

Chemilum. Quantification of cpY31 uncaging

From uncaging on p. 20, a pan and pY31 gel were run. pY31 paxillin (semisynthesized) was used as an internal standard.

1° mouse anti-pan pax (1/11,000)
2° goat anti-mouse HRP (1/10,000)

1° rabbit anti-pY31 pax (1/2,000)
2° goat anti-rabbit HRP (1/20,000)



exposed \approx 2 min. 4 3 2 1

exposed = 2 min. 4 3 2 1

- In
- 1 6.4 μ L benchmark protein ladder
 - 2 6.4 μ L semisynthetic cpY31 (see p. 6-19)
 - 3 6.4 μ L semisynthetic cpY31 uncaged for 90 s.
 - 4 3 μ L of 1/5 semisynthetic pY31

To quantify the density of selected bands:

- zoom in on area w/ bands (zoom icon on top left of screen)
- Under volume select

Calculations of % uncaged based on band density quantification: Oct. 18, 2005

w/ pan 1° ab

		density	st. dev.	density - bckgd	% of internal standard pY31 pax ↓
pY31	U3	7711	(362)	5500	1
uncaged	U2	5508	(293)	3297	0.599
caged	U1	5615	(276)	3404	0.619
bckgrnd	B1	2211	(6)	-	-

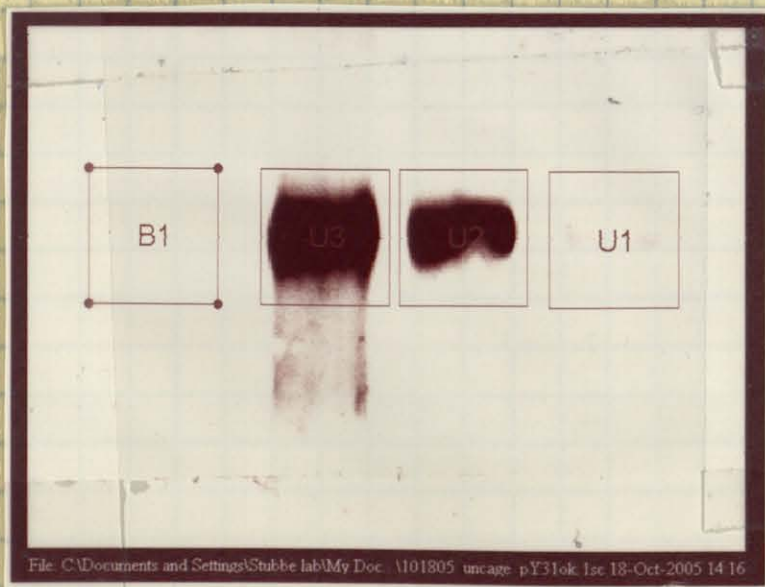
w/ pY31 1° ab

		density	st. dev.	density - bckgd	% of standard
pY31	U3	10650	(637)	8378	1
uncaged	U2	6660	(438)	4388	0.524
caged	U1	2609	(50)	337	0.040
bckgd	B1	2272	(9)		

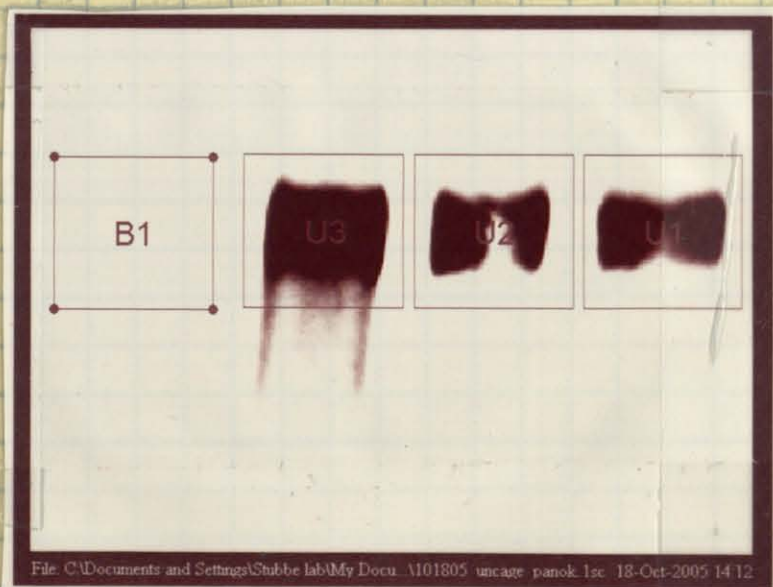
caged: % uncaged = $0.040 / 0.619 = 0.06499$ w/ st. dev. = $6.5 \pm 0.2 \%$

90 s. uncaged: % uncaged = $0.524 / 0.599 = 0.8737$ w/ st. dev. = $87 \pm 4 \%$

To propagate the standard deviation from each measurement, the % of standard was measured for each band (w/ both pan and pY31 ab) using the density for + and - the st. dev. (see excel spreadsheet for details).



pY31 1° ab



pan pax 1° ab