

Repeat1

cropped

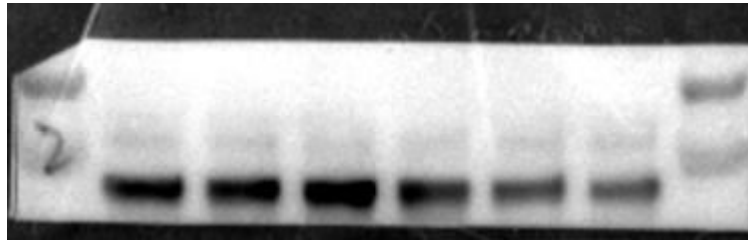
mk	nc	nc	nc	siR NA	siR NA	siR NA

Fn (ab23750)  
270kd  
R



270kd

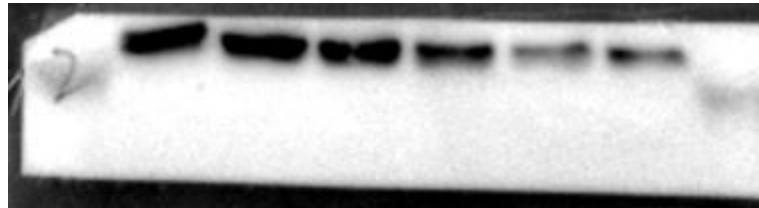
INHBA(A5232)  
45KD  
R



66kd

52kd

Snail (A11794)  
34KD  
R



30kd

GAPDH (6004-  
1-Ig)  
37KD  
M

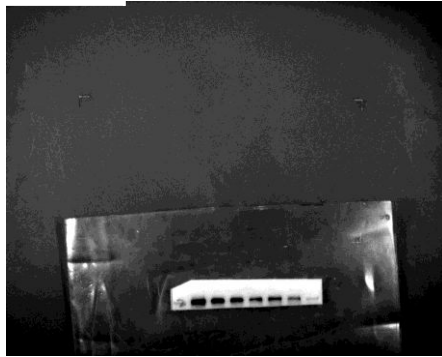


37kd

NRK-49F cells were transfected with nonsense control (NC) or INHBA siRNA (siINHBA). After 24h, cell lysates were collected and Western blot analysis was performed to measure the expression of INHBA, FN and Snail.

Fn (ab23750)  
270kd  
R

original



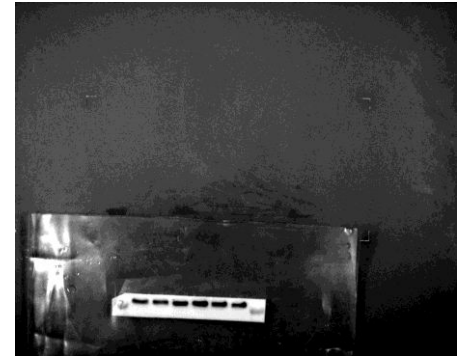
merged

GAPDH (6004-  
1-Ig)  
37KD  
M

original

merged

INHBA(A5232)  
45KD  
R



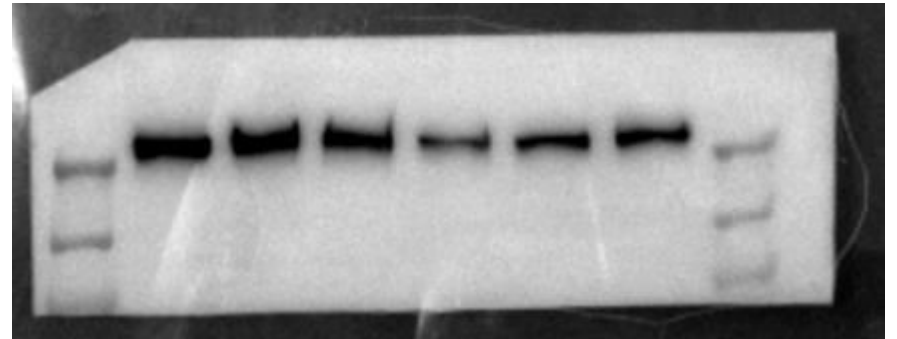
Snail (A11794)  
34KD  
R



Repeat2

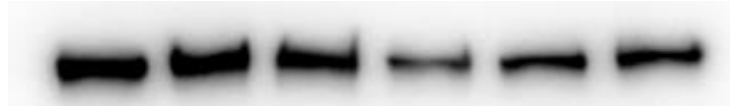
cropped

mk	Nc	Nc	nc	siR NA	siR NA	siR NA	mk
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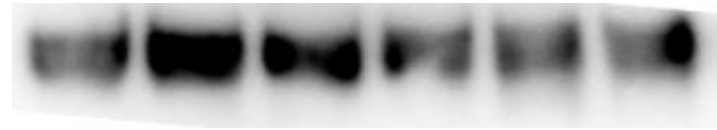


270kd  
170kd  
130kd

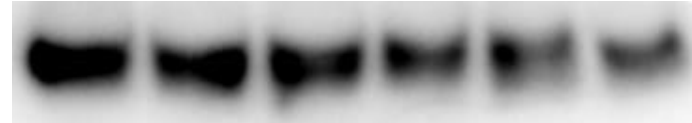
**Fn (ab23750)**  
270kd  
R



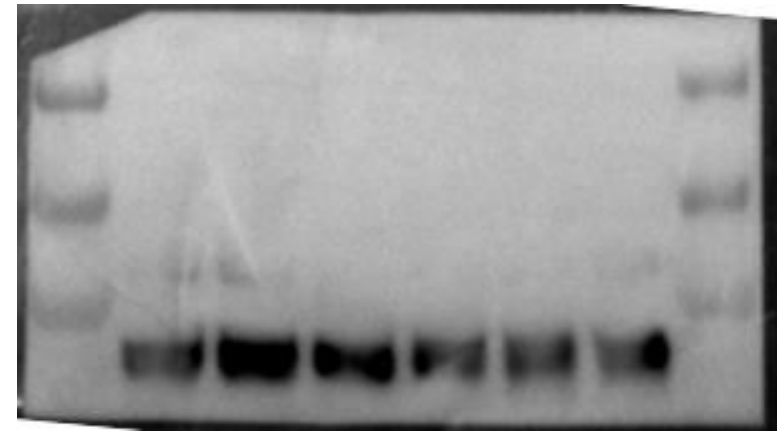
**INHBA(A5232)**  
45KD  
R



**Snail (A11794)**  
34KD  
R



**GAPDH (6004-1-Ig)**  
37KD  
M



90kd  
66kd  
52kd



30kd



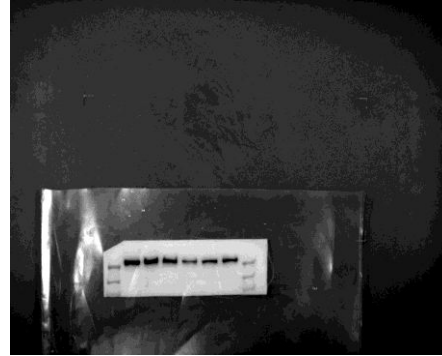
37kd

Fn (ab23750)  
270kd  
R

original



merged



GAPDH (6004-  
1-Ig)  
37KD  
M

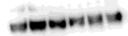
original



merged



INHBA(A5232)  
45KD  
R



Snail (A11794)  
34KD  
R



Repeat3

cropped

mk	Nc	Nc	siR NA	siR NA



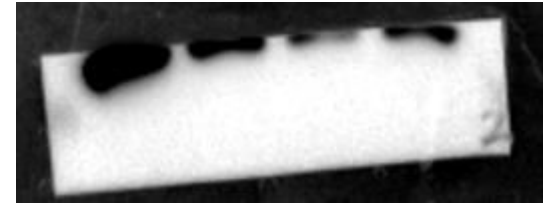
270kd

Fn (ab23750)  
270kd  
R



52kd

INHBA(A5232)  
45KD  
R



30kd

Snail (A11794)  
34KD  
R



37kd

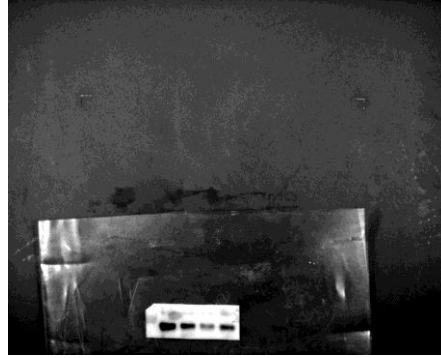
GAPDH (6004-  
1-Ig)  
37KD  
M

INHBA(A5232)  
45KD  
R

original



merged

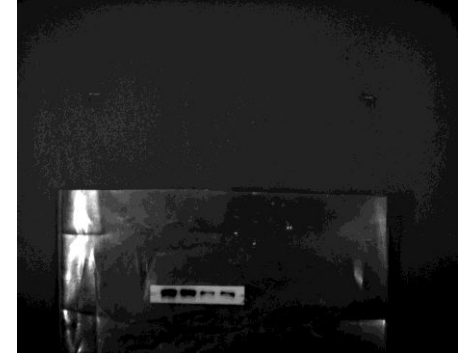


Fn (ab23750)  
270kd  
R

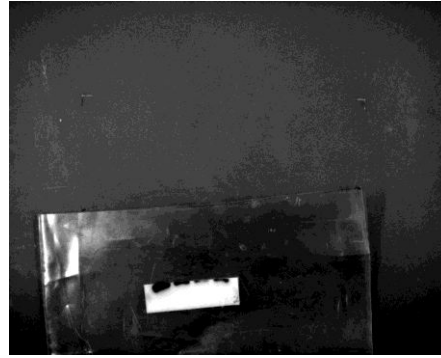
original



merged



Snail (A11794)  
34KD  
R



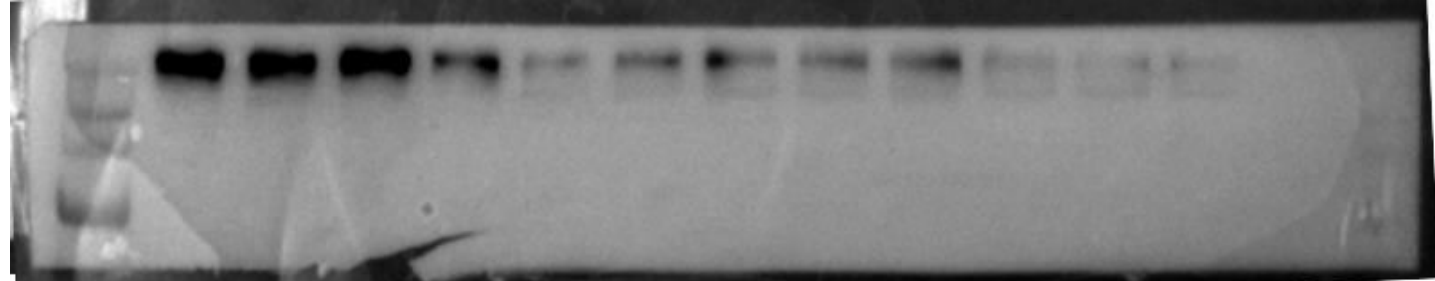
GAPDH (6004-  
1-Ig)  
37KD  
M



Repeat1  
cropped

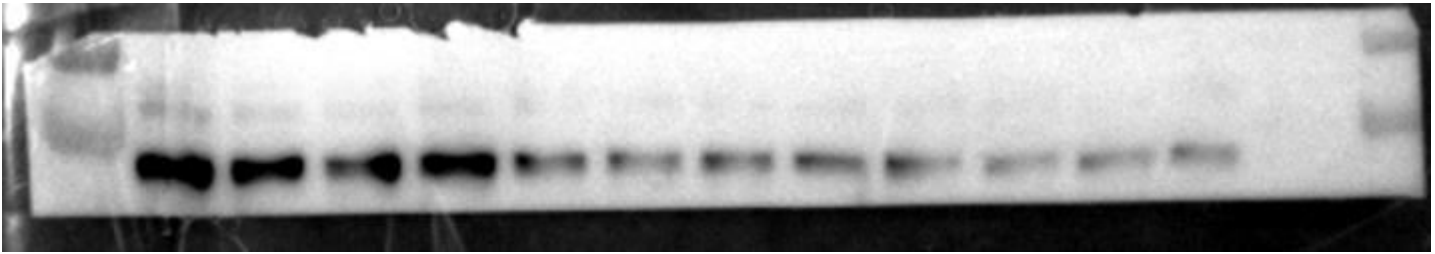
mk	Nc			NC			siRNA			siRNA			mk
	dms0	dms0	dms0	Tan-I	Tan-I	Tan-I	dms0	dms0	dms0	Tan-I	Tan-I	Tan-I	

Fn (ab23750)  
270kd  
R



270kd  
170kd  
130kd  
90kd

INHBA(A5232)  
45KD  
R



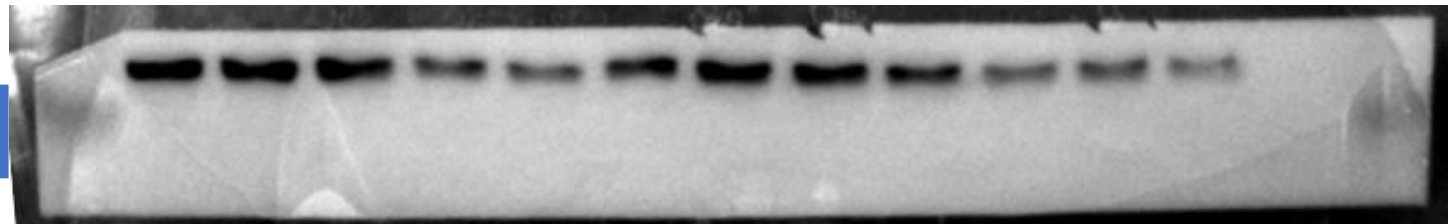
66kd  
52kd

GAPDH (6004-1-Ig)  
37KD  
M



37kd

Snail (A11794)  
34KD  
R



30kd

NRF-49F cells were transfected with NC or siINHBA. On the second day, cells were treated 50  $\mu$ M of Tan-I for another 24h. Cell lysates were collected and Western blot analysis was performed to measure the expression of INHBA, FN and Snail.



Fn (ab23750)  
270kd  
R

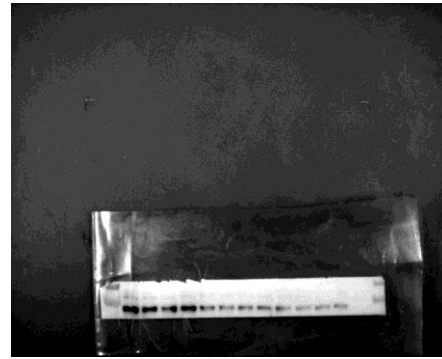
original



merged



INHBA(A5232)  
45KD  
R



Snail (A11794)  
34KD  
R

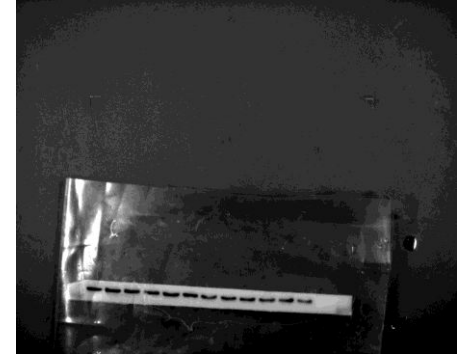


GAPDH (6004-1-Ig)  
37KD  
M

original



merged



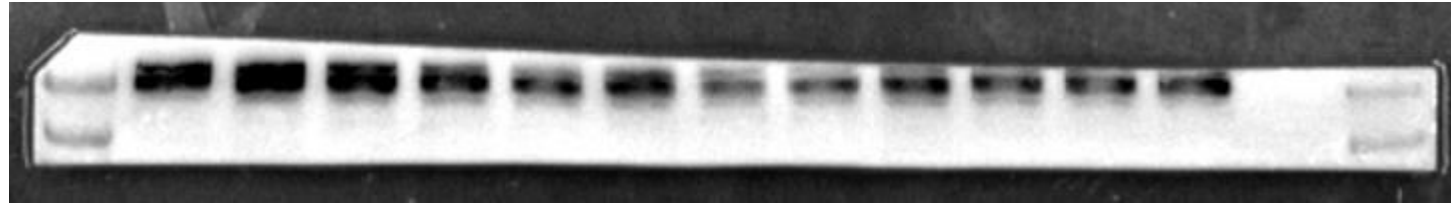


Repeat2

cropped

mk	Nc			NC			siRNA			siRNA			mk	
	dms0	dms0	dms0	Tan-I	Tan-I	Tan-I	dms0	dms0	dms0	Tan-I	Tan-I	Tan-I		

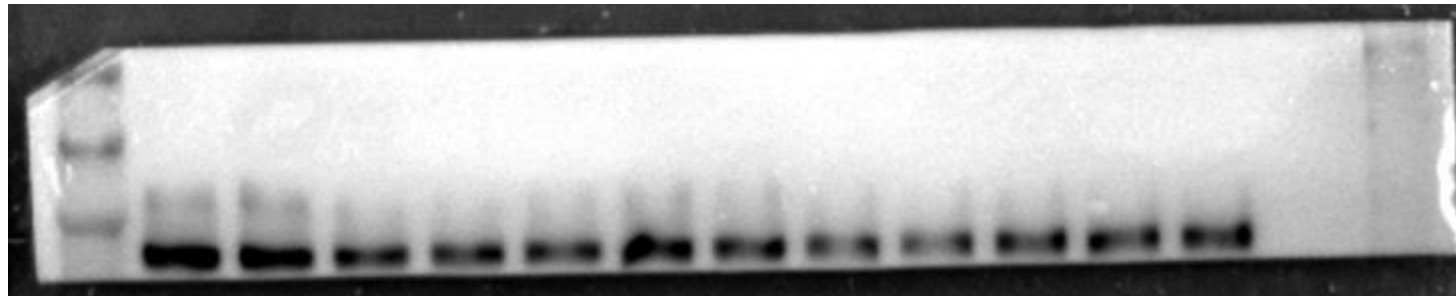
Fn (ab23750)  
270kd  
R



270kd

170kd

INHBA(A5232)  
45KD  
R

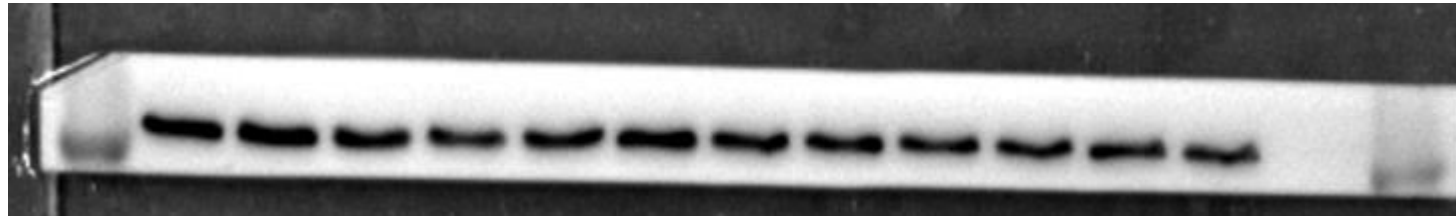


90kd

66kd

52kd

GAPDH (6004-1-Ig)  
37KD  
M



37kd

Snail (A11794)  
34KD  
R



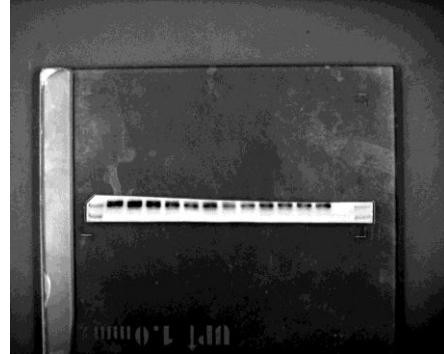
30kd

Fn (ab23750)  
270kd  
R

original

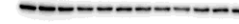


merged



Snail (A11794)  
34KD  
R

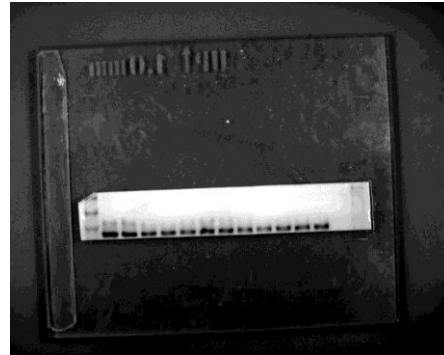
original



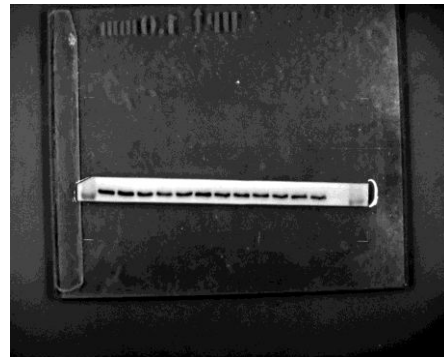
merged



INHBA(A5232)  
45KD  
R



GAPDH (6004-1-Ig)  
37KD  
M



Repeat3

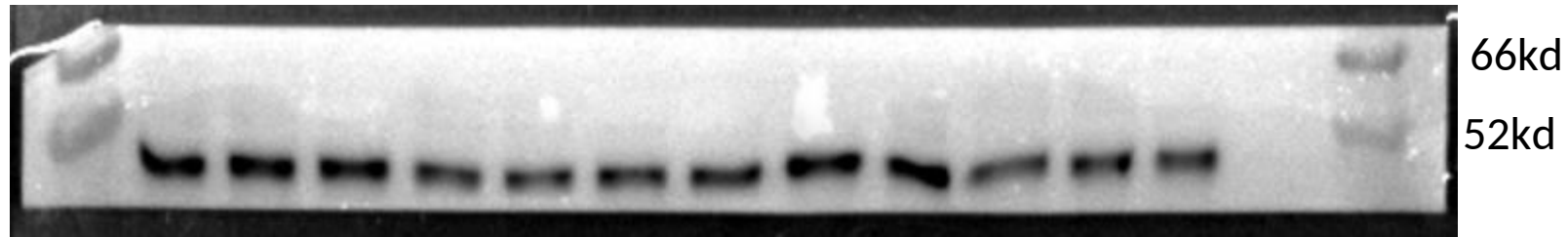
cropped

mk	Nc			NC			siRNA			siRNA				mk
	dms0	dms0	dms0	Tan-I	Tan-I	Tan-I	dms0	dms0	dms0	Tan-I	Tan-I	Tan-I		

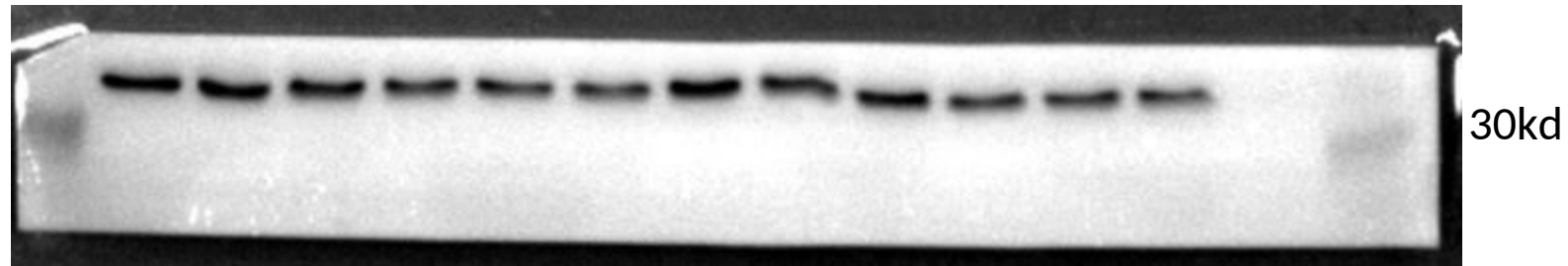
Fn (ab23750)  
270kd  
R



INHBA(A5232)  
45KD  
R



Snail (A11794)  
34KD  
R



GAPDH (6004-1-Ig)  
37KD  
M

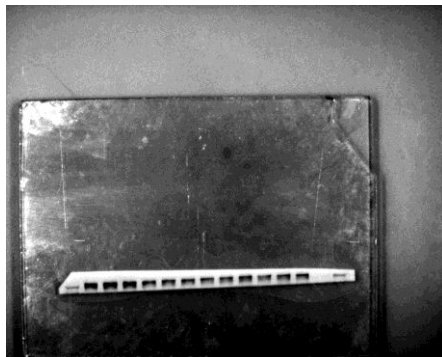


Fn (ab23750)  
270kd  
R

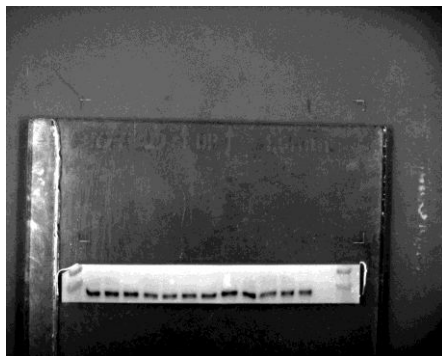
original



merged



INHBA(A5232)  
45KD  
R



Snail (A11794)  
34KD  
R



GAPDH (6004-1-Ig)  
37KD  
M

original



merged

