

EXHIBIT 11

LABORATORY NOTEBOOK

No. 325

APPLIED MOLECULAR GENETICS, INC.

LIN
EXHIBIT
116

AM 17 009862
CONFIDENTIAL
SUBJECT TO PROTECTIVE ORDER

FRITSCH v. LIN
INTER NOS 102,096
102,097, 107,714
LIN DOC NO L01099

Trial Exhibit ATT
97-10814-WGY

Trial Exhibit 217
97-10814-WGY

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ISSUED TO Jeri hane

ON 2/11 1983

DEPARTMENT _____

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— SCIENTIFIC NOTEBOOK CO. —
5007 WEST DONNA DRIVE
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FRITSCH v. LIN
INTER NOS 102,096
102,097, 102,314
LIN DOC NO L01100

Project No. _____
 Book No. _____ TITLE Solution Competition - ^{① Jeffs human epo}
^{② monkey sera}
^{③ 2001-2002}
^{④ HARRIS} ^{⑤ Duhan}

m Page No. _____

• 1/25/84

A variety of epo sources were tested for their ability to compete with standard epo in a solution phase against the anti-peptide Ab.

Procedure

- 1) Add 5 μ l trasytol / tube
- 2) Add sample to tubes along with an appropriate amt. of PBS+1% BSA so the final volume is 300 μ l
- 3) Add 50 μ l of a 1:2500 dilution of 8C-180 #17.0.6 to each tube
- 4) Incubate at 37° for 1 1/2 hrs.
- 5) Add 15 μ l I-125 epo to each tube. 18 μ l = 14,117 cpm.
- 6) Incubate at 40° for ~20 hrs
- 7) Add STAPH (150 μ l / tube: except monkey sera gets 600 μ l, 400 μ l, ~~200~~ 300 μ l depending on the size of the sample)
- 8) Proceed as usual

Results

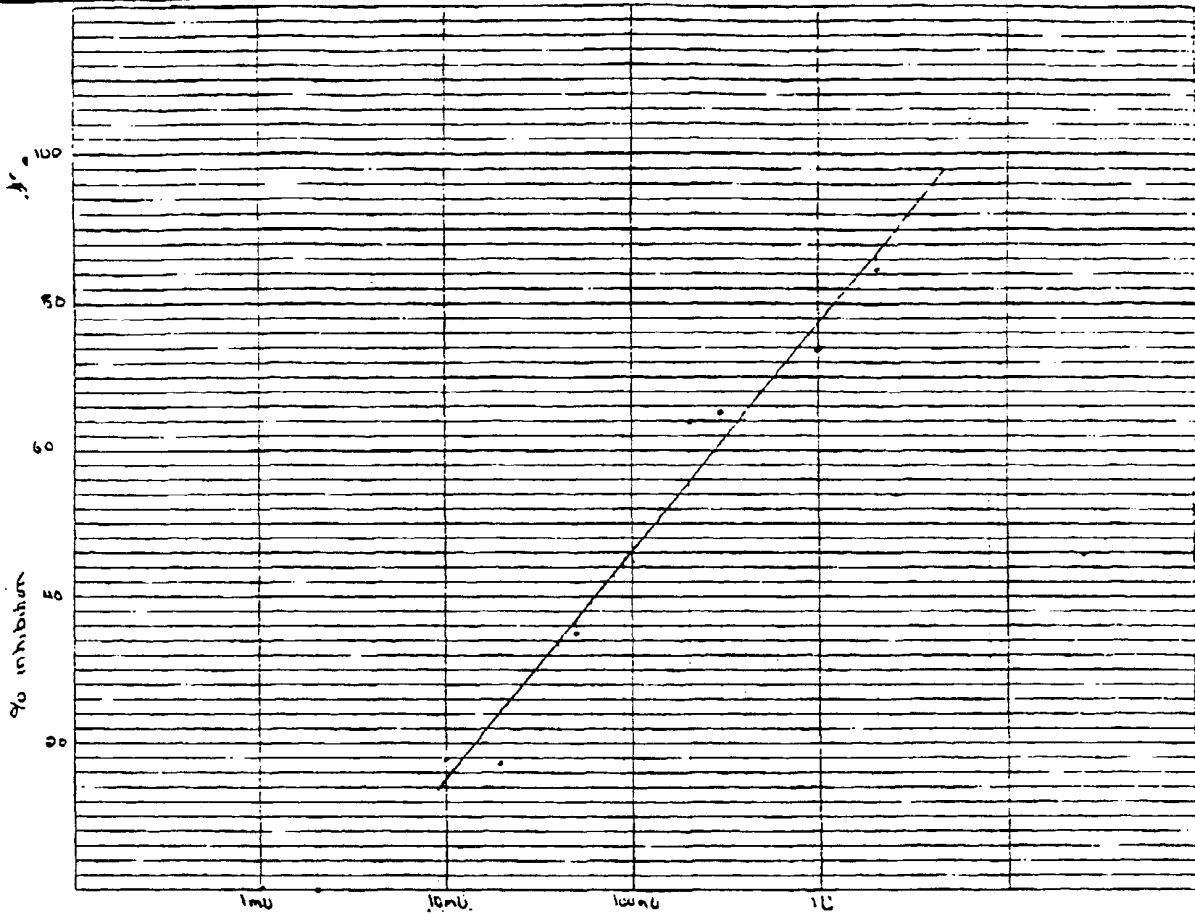
Sample	EPo or expected EPo	PBS+1% BSA	CPM	CPM - Pre-immune	% inh.	calculated epo EPo/ml
0	-	240	4883	4464		
0	-	240	5120			
5 μ l D = 1 ml		235	5068	4530	0	
10 μ l D = 2 ml		235	5176	4638	0	
5 μ l C = 10 ml		235	4248	3710	16.9	
10 μ l C = 20 ml		230	4305	3767	15.6	
2.5 μ l B = 50 ml		237	3446	2908	34.9	
5 μ l B = 100 ml		235	3018	2450	44.4	
10 μ l B = 200 ml		230	2142	1604	64.1	
15 μ l B = 300 ml		225	2075	1537	65.6	
5 μ l A = 1 U		235	1706	1162	74.0	
10 μ l A = 2 U		230	1231	693	84.5	
Pre-immune		240	441	538	Pre-immune	
Pre-immune		240	635			
human = E7.6 μ l 15.1		180	5252	4714	0	0
" 120 μ l 30.2		120	3995	3457	22.6	19 ml
human = M.6 μ l 0		180	6096	5558	0	0
" 120 μ l 0		120	6050	5512	0	0

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 102,097, 102,334
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FRITTSCH v. LIN 102,097, 102,334
 INTER NOS 102,096 LIN DOC NO L01102

page 77) using α -crude Ab in a 50μ competition showed the titer of E7 to be 35.4 mU/ml (before concentration). The mock was negative. This assay gives us 22.3 mU/ml for E7 and 0 for the mock.

Monkey EPO samples - monkey ③ is old monkey #3 which gives a titer of 1.34 mU/ml in genes assay. monkey #2 is from '183 and gives a titer of 20 mU/ml using anti-crude Ab. Monkey control is monkey #1 '183 which gives a titer of 50 mU/ml.

All monkey samples are negative w/ anti-peptide Ab.

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From Page No	Epo or expected EPO	PBSto. 17285A	CPM	CPM-pre	% inh	calculated EPO	EPO/ml
9 monkey @ 50λ		190	5824	5286	0	0	
20 25λ		215	5375	4837		0	
" 10λ		230	5650	5112		0	
2 monkey @ 50λ		190	6794	6256		0	
3 25λ		215	6270	5732		0	
" 10λ		230	6614	6076		0	
5 monkey-C 50λ		190	6092	5554		0	
" 25λ		215	7008	6470		0	
" 10λ		230	6100	5562		0	
x-therapeutics EPO							
10λ 1 U		230	1365	827	81.5	1.3 U	130 u/ml
20λ 1:10 200mU		220	1941	1403	68.6	510 mU	255 u/ml
10λ 1:10 100mU		230	2597	2059	53.9	190 mU	190 u/ml
0λ 1:10 50mU		235	2886	2348	47.4	110 mU	220 u/ml
2λ 1:10 20mU		238	4040	3502	21.6	18 mU	90 u/ml
Hankins 30u/ml							Am = 177 u/ml
10λ 1:10 300mU		140	3829	2091	53.2	180 mU	180 u/ml
30λ 1:10 150mU		190	3130	2592	41.9	72 mU	144 u/ml
25λ 1:10 75mU		215	3451	2913	34.7	44 mU	17.6 u/ml
Dukes 598 U/700λ							Am = 16.7 u/ml
20λ 1:100 171 mU		220	5360	5022	0	0	
10λ 1:100 85 mU		230	5782	5244	0	0	
5λ 1:100 42.7 mU		235	6203	5665	0	0	

Human EPO samples - E7 = 293 cell transfected w/ a 6 kb piece containing the sequence for human epo. M = mock transfection. Both samples were concentrated 7.1x before assaying. Previous results (book #340 - page 77) using α-crude Ab in a soln competition showed the titer of E7 to be 35.4 mU/ml (before concentration). The mock was negative. This assay gives us 22.3 mU/ml for E7 and 0 for the mock.

Monkey EPO samples = monkey 3 is old monkey #3 which gives a titer of 1.3 u/ml in genes assay. monkey #2 is from '83 and gives a titer of 500 mU/ml using anti-crude Ab. Monkey Control is monkey #1 '83 which gave a titer of 50 mU/ml. All monkey samples are negative w/ anti-peptide Ab.

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FRITSCHE v. LIN 102,097, 102,334
 INTER NOS 102,096 LIN DOC NO 101103

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Book No. _____ TITLE _____

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Sum Page No. _____

⇒ Therapeutics - Their titer is supposed to be 100 u/ml we get ~ 2x higher in the anti-crude assay and an avg of ~ 1.8x higher titer here using the anti-peptide.

⇒ Hankins material - a partially-purified mouse erythro leukemia supernatant from 483. The titer is 30 u/ml using the anti-crude. Its negative w/ the MCA in a plate binding. Our average value is 16.2 u/ml in the anti-peptide assay.

⇒ Duker's material - T-8-30 - fraction # 30 - peak from HA column ~ 50% pure. In vivo assay is 854 mu/ml. Anti-crude is ~ 1/3 that. Anti-peptide is negative.

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Project No. _____ Double Ab western - ① material from
 Book No. _____ TITLE Wart ② CHO cell material

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3/20/84 - We will be running two double Ab westerns. The first will be examining cell culture material from wart. We have a new preparation of his 1w32 line as well as a new line called 20 in 7.10 - 7/26/83 which is supposed to exhibit similar activity to the 1w32 line. The second western will be examining material from CHO cells (monkey ep)

Immunoprecipitation

- 1) Add 75% trisylol per tube.
- 2) Add sample, Ab- and T-12 Sepa (for ~ calculation of recovery) ^{30x = 31,504cpm}
- 3) Incubate overnight at 4°
- 4) Precipitate with 75% of 1:5 protein-A sepharose. Wash as usual
- 5) Count
- 6) Add 85% tube of 3x LSB. Boil ~5 min. Pellet. Remove supernatant. Calculate recovery.
- 7) Add dye & Ficoll. Prepare the remaining sample for the gel. Boil ~5 min
- 8) Electrophorese overnight at ~30V on a 12.5% polyacrylamide gel
- 9) Transfer to nitrocellulose at 60V for 6 hrs
- 10) Block w/ 10% horse serum add RT for ~ 1 1/2 hrs
- 11) Add mCA (protein A-sepharose #8) at 10 ug/ml in PBS + 5% horse serum w/ azide, trisylol and PMSF per usual. Incubate overnight
- 12) Wash 3x w/ PBS- pH 7.6
- 13) Add biotinylated horse anti mouse (1 drop/10 ml). Incubate at RT ~45 minutes.
- 14) Wash 3x w/ PBS- pH 7.6
- 15) Add ABC (2 drops/10 ml). Incubate 60 minutes at RT.
- 16) Wash 3x w/ PBS-
- 17) Add color development reagent.

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Results

Immunoprecipitation: from RIA done in book # 569 p 3/13-3/14/54

Sample	vol	CEP07	Ab	(2) T-125	cpm in pellet	% (1)	cpm in L58 sup	cpm in L58 pellet	% recovery	ng loaded on gel
2. 1W-32-Sx conc as usual - 1/2 the act. as usual (line 1)	3x1ml	233ng	3x1ml (9x total)	3x2 in 2 of the 3 tubes	8290	13.2 (37.6)	6150	1735	74.2	6.5 ng
3. 201ng - new line 2 - 200 ng 10x conc	1ml	300ng	9x	30x	5285	16.8 (47.9)	4575	750	86.6	12.4 ng
4. 1W-32 old batch	1ml	15.4ng	9x	30x	90	-	-	-	-	-
5. 201: 1W-32-10x	1.5ml	0	9x	30x	3870	12.3 (35)	3105	545	80.2	2
6. genes code epo	17x1 1:10	90ng	9x	30x	8055	25.6 (72.9)	6365	1100	79.0	23.0 ng
7. genes code epo	5.5x1 1:10	200ng	9x	30x	10525	33.4 (95.2)	8450	1255	80.6	15.3 ng
8. genes code epo	5.5x1 1:10	200ng	9x	30x	45	0	-	-	-	-
2 CHO-line 3 - 20 u/ml	0.2ml	25ng	6x	30x	11045	35.1 (100%)	8745	1790	79.2	22.2 ng
3 CHO-line 3 - 20 u/ml	0.2ml	54ng	8x	30x	9715	30.8 (97.8)	8190	1635	84.3	62.1
4 CHO-line 3 - 20 u/ml	0.2ml	186ng	10x	30x	6370	20.2 (67.6)	5720	950	89.8	101.3
5. control CHO	0.2ml	0	8x	-	-	-	-	-	-	-
6. CHO-line 3	0.2ml	186ng	8x	-	-	-	-	-	-	-

(1) The first # here is the % precipitation based on the 30,000 cpm in the starting material. The second number in parenthesis is the % precipitation using the 35.1% of the total in the first CHO sample as 100%. This number approximates the maximum precipitability of the system and is used to calculate the final recovery (i.e. ng loaded onto gel = % recovery in extraction x the % precipitation possible (if 35.1% of the total = 100%) x starting [cpm]).

(2) Ab: SC-204 + 3089 + b #3 - used undiluted in the amts mentioned. Comments - the % immunoprecipitability of the 1W-32 line is lower than usual - in fact the background is almost as high as the 'positives' Genes code epo precipitated as usual - as did the lysates of the CHO material - the larger amts of CHO material didn't compete and lower the % precipitability of the T-125 epas as well. As usual - however this had happened before.

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Date: FRITSCH v. LIN
INTER NOS 102,096
102,097, 102,334
LIN DOC NO L01106

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Book No. _____

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12.5% PAGE -

Gel 1

1. low mw prestained marker	15.1
2. 1u32 - line 1 new material	6.5ng
3. 201nn - new line 2	12.40g
4. 1u32 - old material + PI sera	0
5. 201-1u32 + Immure sera	0
6. genes crude epa	23
7. " "	153
8. " + PI sera	0
9. genes crude epa 17.1'10 - No Ab	40ng
10. " 8.5'1'10 No Ab	20ng

Gel 2

2. low mw prestained marker	15.1
3. CHO line 3	22.2ng
4. iCHO line 3	62.1ng
5. CHO line 3	106.3ng
6. control CHO	0
7. CHO line 3 + PI sera	0
8. Genes crude epa 1'10 No Ab	20ng

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Results-

Gel #1 - can see material which co-migrates with authentic epa in lanes 2 and 3 - although fainter than was hoped. However based on the arcs that were theoretically loaded on the gel - the intensity of the bands is close.

Gel #2 - can see material in lanes 2-4 - none in the control lanes 5 & 6. This specific material is a very broad band which seems to have its average migration distance slightly less than authentic epa indicating an average MW above the control epa.

From some rough MW determinations it appears that the range of the CHO cell material is 30,000 - 35,000 - with the mode of intensity at about 34,000.

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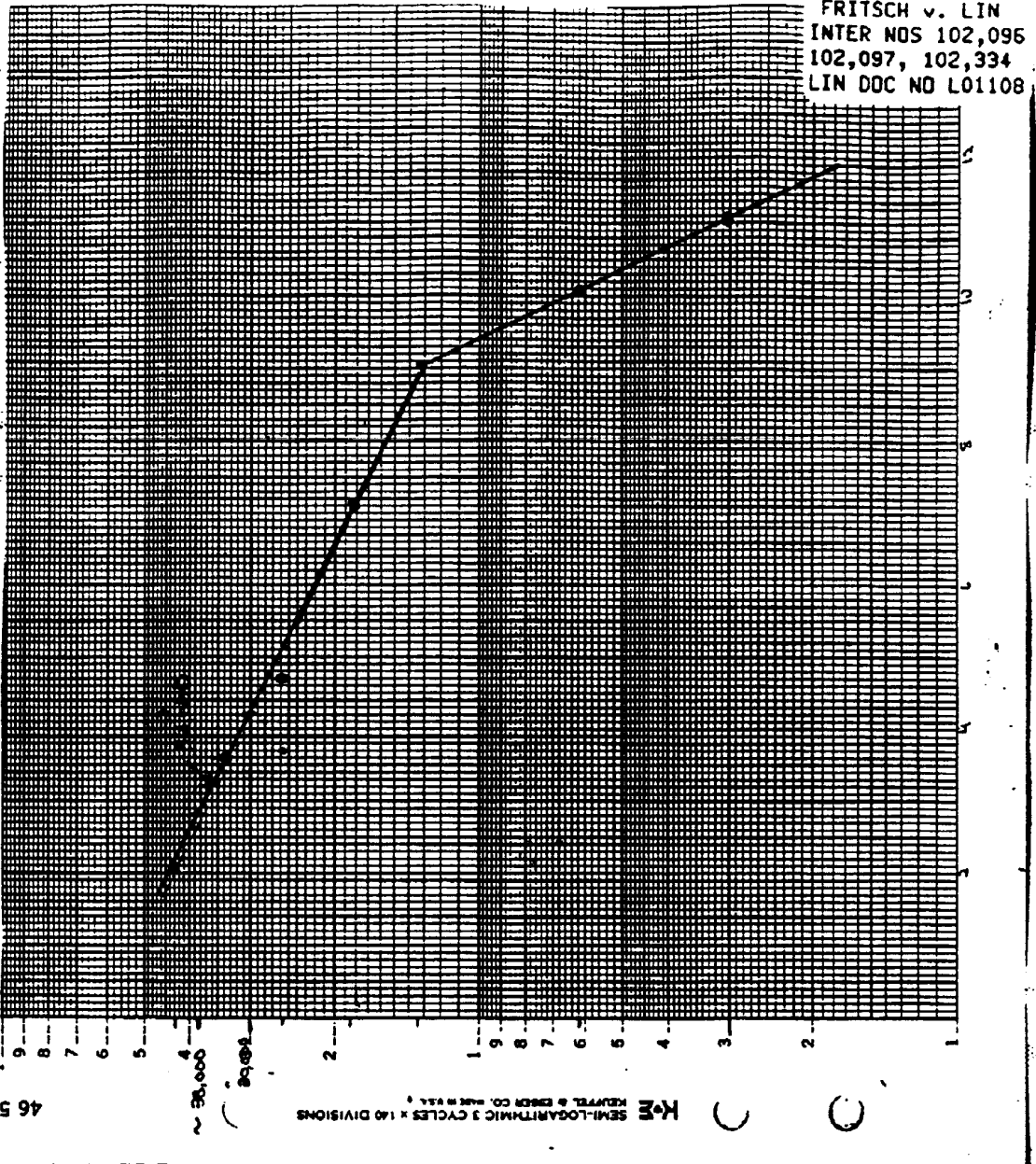
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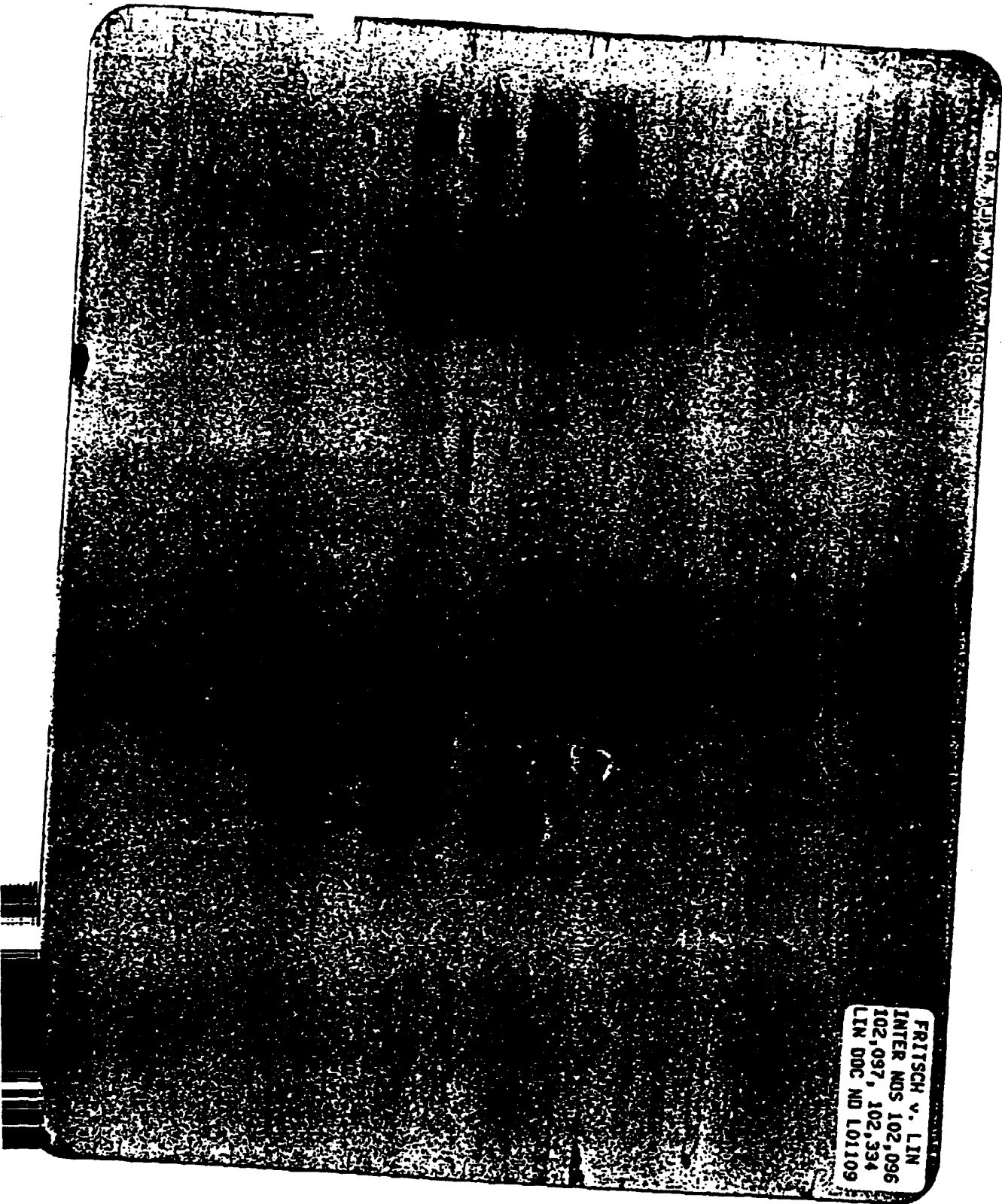
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migration - m.w. markers - 2.1cm, 4.7cm, 7.1cm, 9.05cm, 10.1cm, and 11.11cm. Authentic epa migrates with mode intensity at 3.6cm - The CHO cell material ranges from 2.5 - 4.2cm with the mode intensity at 3.2cm

Conclusion -

gel #1 - we are picking up material which co-migrates with authentic epa. from various cell lines. Although the intensity of the band isn't optimal - we will photograph this western since it shows reaction with positive - no reaction w/ negative

gel #2 - we are visualizing a broad specific band from the CHO cell material. This band is epa related since the Ab reaction is specific above background. The broad band ranges in m.w. from ~36,000 to ~38,000 with the mode of intensity at ~36,000. The band is quite continuous - no distinct bands within. In regard to the heterogeneity there are two possible explanations: A) carbohydrate heterogeneity - or B) protein heterogeneity - probably at the C-terminus - since the visualization is done with mAb. why the size includes material bigger than epa is also known - but the same 2 possibilities - may apply in conjunction - protein heterogeneity (maybe in processing) protein and carbohydrate will need to test this hypothesis.

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102,097, 102,334
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