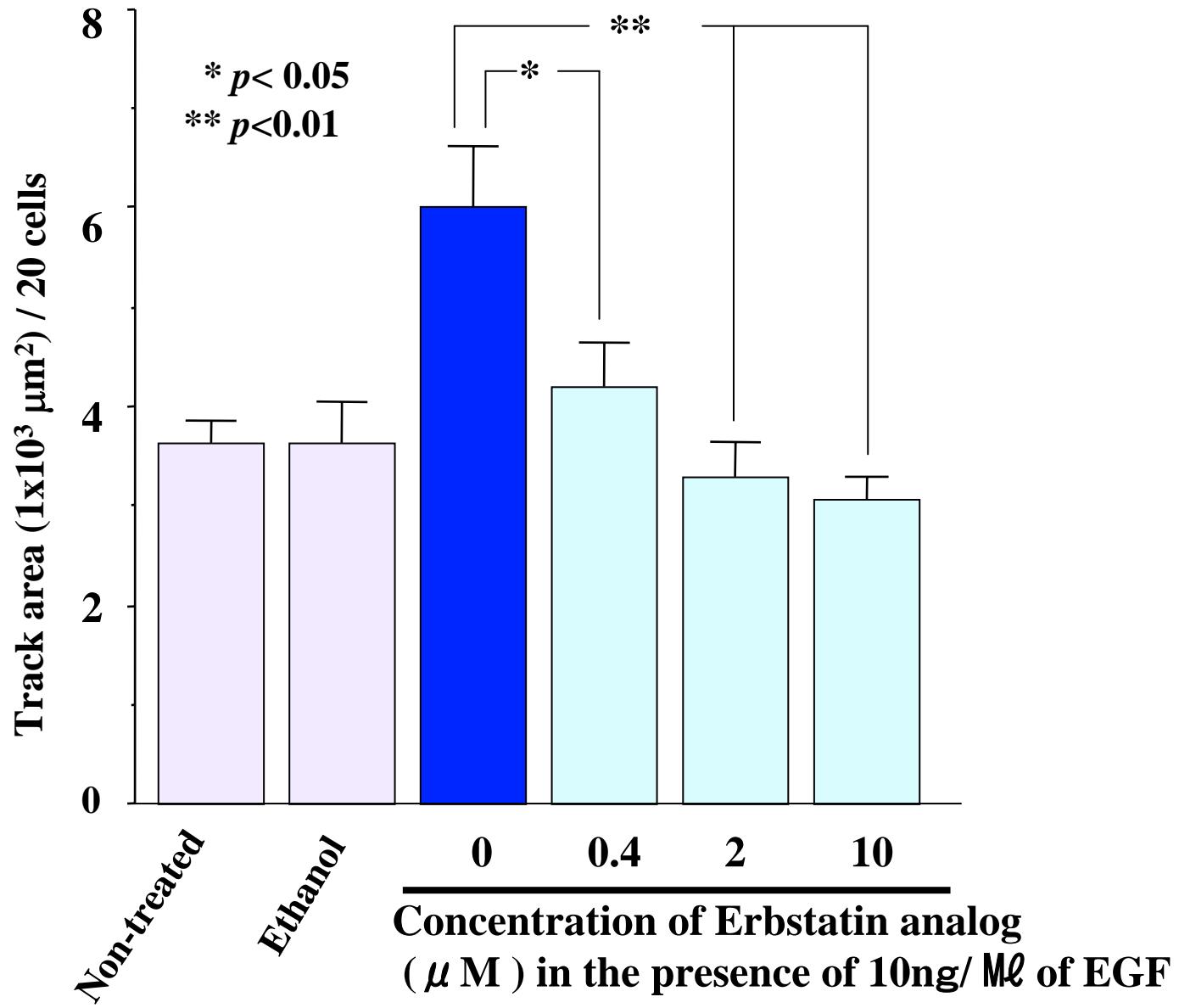
EGF刺激による運動能の促進(金コロイド法による評価)



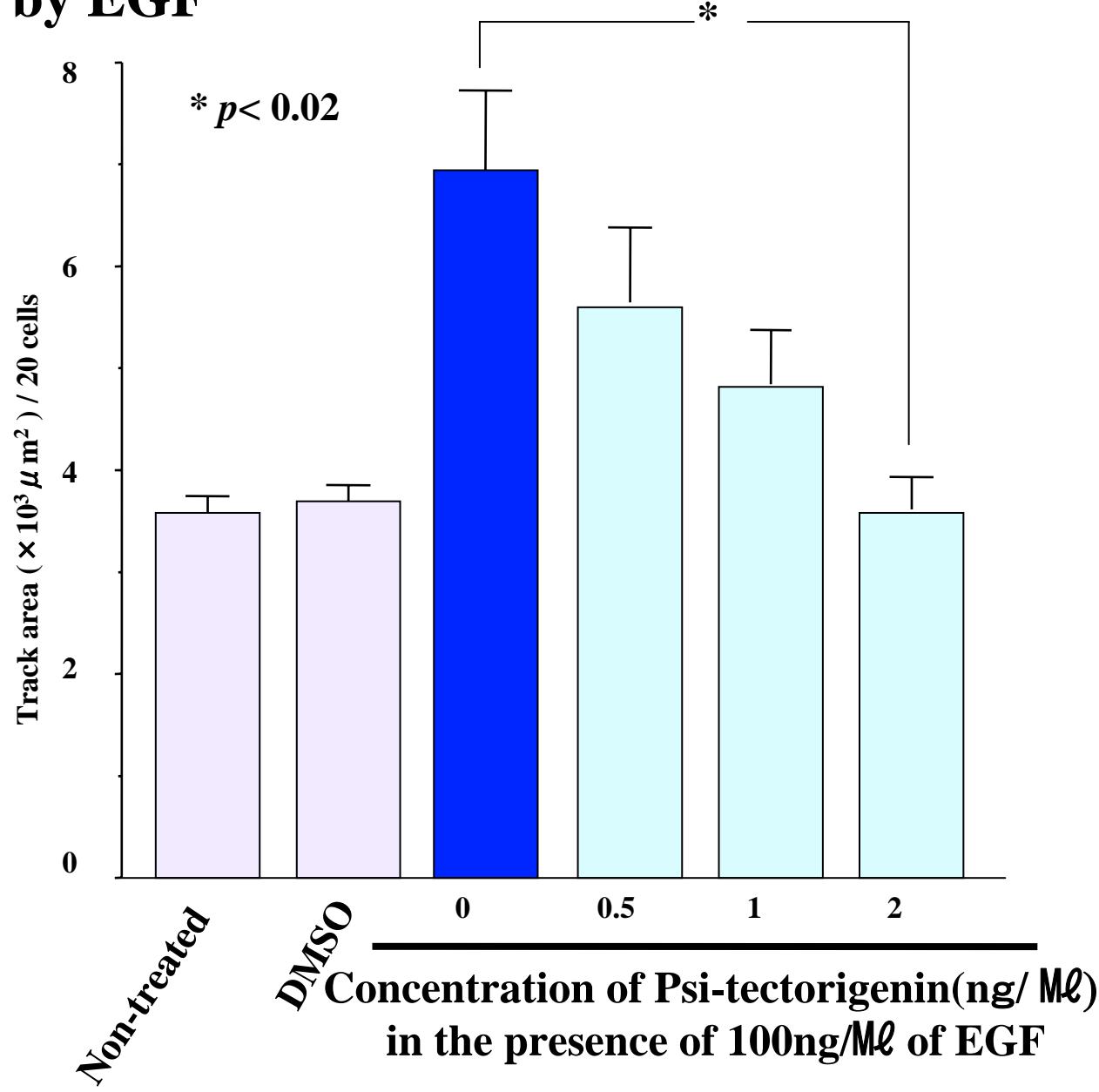
# Enhanced effect of EGF stimulation on the motility of Ca9-22 and s-1 cells



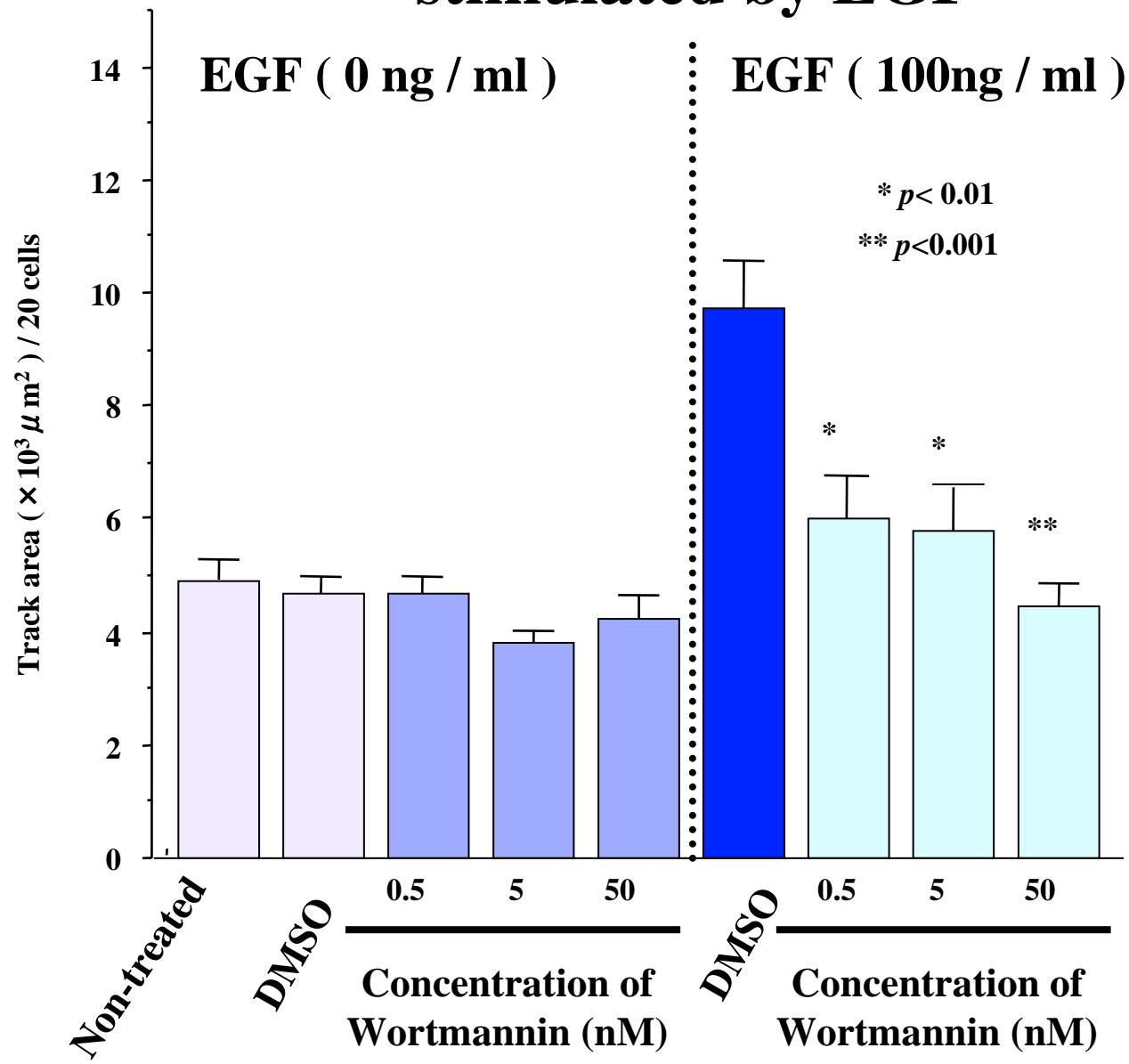
# Effect of Erbstatin analog on the motility of s-1 cells stimulated by EGF



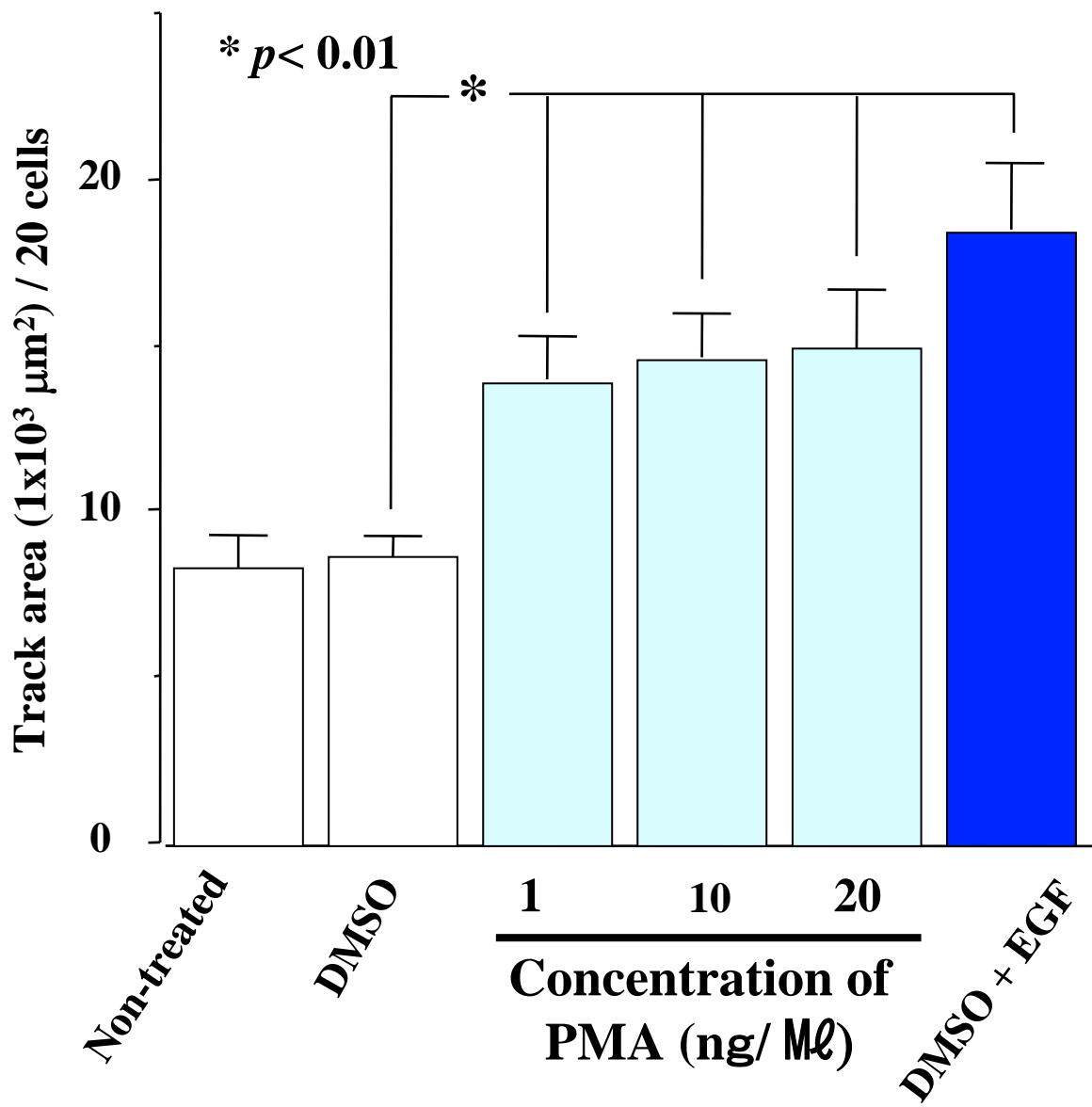
# Effect of Psi-tectorigenin on the motility of s-1 cells stimulated by EGF



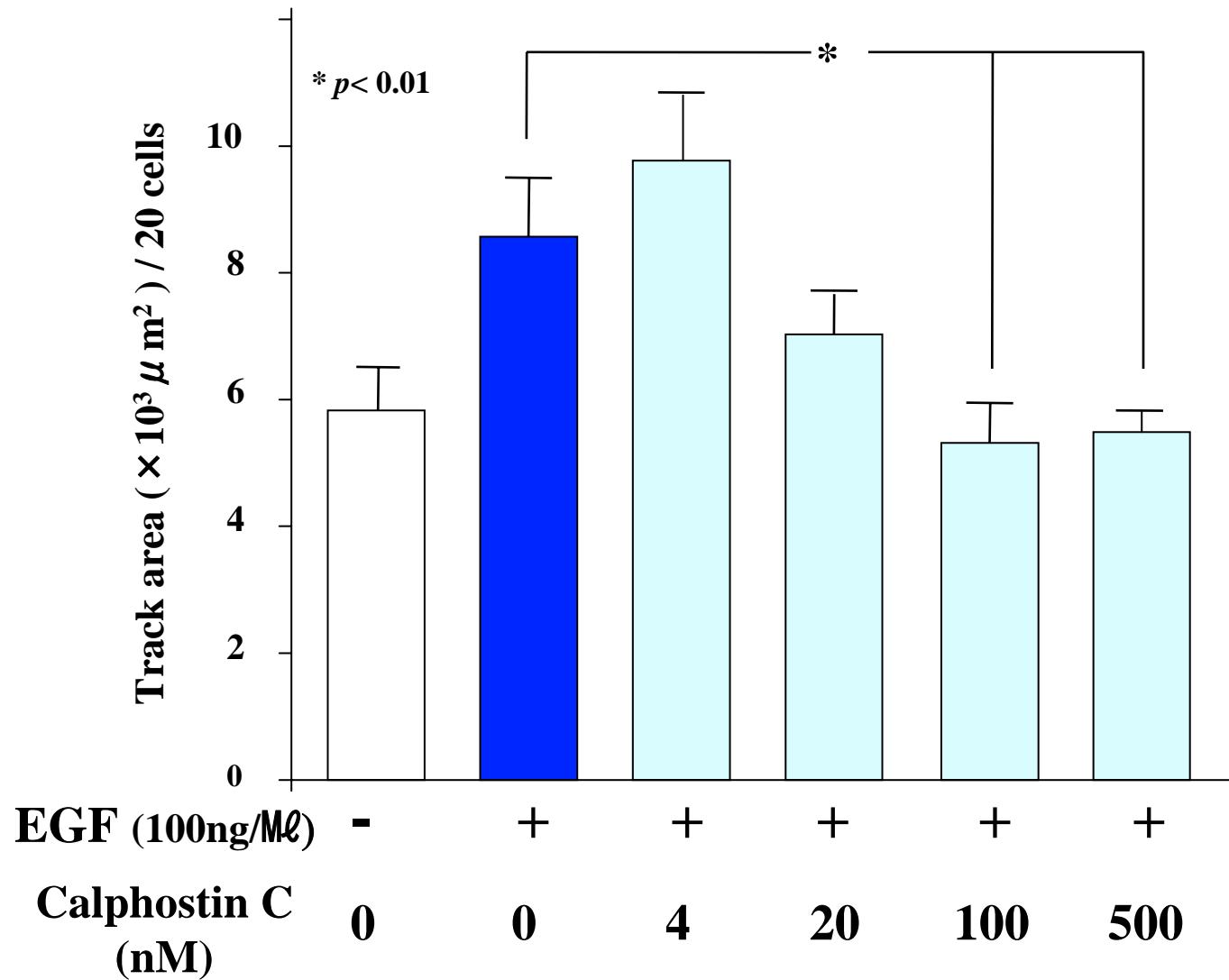
# Effect of Wortmannin on the motility of s-1 cells stimulated by EGF



# Effect of PKC activator on the motility of s-1 cells

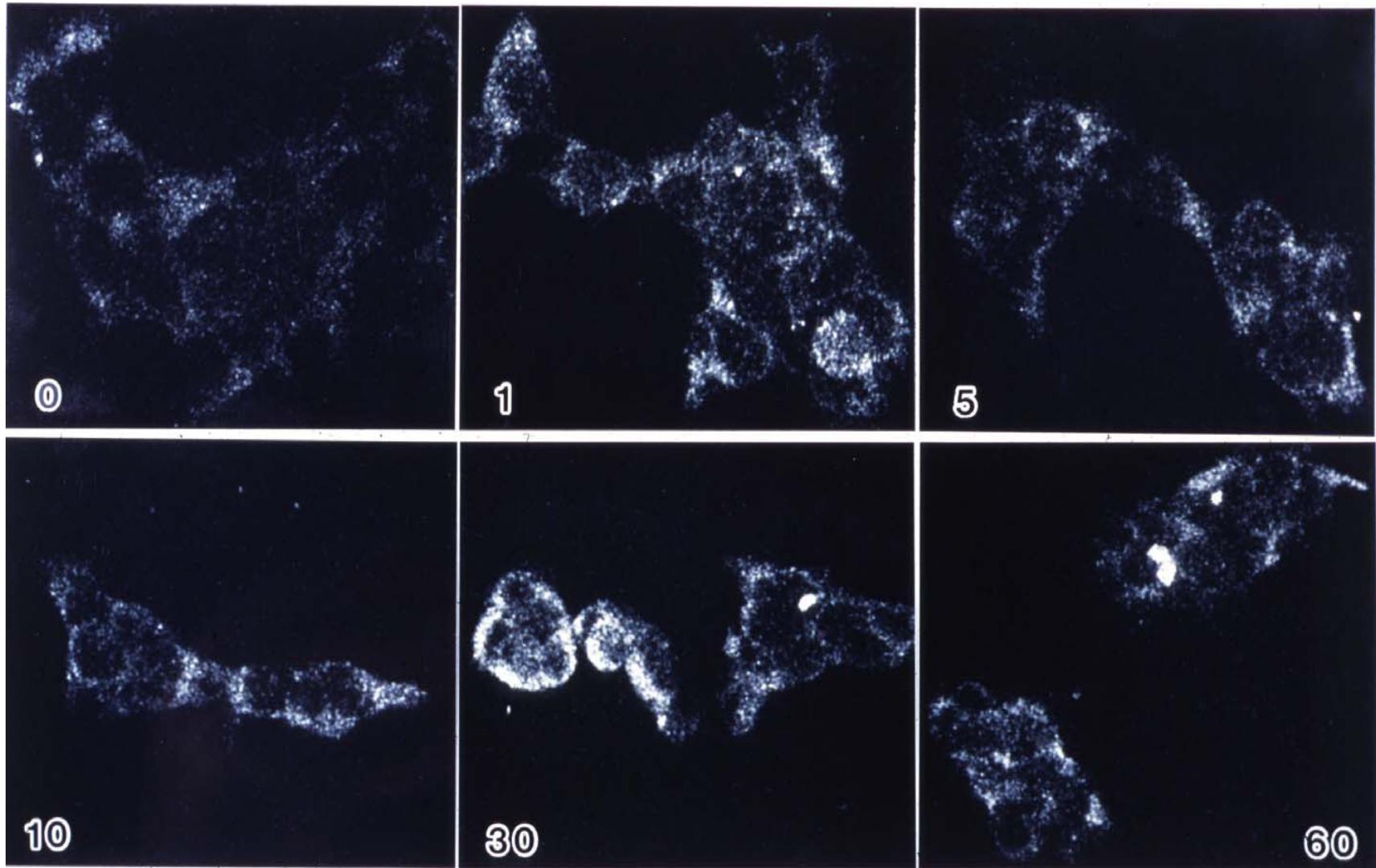


# Effect of PKC inhibitor on the motility of s-1 cells stimulated by EGF

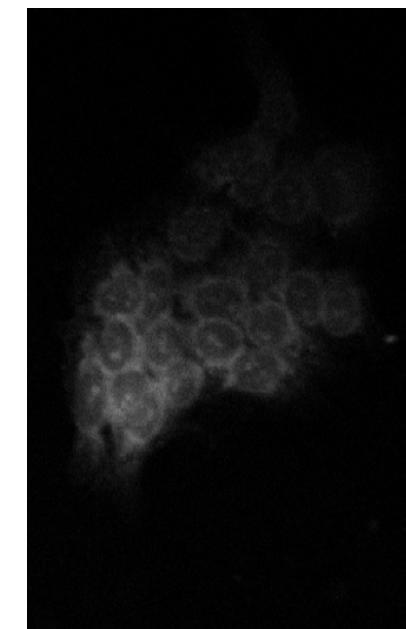
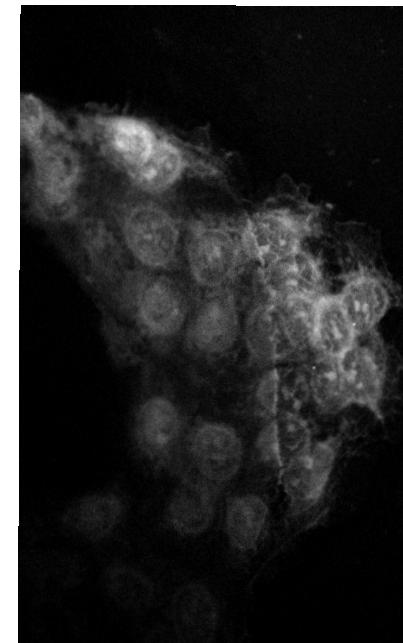
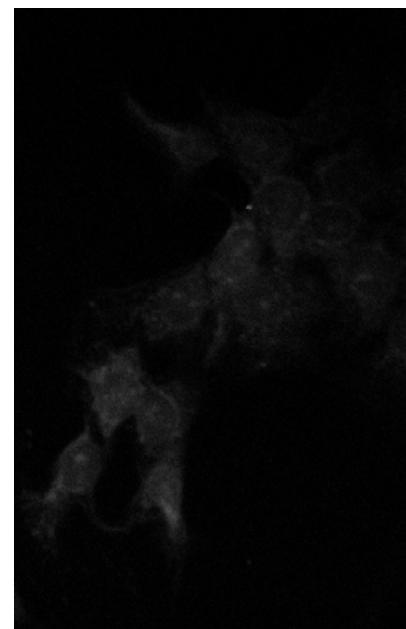


# Translocation of PKC in EGF stimulated s-1 cells

pan PKC MoAb staining



**nPKC- $\delta$**



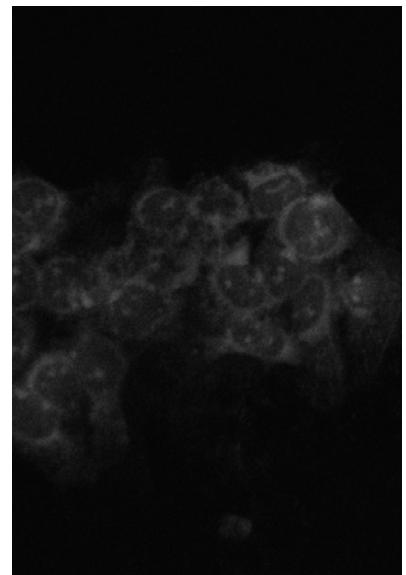
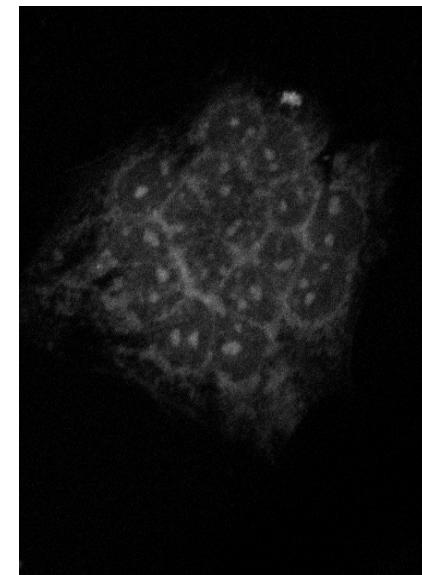
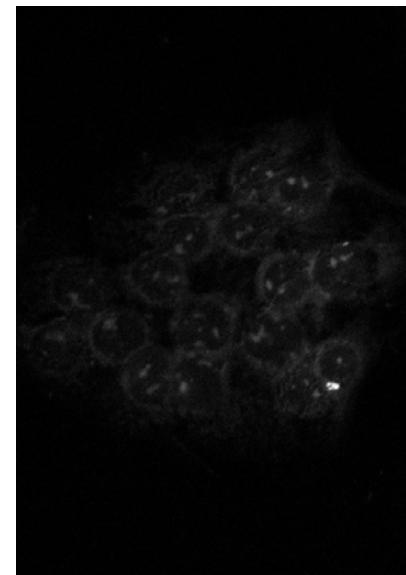
**EGF stimulation**

**1**

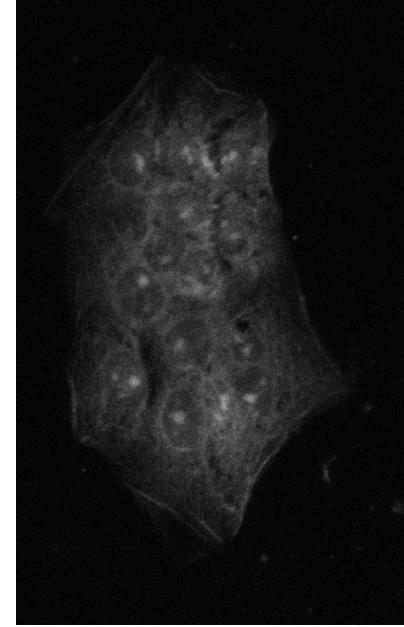
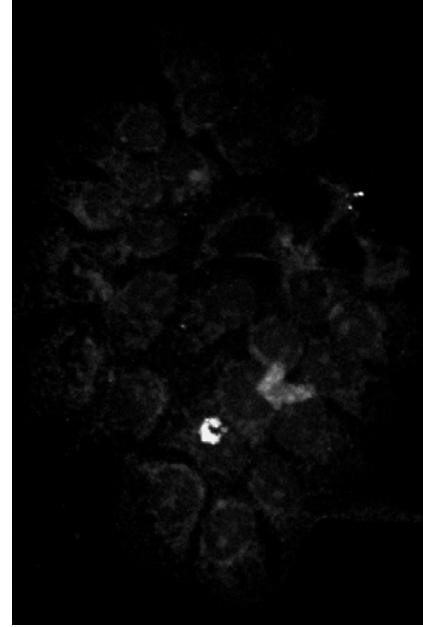
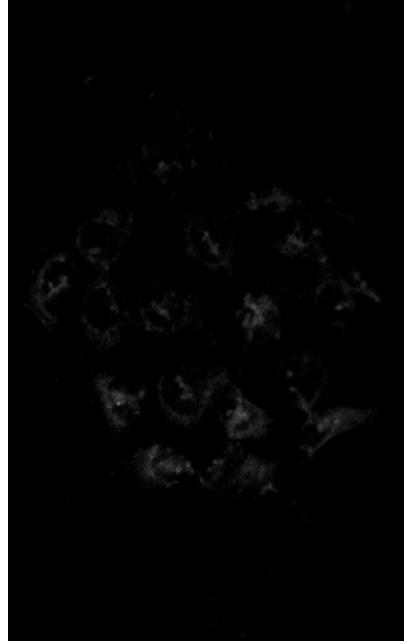
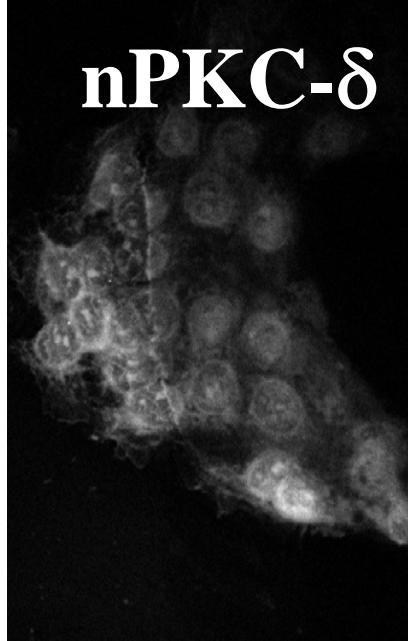
**10**

**60 (min)**

**aPKC- $\zeta$**



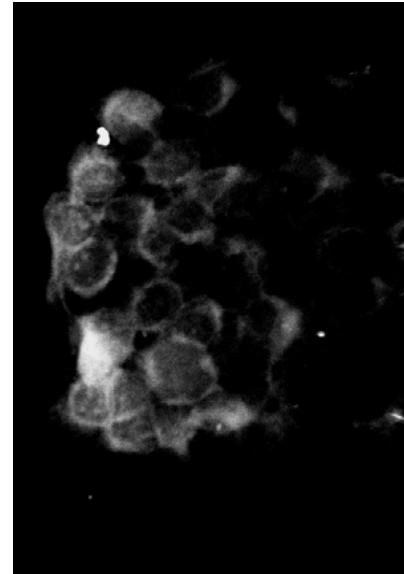
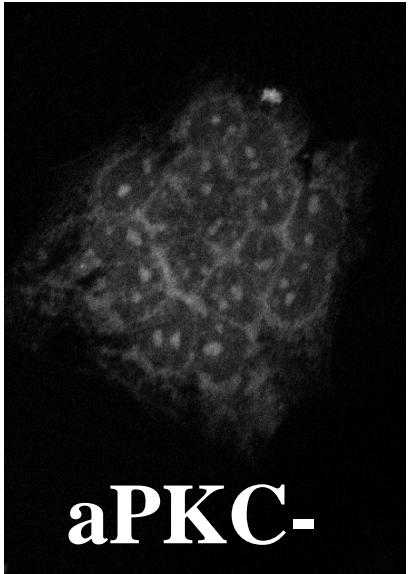
**nPKC- $\delta$**



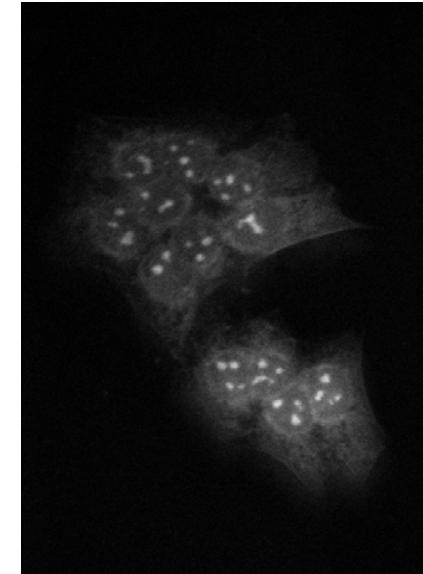
**EGF**

**+**

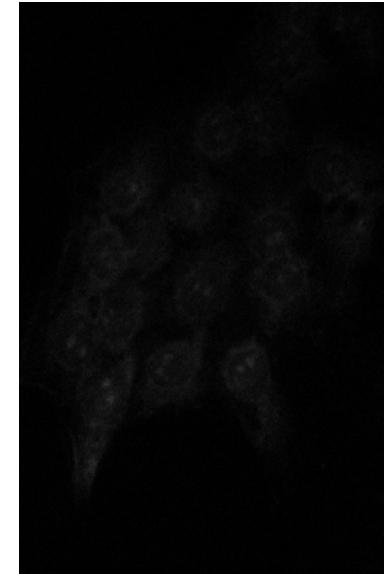
**Erbstatin**



**Psi-tectorigenin**

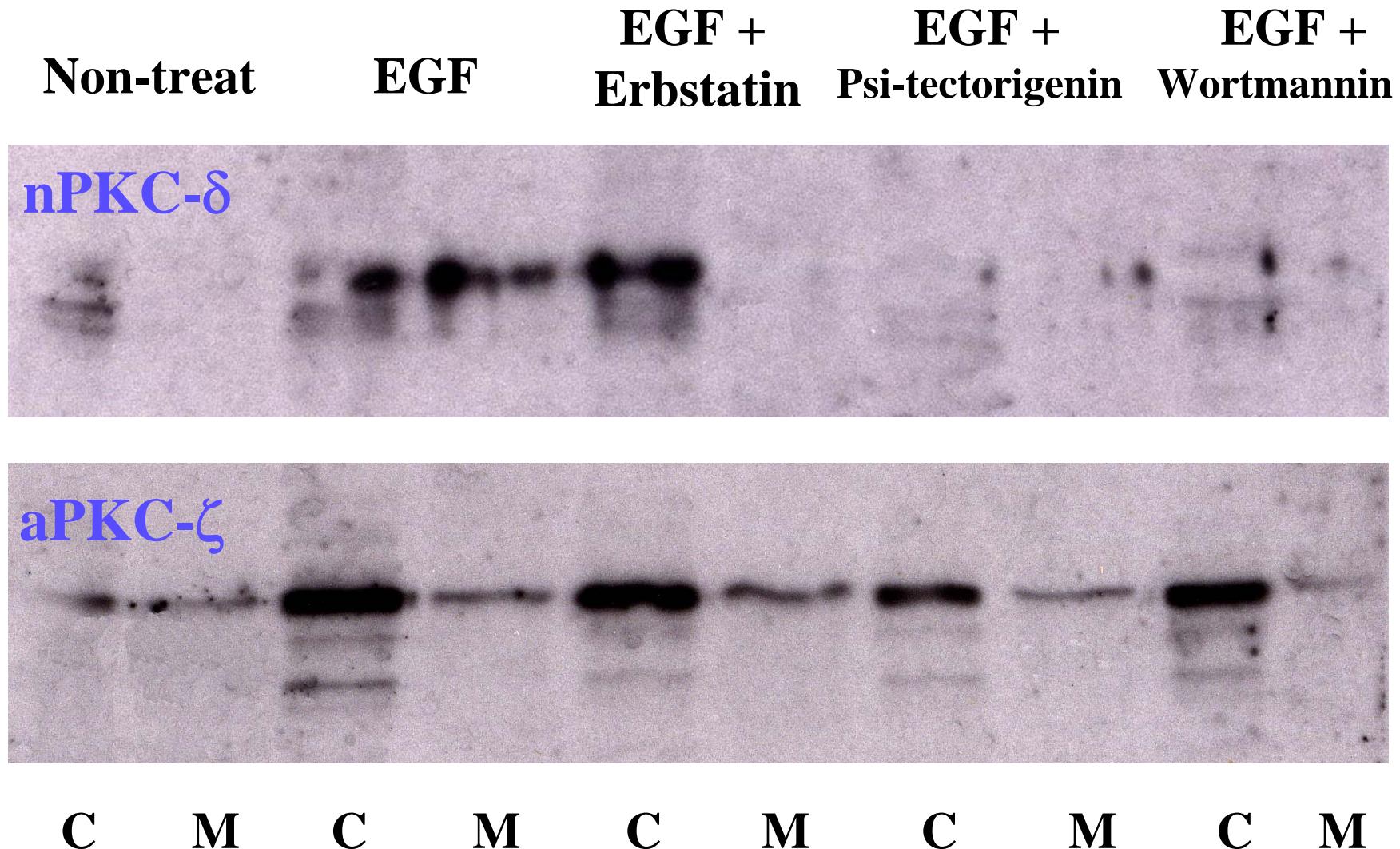


**Wortmannin**

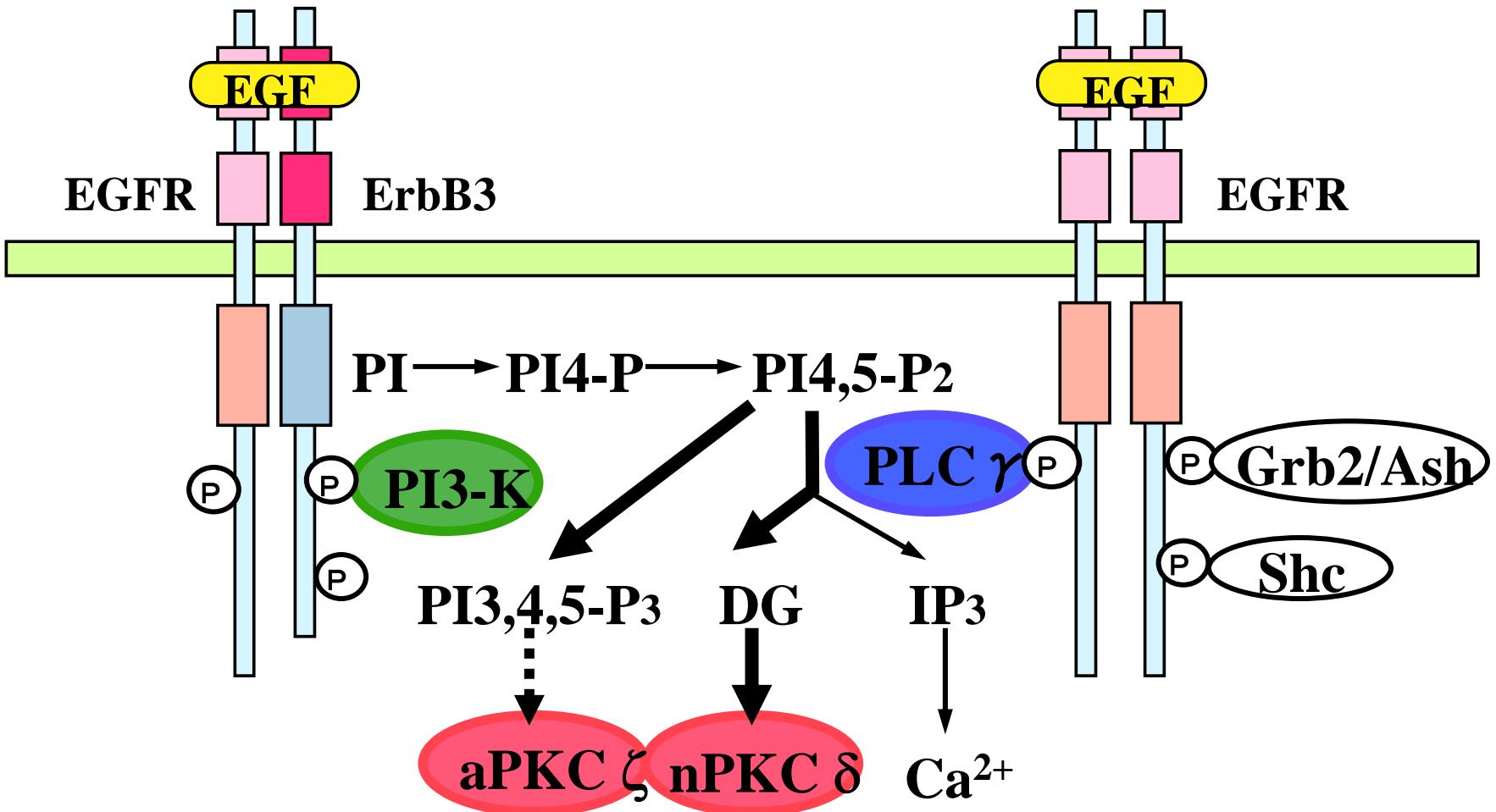


**aPKC-**

# Localization of intracellular PKC- $\delta$ and PKC- $\zeta$ in EGF and signal transduction inhibitors treated s-1 cells.



# Possible Signal Transduction Pathways Involved in Cell Motility Induced by EGF



# EGF受容体を分子標的とした薬剤

	薬剤	標的分子・特徴	
低分子化合物	ZD1839	EGF-R Tyrosine K	Iressa
	OSI-774	EGF-R Tyrosine K	PI3K MAPK
	EKB-569	EGF-R Tyrosine K	
	Cl-1033	EGF-R Tyrosine K	
	PKI-116	EGF-R Tyrosine K	
	PD183805	EGF-R Tyrosine K	
	CGP-59362A	EGF-R Tyrosine K	
抗体	IMC-C225	キメラ抗EGF-R 抗体	Cetuximab
	h-R3	ヒト化抗EGF-R 抗体	
	MDX-447	抗EGF-R, CD64抗体	
	ABX-EGF	ヒト化抗 EGF-R抗体	

# Tumorigenicity of ER-1 cells treated with EGF

- 24 hours treatment

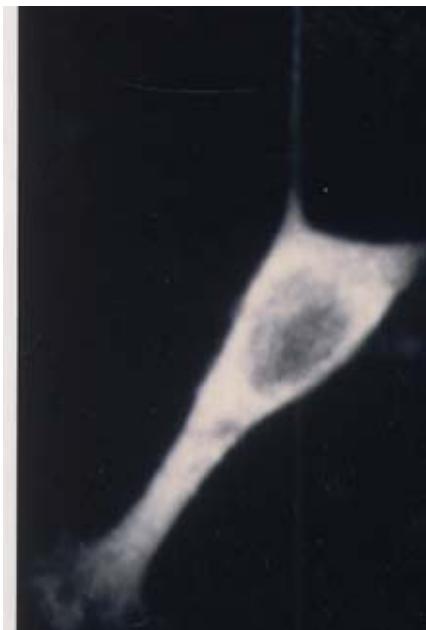
EGF	Tumor take
None	0/5
100ng/ml	5/5
100ng/ml (4 days EGF free)	0/5

- 1 month treatment

EGF	Tumor take
None	0/5
100ng/ml (1 month EGF free)	4/5
100ng/ml (2 month EGF free)	5/5

ER-1 cells 1x10<sup>5</sup> ip

DCFDA

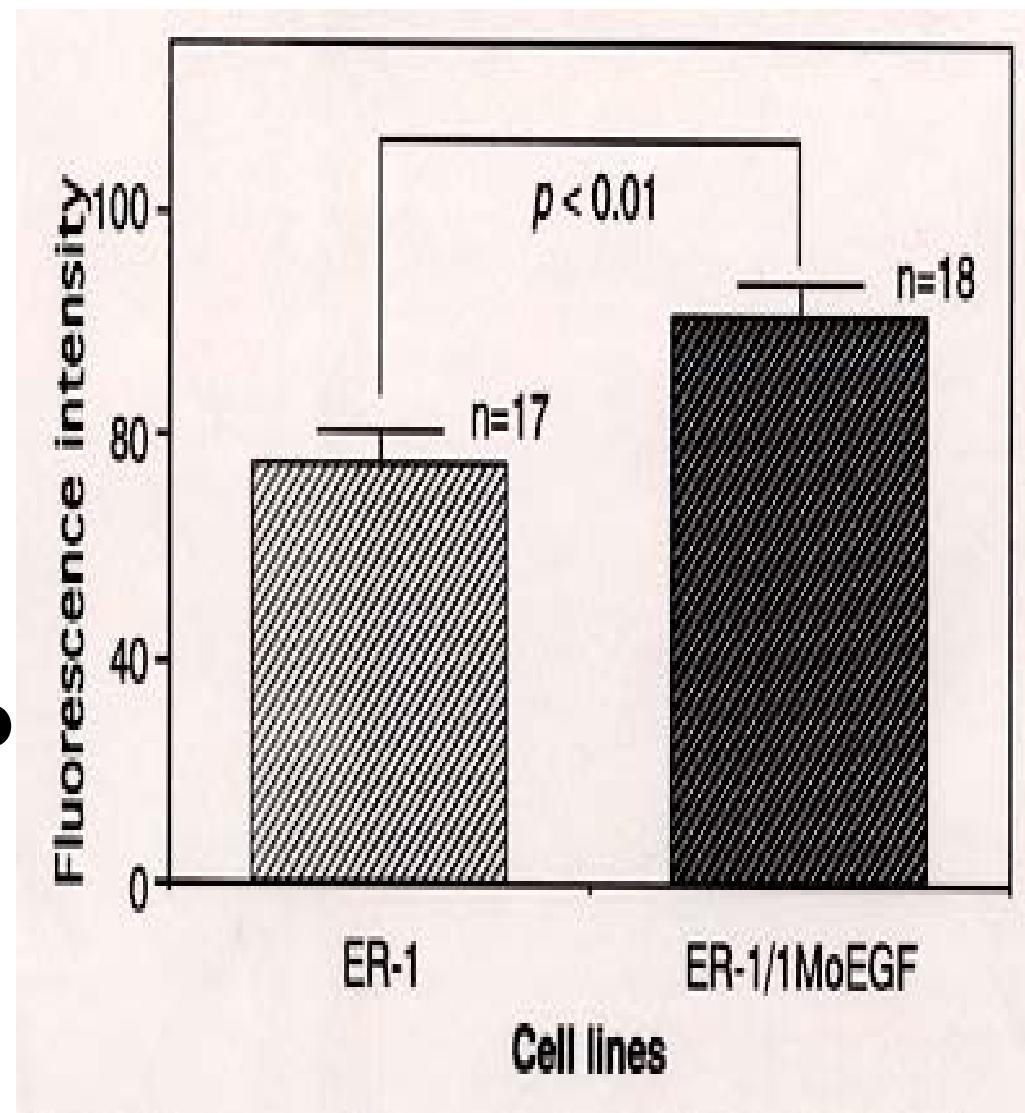


ER-1

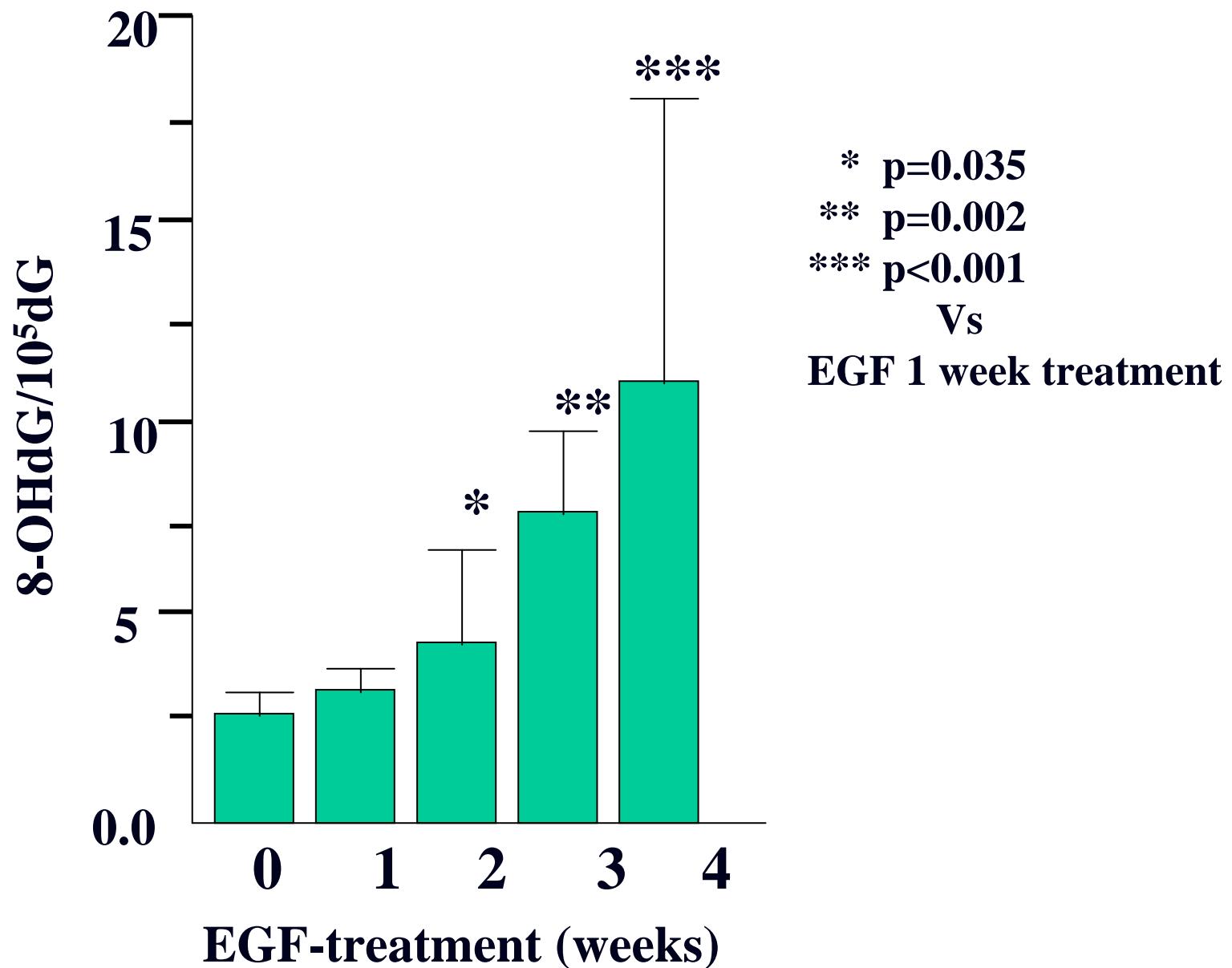


EGF 1 mo  
treated  
ER-1

## Intracellular oxidized state in ER-1 cells



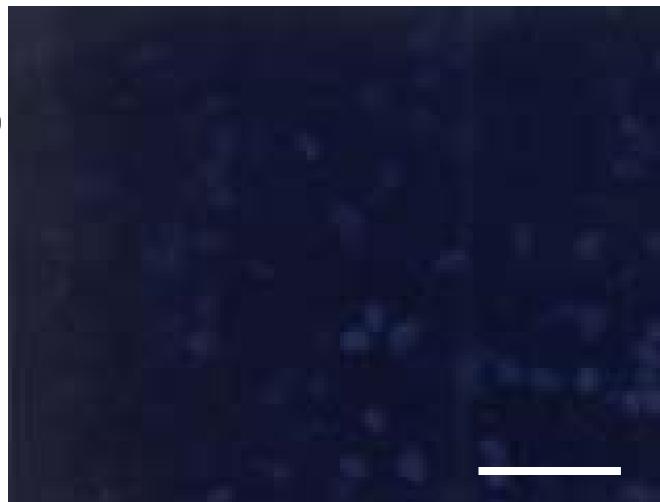
# Levels of 8-OHdG in ER-1 cells depending on the length of EGF treatment



# Inhibition of EGF-induced intracellular peroxides by N-acetylcysteine (antioxidant)

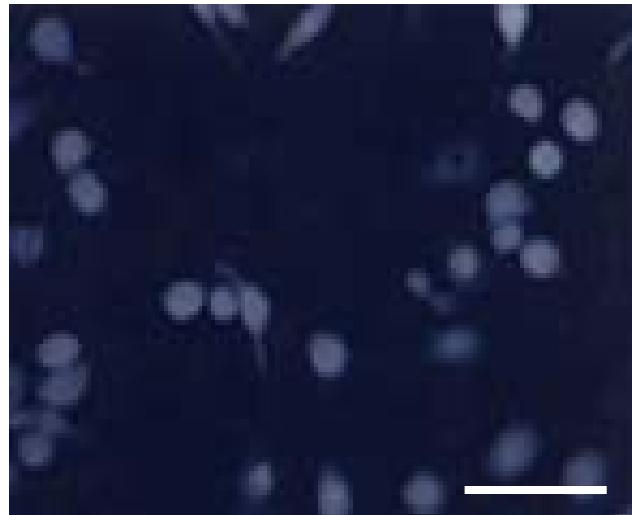
Using of DCFDA

EGF (-)  
100ng/ml



NAC (-)  
10mM

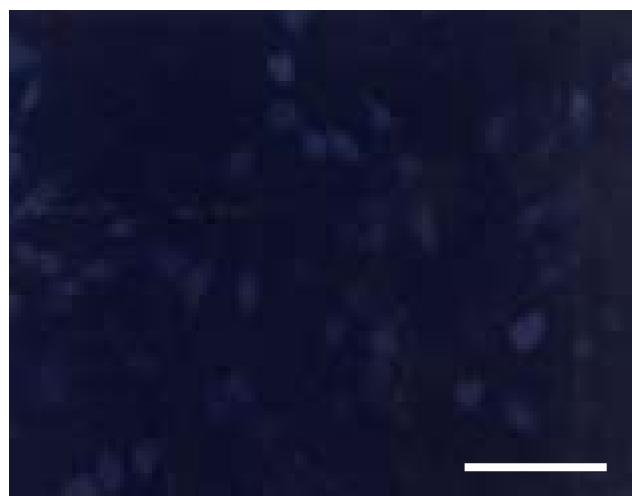
EGF (+)  
NAC (-)



EGF (-)  
NAC (+)



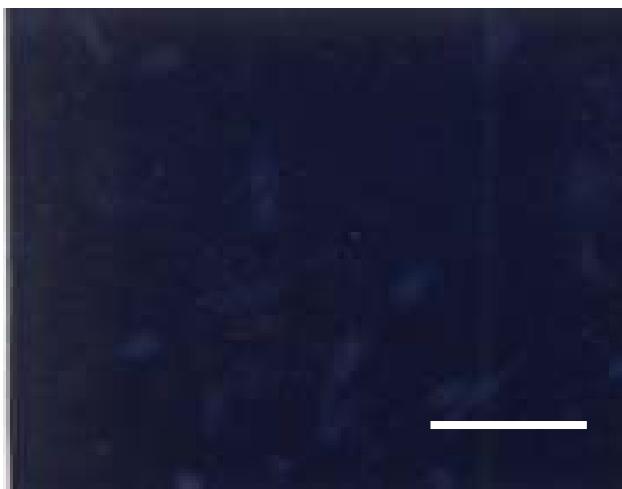
EGF (+)  
NAC (+)



# Inhibition of EGF-induced intracellular peroxides by sodium selenite (antioxidant)

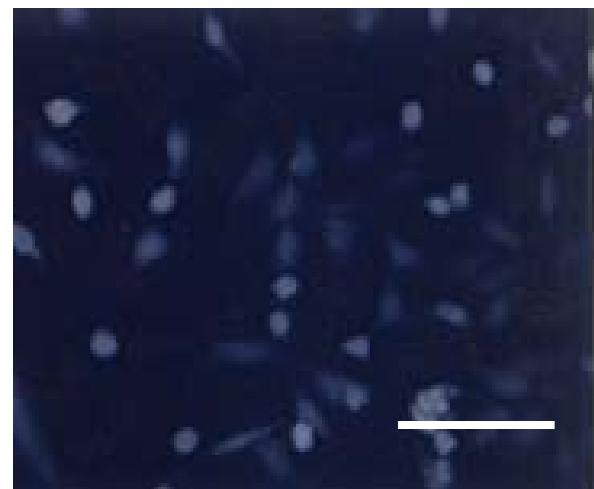
Using of DCFDA

EGF(-)  
100ng/ml



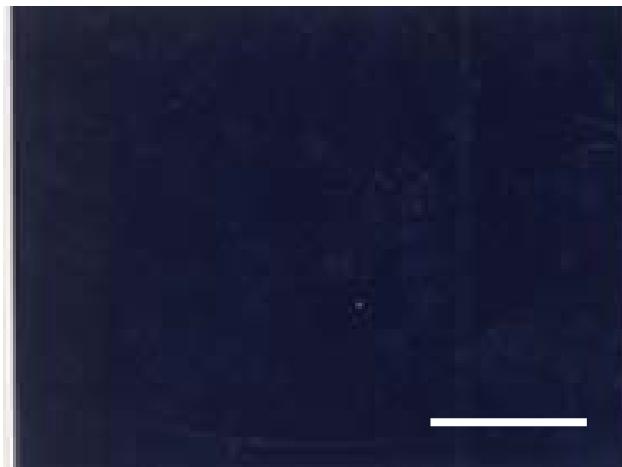
Se (-)  
100ng/ml

EGF(+)  
Se (-)



EGF(-)

Se (+)



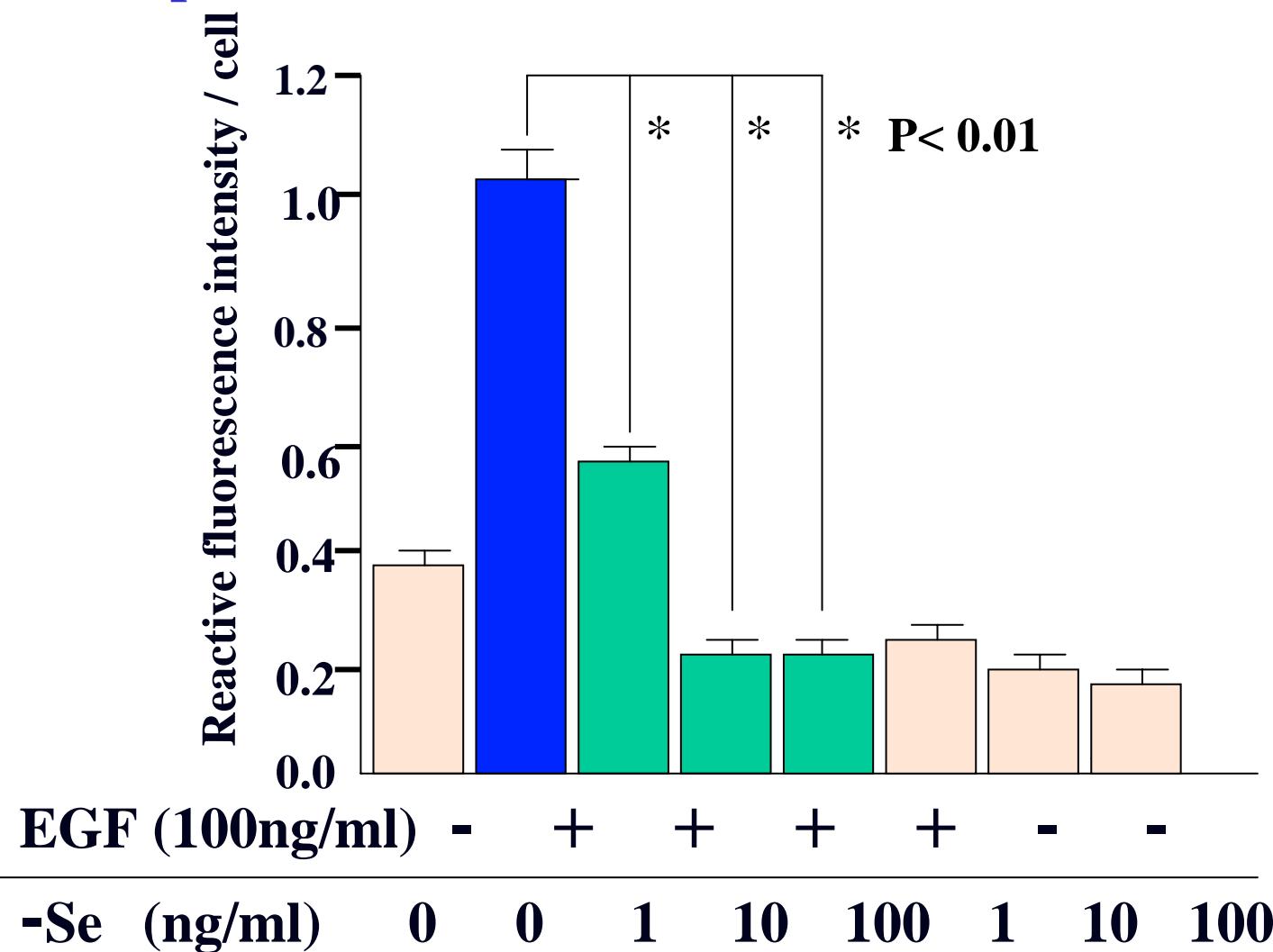
EGF(+)

Se (+)



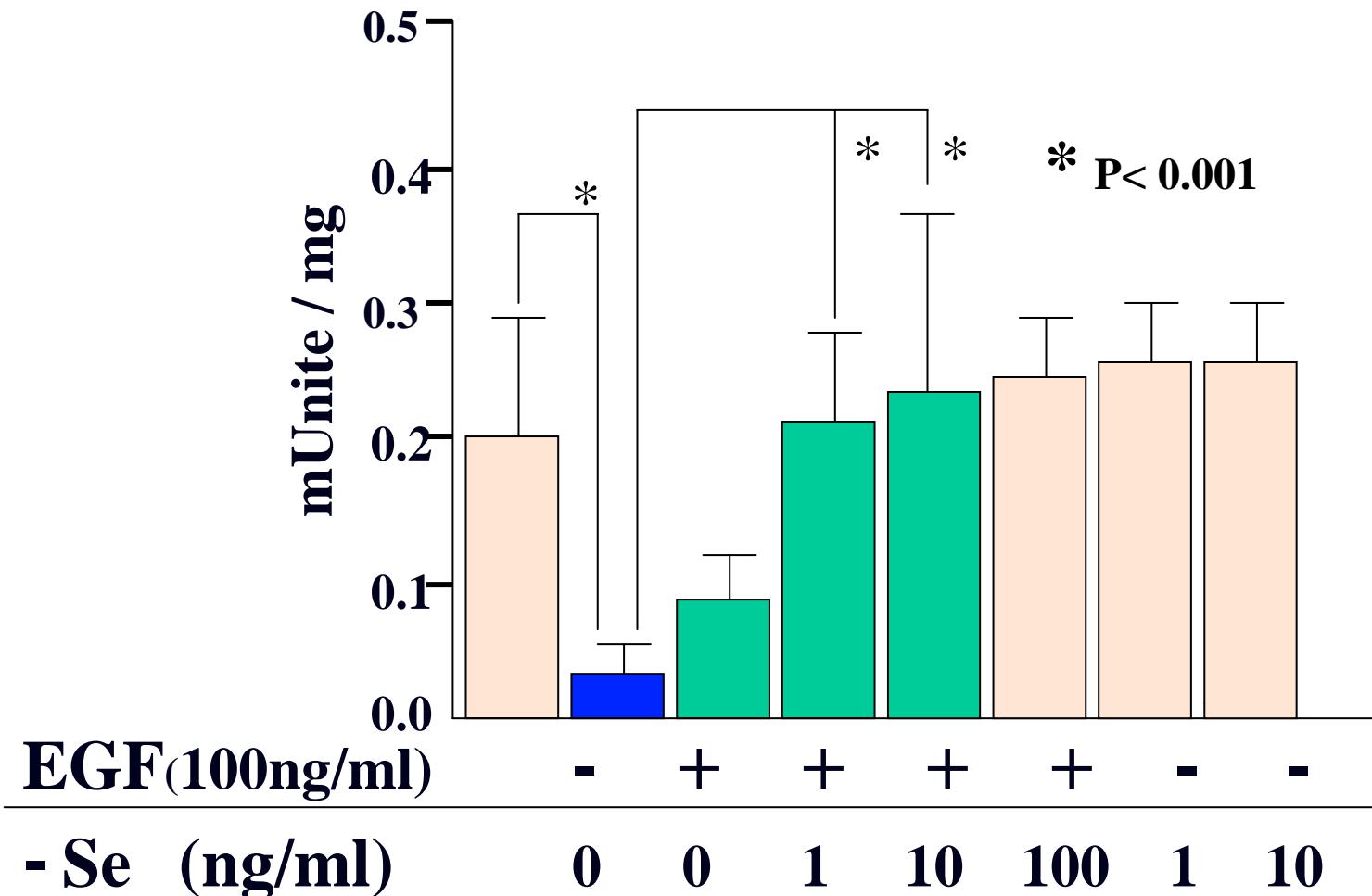
# Inhibition of EGF-induced intracellular peroxides by sodium selenite

(Levels of peroxidase)

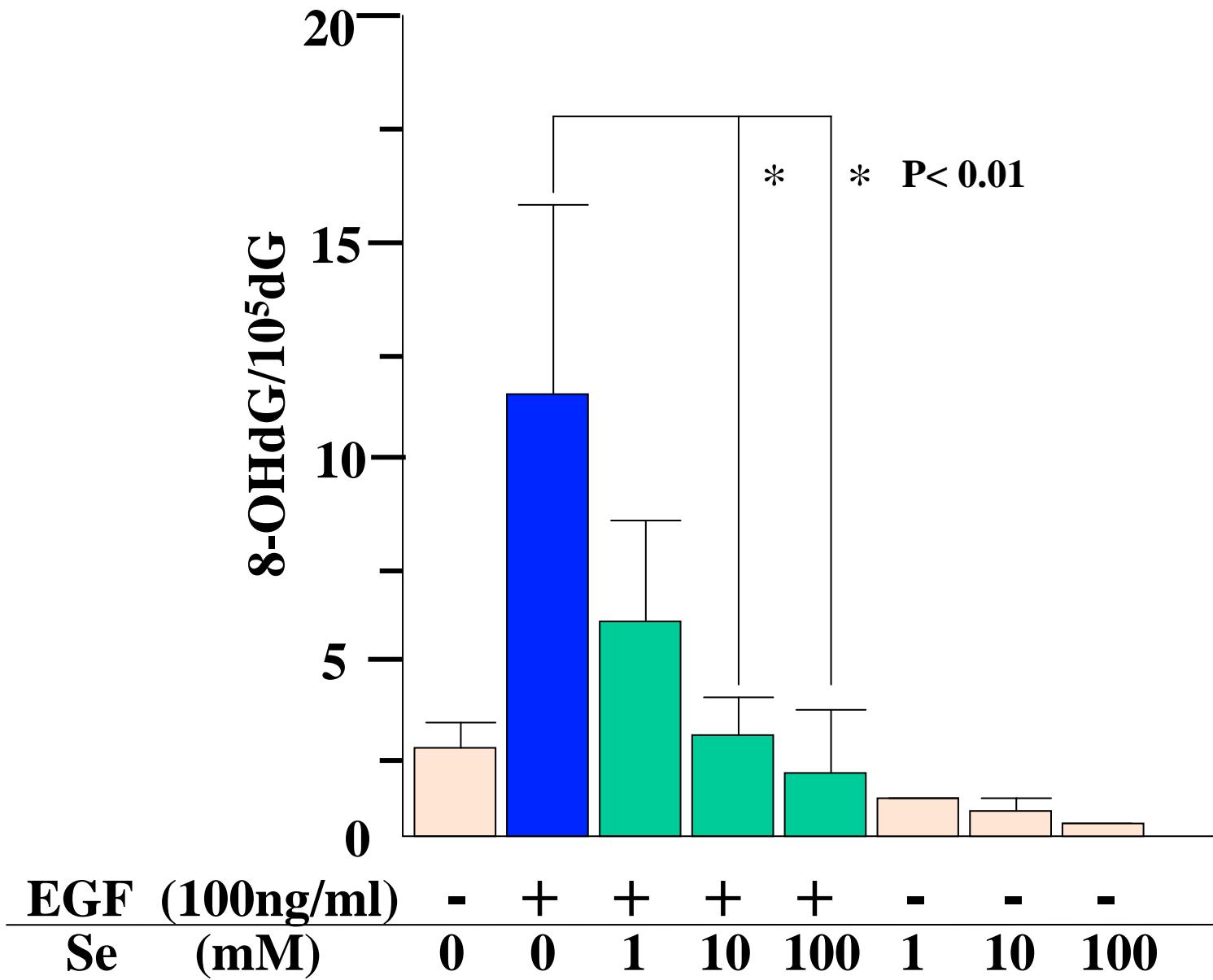


# Inhibition of EGF-induced intracellular peroxides by sodium selenite

(Levels of glutathione peroxidase activity)

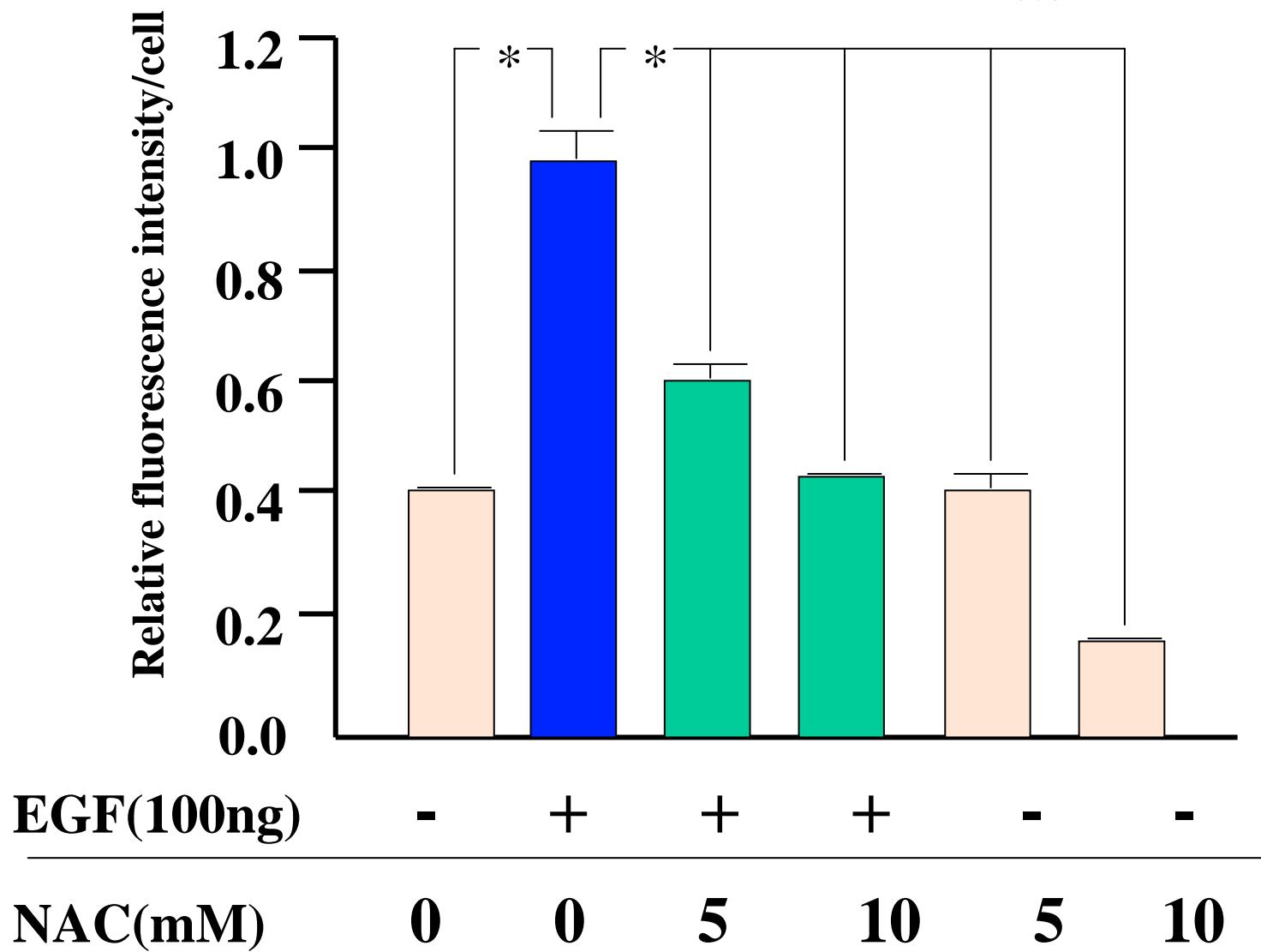


# Levels of 8-OHdG in ER-1 cells treated with EGF and /or sodium selenite (Se)

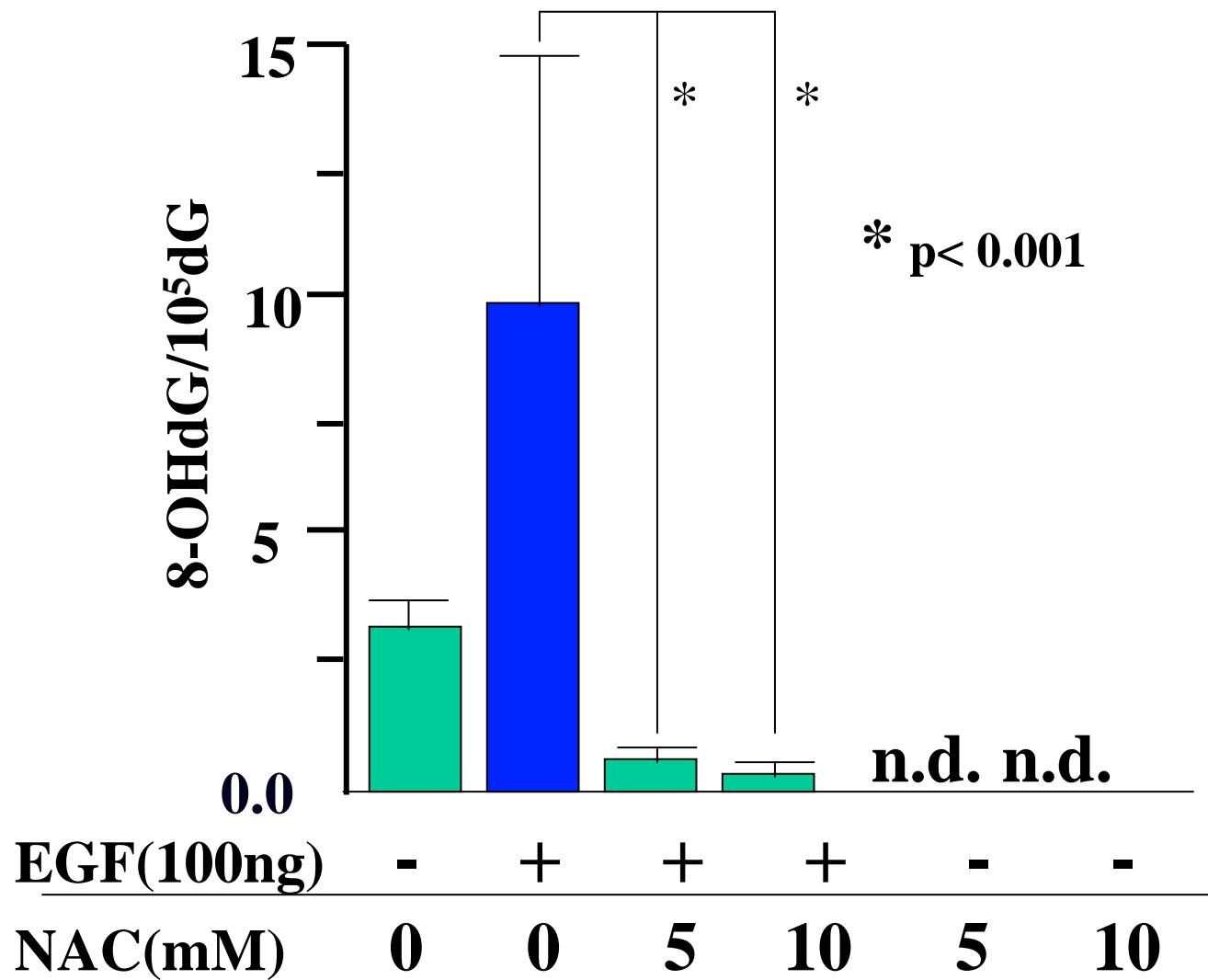


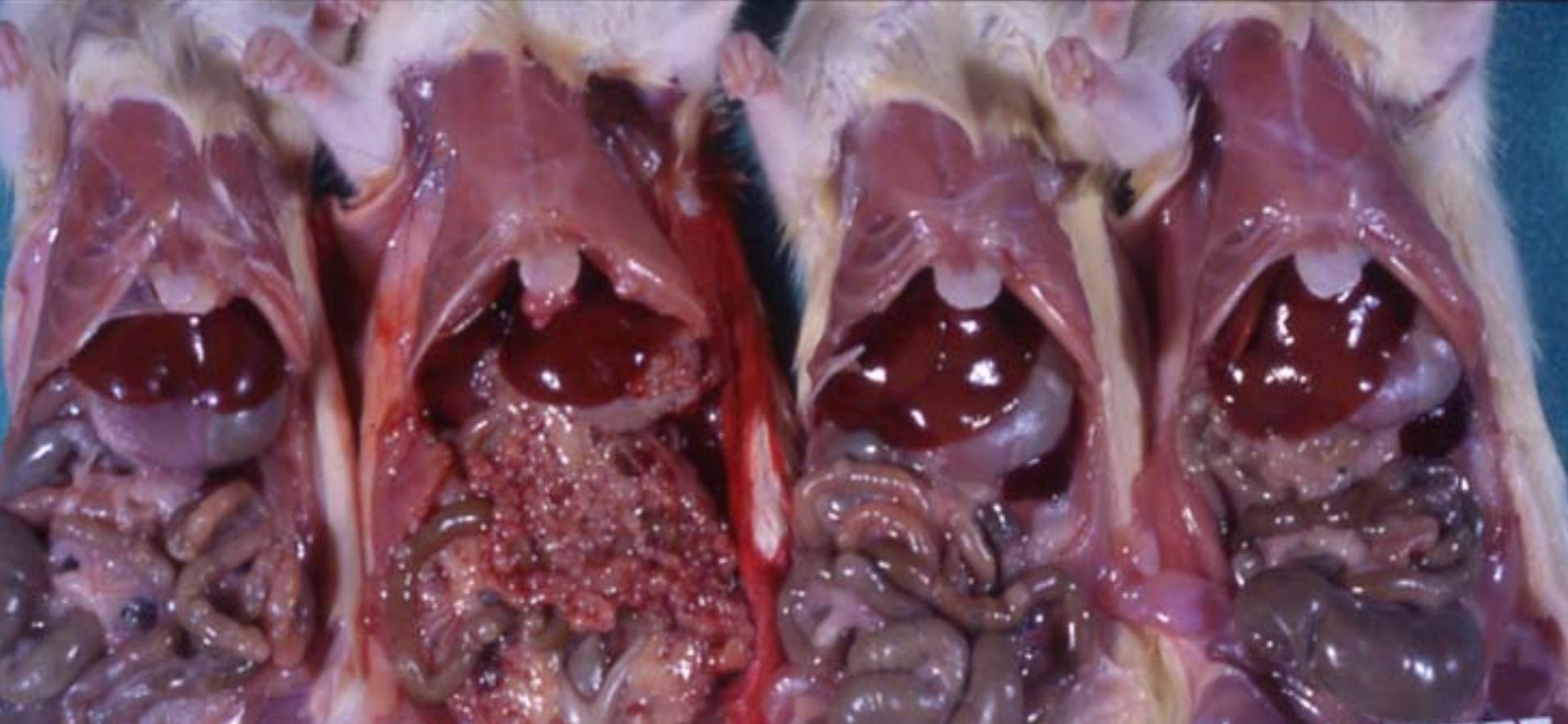
# Inhibition of EGF-induced intracellular peroxides by NAC

\* P< 0.01



# Levels of 8-OHdG in ER-1 cells treated with EGF and /or N-acetylcysteine (NAC)





**EGF (-)**  
**Se (-)**

**EGF (+)**  
**Se (-)**

**EGF (-)**  
**Se (+)**

**EGF (+)**  
**Se (+)**

**EGF(100ng/ml) +/- Sodium Se(100ng/ml) 1 mo**

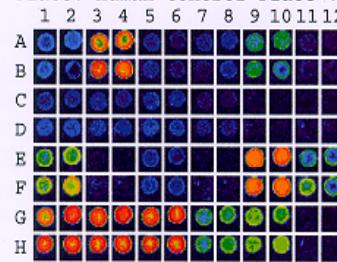
# **Tumorigenicity of ER-1 cells treated with EGF in the presence or absence of NAC or Selenite**

<b>Treatment</b>			
<b>EGF</b> <b>ng/ml</b>	<b>NAC</b> <b>mM</b>	<b>Selenite</b> <b>ng/ml</b>	<b>Tumor take</b>
0	0		0/6
100	0		6/6
100	5		1/6
100	10		1/6
0	5		0/5
0	10		0/5
0	0		0/6
100	0		6/6
100	1		6/6
100	10		2/6
100	100		1/6
0	1		0/6
0	10		1/6
0	100		1/6

# Micro-Array Analysis ---- Comparison Between S-1 and I-3 cells

CY3: s1.kuraboshibata (123B6326)

Plate: Human Control Plate v3 (021Y9970)



CY5: I3.kuraboshibata (123I6327)

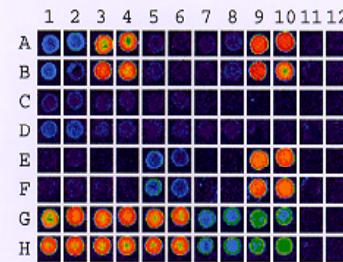


Plate: UGV212.IN.A-A.02148718 (0210A5MJ)

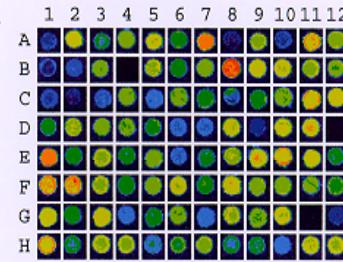
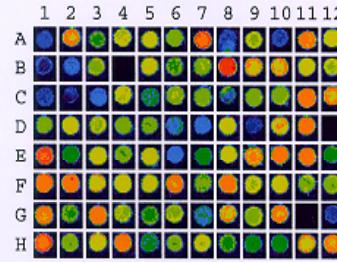


Plate: UGV212.IN.A-A.021B8719 (021WA4V3)

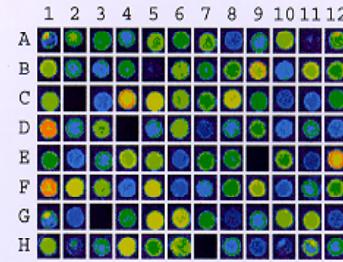
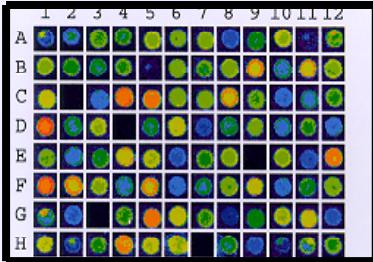
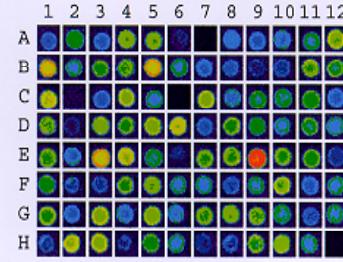
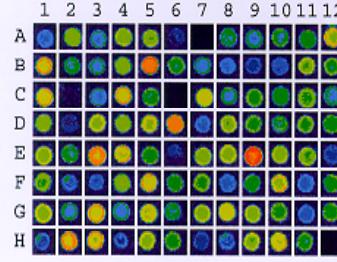


Plate: UGV212.IN.A-A.021Q8720 (0213A4V4)

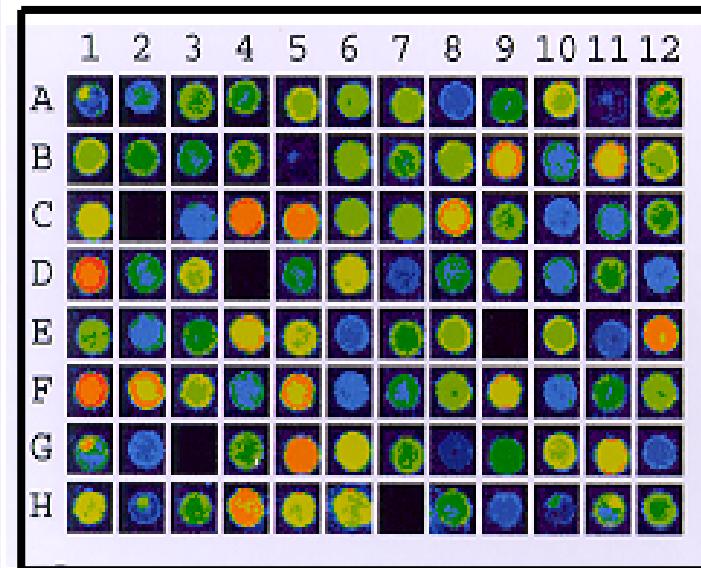


S-1

CY3(Red)

I-3

CY5(Green)



# Comparison of mRNA Expression between S-1 and I-3 cells

matrix metalloproteinase 1 (interstitial collagenase)	spermidine/spermine N1-acetyltransferase
proteasome (prosome, macropain) subunit, beta type, 9 (large multifunctional protease 2)	M-phase phosphoprotein 4
heparin-binding growth factor binding protein	MHC class I region ORF
keratin 13	ESTs, Moderately similar to ORF derived from protease and integrase coding regions [H.sapiens]
caveolin 1, caveolae protein, 22kD	requiem, apoptosis response zinc finger gene
protein kinase, cAMP-dependent, catalytic, alpha	Ste-20 related kinase
matrilin 2	transcription factor AP-2 alpha (activating enhancer-binding protein 2 alpha)
ESTs	high-mobility group (nonhistone chromosomal) protein 2
serine/threonine kinase 15	keratin 6B
annexin A1	matrix metalloproteinase 13 (collagenase 3)
KIAA0069 protein	Incyte EST
sperm specific antigen 2	coagulation factor III (thromboplastin, tissue factor)
ataxia-telangiectasia group D-associated protein	contactin 1
centromere protein A (17kD)	serine/threonine kinase 17a (apoptosis-inducing)
cadherin 13, H-cadherin (heart)	low density lipoprotein receptor (familial hypercholesterolemia)
karyopherin alpha 2 (RAG cohort 1, importin alpha 1)	laminin, gamma 1 (formerly LAMB2)
integrin, alpha 6	synaptophysin-like protein
EGF-containing fibulin-like extracellular matrix protein 1	butyrophilin, subfamily 3, member A2

**Genome\_Systems**

Fab: 7/26/2000 Hyb: 10/13/2000 Scan: 10/13/2000 Analyze: 10/13/2000 Ready: 10/17/2000

GEM Notes:

CY3: si10.kurabo.shibata (123I8109)

Plate: Human Control Plate v3 (021Y9970)

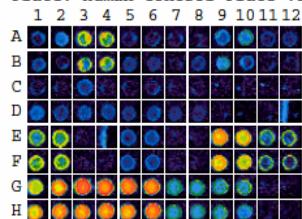


Plate: UGV2.27.IN.A-A.02148718 (0210A5MJ)

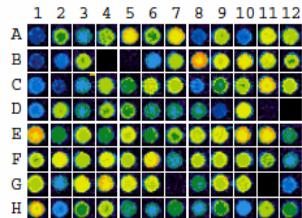


Plate: UGV2.27.IN.A-A.021B8719 (021WA4V3)

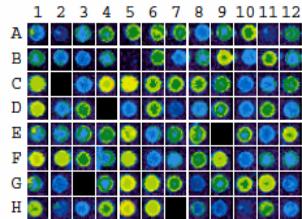
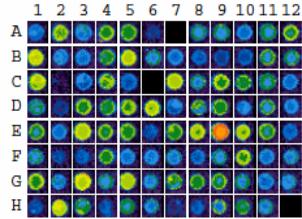
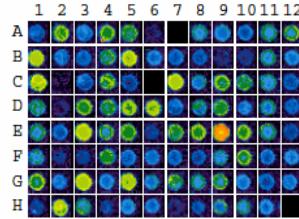
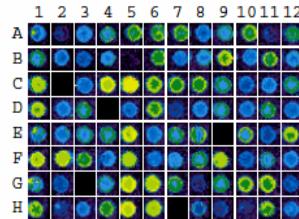
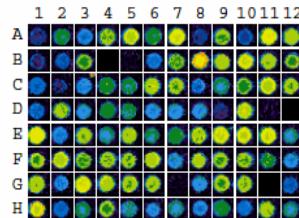
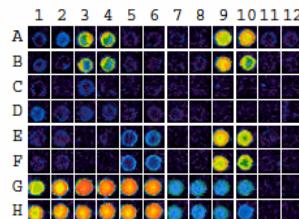


Plate: UGV2.14.IN.A-A.021Q8720 (0213A4V4)

**IncyteGenomics**

CY5: siegf10.kurabo.shibata (123X8110)

**Genome\_Systems**

Fab: 7/26/2000 Hyb: 10/13/2000 Scan: 10/13/2000 Analyze: 10/13/2000 Ready: 10/17/2000

GEM Notes:

CY3: si10.kurabo.shibata (123I8109)

Plate: UGV2.27.IN.A-A.021X8721 (021AA4V5)

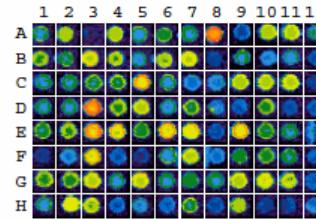


Plate: UGV2.27.IN.A-A.02148722 (021HA4V6)

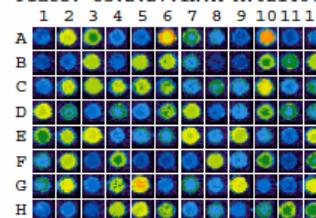


Plate: UGV2.27.IN.A-A.021B8723 (021OA4V7)

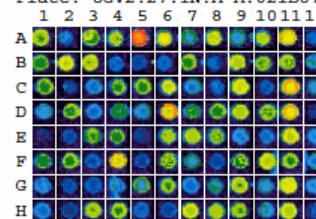
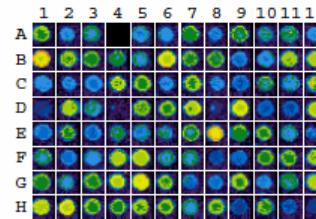


Plate: UGV2.27.IN.A-A.021I8724 (021VA4V8)

**IncyteGen**

CY5: siegf10.kurabo.shibata (123X8110)

