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: «

(EGFR)

-2 (COX2)

»

6



2017

ΟΡΚΟΣ
ΙΠΠΟΚΡΑΤΟΥΣ



ὈΜΝΥΜΙ ΔΠΟΛΛΩΝΔ ΗΙΤΡΟΝ
ΚΑΙ ΥΓΙΕΙΩΝ ΚΑΙ ΠΑΝΑΚΕΙΩΝ ΚΑΙ
ΘΕΟΥΣ ΠΑΝΤΑΣ ΤΕ ΚΑΙ ΠΑΣΑΣΙΣΤΟΡΑΣ
ΠΟΙΕΥΜΕΝΟΣ ΕΠΙΤΕΛΕΑ ΠΟΙΗΣΙΝΚΑΤΑ
ΔΥΝΑΜΙΝ ΚΑΙ ΚΡΙΣΙΝ ΕΜΗΝ ΟΡΚΩΝ
ΤΩΝΔΕ ΚΑΙ ΖΥΓΓΡΑΦΗΝ ΤΗΝΔΕ ΗΤΗΣ
ΔΙΦΘΑΙΜΕΝ ΤΟΝΔΙΔΑΧΗΝΤΑ ΜΕ ΤΗΝ
ΤΕΧΝΗΝ ΤΑΥΤΗΝ ΙΣΑ ΓΕΝΕΤΗΣΙΝ ΕΜΟΙΣΙ
ΚΑΙ ΒΙΟΥ ΚΟΙΝΩΣΑΣ ΘΑΙ ΚΑΙ ΧΡΕΩΝ
ΧΡΗΣΙΖΟΝΤΙ ΜΕΤΑΔΟΣΙΝ ΠΟΙΗΣΑΣ ΘΑΙ
ΚΑΙ ΓΕΝΟΣ ΤΟ ΕΣ ΟΥΤΕΩΝ ΑΔΕΛΦΟΙΣ

ΙΣΩΝ ΕΠΙΚΡΙΝΕΣΙΝ ΑΡΡΗΣΙ ΚΑΙ ΔΙΔΑΧΕΙΝ ΤΗΝ ΤΕΧΝΗΝ ΤΑΥΤΗΝ
ΗΝ ΧΡΗΣΙΩΣΙ ΜΑΝΘΑΝΕΙΝ ΑΝΕΥ ΜΙΣΘΟΥ ΚΑΙ ΖΥΓΓΡΑΦΗΣ
ΠΑΡΑΓΓΕΛΙΑΣ ΤΕ ΚΑΙ ΑΚΡΟΗΣΙΟΣ ΚΑΙ ΤΗΣ ΛΟΙΠΗΣ ΑΠΑΣΗΣ
ΜΑΘΗΣΙΟΣ ΜΕΤΑΔΟΣΙΝ ΠΟΙΗΣΑΣ ΘΑΙ ΝΙΟΙΣΙ ΤΕ ΕΜΟΙΣΙ
ΚΑΙ ΤΟΙΣΙ ΤΟΥΕΜΕ ΔΙΔΑΧΑΝΤΟΣ ΚΑΙ ΜΑΘΗΤΑΙΣΙ ΣΥΓΓΕ
ΓΡΑΜΕΝΟΙΣΙ ΤΕ ΚΑΙ ΑΡΚΙΣΜΕΝΟΙΣ ΝΟΜΩ ΙΗΤΡΙΚΟΑΛΛΟ
ΔΕΟΥΔΕΝΙ ΔΙΑΙΤΗ ΜΑΣΙ ΤΕ ΧΡΗΣΙΩΜΑΙ ΕΠ ΟΦΕΛΕΙΗ ΚΑΝΟΝΩΝ
ΚΑΤΑ ΔΥΝΑΜΙΝ ΚΑΙ ΚΡΙΣΙΝ ΕΜΗΝ ΕΠΙΔΗΑΣΕΙ ΔΕ ΚΑΙ ΔΑΙ
ΚΙΗ ΚΙΡΖΕΙΝ ΟΥΔ ΔΩΣΩ ΔΕ ΟΥΔΕ ΒΟΗΓΗΣΟΜΑΙ ΣΥΜΒΟΥΛΙΗΝ
ΤΟΙΗΝΔΕ ΟΜΟΙΟΣ ΔΕ ΟΥΔΕ ΓΥΝΑΙΚΙ ΠΕΣΣΟΝ ΘΕΟΡΙΟΝ ΔΩΣΩ
ΑΓΝΟΣ ΔΕ ΚΑΙ ΟΣΙΟΣ ΔΙΑΤΗΡΗΣΩ ΒΙΟΝ ΤΩΝ ΕΜΩΝ ΚΑΙ ΤΕΧΝΗΝ
ΤΗΝ ΕΜΗΝΟΥΤΕΜΕΩ ΔΕ ΟΥΔΕ ΜΗΝ ΑΙΘΙΟΝΤΑΣ ΕΚΧΩΡΗΣΩ ΔΕ
ΕΡΓΑ ΤΗΣ ΑΝΔΡΑΣΙ ΠΡΗΣΙΟΣ ΤΗΣ ΔΕ ΕΣ ΟΙΚΙΑΣ ΔΕ ΟΚΟΣΑΣ ΑΝ
ΕΣΩ ΕΣΕΛΕΥΣΟΜΑΙ ΕΠ ΟΦΕΛΕΙΗ ΚΑΜΝΩΝΤΩΝ ΕΚΤΟΣ ΕΩΝ ΠΑΣΗΣ
ΑΔΙΚΗΣ ΕΚΟΥΣΙΗΣ ΚΑΙ ΘΕΟΡΙΗΣ ΤΗΣ ΤΕ ΑΛΛΗΣ ΚΑΙ ΑΦΡΟΔΙΣΙ
ΩΝ ΕΡΓΩΝ ΕΠΙΤΕ ΓΥΝΑΙΚΕΙΩΝ ΣΩΜΑΤΩΝ ΚΑΙ ΑΝΔΡΩΝ ΕΛΕΥ
ΘΕΡΩΝΤΕ ΚΑΙ ΔΟΥΛΩΝ Δ Δ ΑΝ ΕΝ ΘΕΡΑΠΗΗ ΚΑΙ ΙΔΩ Η Δ
ΚΟΥΣΩ Η ΚΑΙ ΑΝΕΥ ΘΕΡΑΠΗΗΣ ΚΑΤΑ ΒΙΟΝ ΑΝΘΡΩΠΩΝ ΑΜΗ
ΧΡΗ ΠΟΤΕ ΕΚΑΔΑΕΕΣΘΑΙ ΕΞΩΣΙΓΗΣΟΜΑΙ ΑΡΡΗΤΑ ΗΓΕΝΥΜΕΝΟΣ
ΕΙΝΔΙ ΤΑ ΤΟΙΑΥΤΑ ΟΡΚΩΝ ΜΕΝ ΟΥΝ ΜΟΙ ΤΟΥΔΕ ΕΠΙΤΕΛΕΑ
ΠΟΙΩΝΤΙ ΚΑΙ ΜΗ ΖΥΓΧΕΟΝΤΙ ΕΙΗ ΕΠΑΥΡΑΣΘΑΙ ΚΑΙ ΒΙΟΥ
ΚΑΙ ΤΕΧΝΗΣ ΔΟΧΑΖΟΜΕΝΩ ΠΑΡΑ ΠΑΣΙΝ ΑΝΘΡΩΠΟΙΣ ΕΙΣ ΤΩΝ
ΔΙΕΙ ΧΡΟΝΩΝ ΠΑΡΑ ΒΑΙΝΩΝΤΙ ΔΕ ΚΑΚΕΠΙΟΡΚΟΥΝΤΙ ΤΑΝΑΝΤΙΑ
ΤΟΥΤΕΩΝ

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:
/ : 28/05/1975
. :
: 6977888775
E-mail: mzktann@yahoo.gr

2001

2001 (8,4:)
: << **Crohn**
K >> (: 10)

7/2014 : **ó**

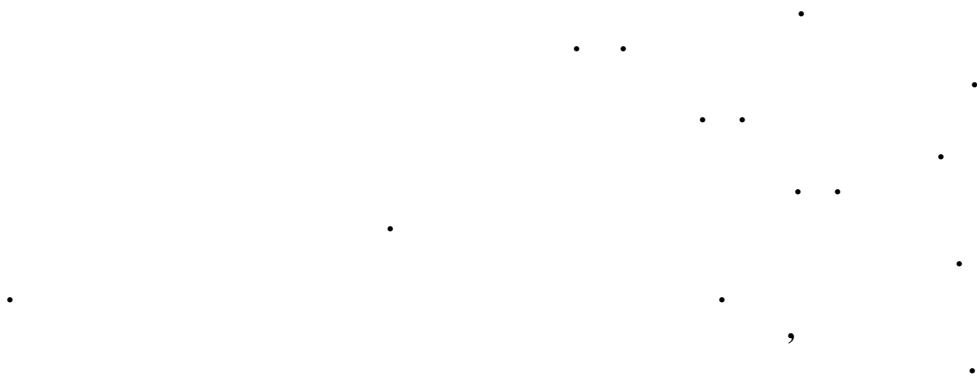
10/2011 **9/2012** **6/2014:**
8/2012:

9/2011:
10/2007 **2010:**
3/2007 **9/2007:**

2004-2007:
2003-2004:
2002-2003: -

- 1.**
- 2.**

2006 **2006:** / **(ECDL Core)**
:



202, (.2 .5343/1932) ,

, 875.000

20

44,5/100.000

, 35,6/100.000

5,7/100.000

2,1/100.000

5

()

Crohn, to 2-3%

50

, / , .
(APC),
,
-Cadherin .
, - c-myc ras,
17 p53,

.
DNA,
(MSI)
(MSI-).
, MSI-
DNA KRAS ABRAF.
/

1.

:

()

APC, c-myc ras,
p53, p16, 3p.

2%

(80%) (10%).

<1 1%, 4%

10% 1 2

>2

2. (Serrated)

, ,
,
, DNA. ,
, 1,3-11% .
4-37%.

3.

, 28%
DALM.
,
,
,

WHO (2010),

Paneth,

MSI

MSI,

(3) (1) (2), (4).

Dukes
 (), () ().
 , (),
 . , :
 (), (), ()
 IV ().
 ➤ / : T (),
 (), (V)
 (R).
 ➤ / :
 (grade), (/CRM)
 (pTNM).
 ➤ / :
 MSI,
 , MSI
 .
 ➤ / : DNA
 , / 18q/DCC.
 ➤ **IV**
 (IV) : ,
 , ,
 , NORs
 .
 ,
 ,
 ,

Astler-Coller

(1954)

		5
	$\frac{1}{4}$	100,00%
B1	.	(-) 67%
B2	.	(-) 54%
C1	.	(+) 43%
C2	.	(+) 23%
D		-

Dukeø (1932)

A	O
B	
C	

C

C1	
C2	

	Tis, No, Mo
	T1, No, Mo 2
	3, , 4, ,
	(), 1, () 2
V	() () 1
Tis	In situ
T1	
T2	
T3	-
4	
No	
N1	1-3
N2	>4
Mo	
M1	

,

(8,7 100.000).

,

6%

,

.

1.
2.
3.
4.
5.
6.
7.

- 1. KRAS
- 2. DNA
- 3. APC ()
- 4. DCC 18
- 5. p53 17

30%
 ~5%
 (<2% <1,5 , >10% ×2,5
) 65%

(30% >30).

>5 mm.

(FAP)

APC

().

5bpAAAGA 3927-3931 10%

APC

1 10.000

APC

1249-1330

(5.000).

(1.000).

(78-163) 5ø

FAP (

, 100,).

(. .
100% 40 ,
, Vater,
. .) (. .
10 ,
,
(), , .).

FAP

- 1. :
- 2. 10-12 . 1-
- 2 35.

(**Lynchø, HNPCC**)

Lynchø

1913 Warthin 15%

Lynchø ,

HT et al, 1971 & Boland et al, 1984,

80%.

Lynch

Lynch

(

44)

Lynch ,



()



35 .

HNPCC

HNPCC 3-6%

12-15%

(4-26%)

hMSH2 6-10%

hMSH2

hMLH1.

hMSH2 hMLH1

16

73 kb hMSH2

16

58 kb hMLH1.

DNA

1:200.

(60%).

1.

1-3 ,

20-25 .

2.

25-35

3.

4.

25

5.

6.

35

(,)

DPCA4/SMAD4

(TGF-). SMAD4

TGF-

Peutz-Jeghers

(95% 25).

LKBI (STK11)

19.

(50% , 40% , 30% 20%

)

PJS

(,),

PJS

PJS, Tomlinson & Houlston (1997)

PJS:

1. 2 , 10,

1 mm.

2. 3 , 25 (1 mm.

3. 10 ,

4. 25 ,

35 .

5. 25 .

6. .

20 .

35

30

•
_____ :
1. (,
)

2.
3.
4.
_____, _____ :
1.
2. .
3.
4.

•
•
_____ ,
_____ . 15%
_____ , _____ .

•
1. _____ :
_____ (_____).
_____,
_____ (APC) _____ 5;21.
_____ 100%,
40 _____ .
_____ ,
_____ .

2. **Gardner:**
(, , , ,
).
.

3. **Turcot:**
.

4. **Lynch):** (HNPCC,
80%
(
, , , ,).

DNA (MLH1, MSH2, MSH6
PMS2).

5. :
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, , .
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6. **Peutz-Jeghars:**
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Crohn:

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50

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1. : ×50
2. : , ,
<60 ,
3. :
50 ,
HNPCC,

1. :
2. , , ..
3. .
400 mg

(sulindac)

-2 (COX-2)

COX-2

PreSAP (Prevention of Colorectal Sporadic

selecoxib

Adenomatous Polyps)

3

selecoxib

3

COX-2

➤

90%

➤

Faecal Occult Blood Test-FOBT).

FOBT

NSAID,

6%

(60), $2/3$

(2 10.000).

()

1.

:

,

.

75%

,

.

2.

:

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.

American Cancer Society

50

:

- 10 .
- (FOBT test)
- 5).
- () 5 .

:

1

<1 mm	3-6		,
	3		, 5 . ,
Ca			3 , 5 .
Ca	40 10		5
Crohn ()	8		1-2

:

2

			,
(HNPCC)	21		, 1-2 40,

(NPCC).

DNA (mismatch repair genes, MMR).

Amsterdam (1991),
Bethesda,

1997

«

HNPCC

»

:

3

Familiar adenomatous polyposis	5q21	APC
Juvenile polyposis	10q22 10q23 18q21	BMPR1 PTEN SMAD4
Cowden syndrome	10q23	PTEN
Peutz-Jeghers syndrome	19p13	STK11
Hereditary mixed polyposis	6q	unknown
Hereditary nonpolyposis colon cancer	2p22 3p21 2q31 7p22 14q24	MSH2 MLH1 PMS1 PMS2 MLH3 (15)

, APC, SH2, MLH1.

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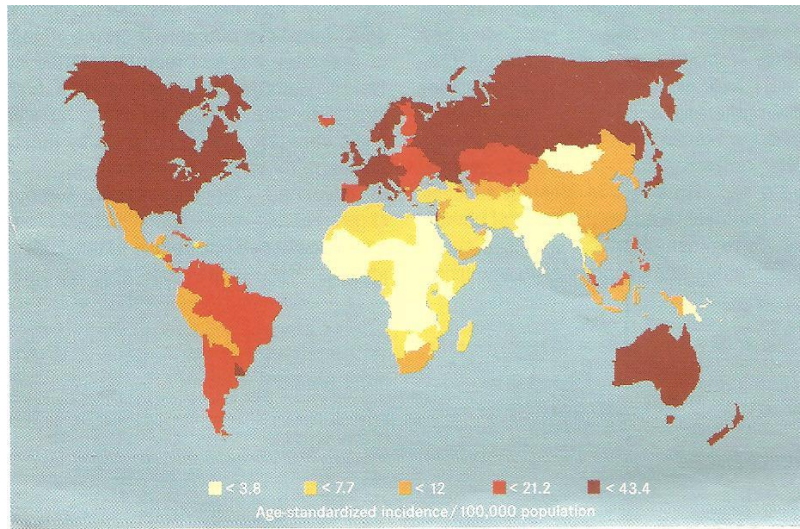
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. . . ,

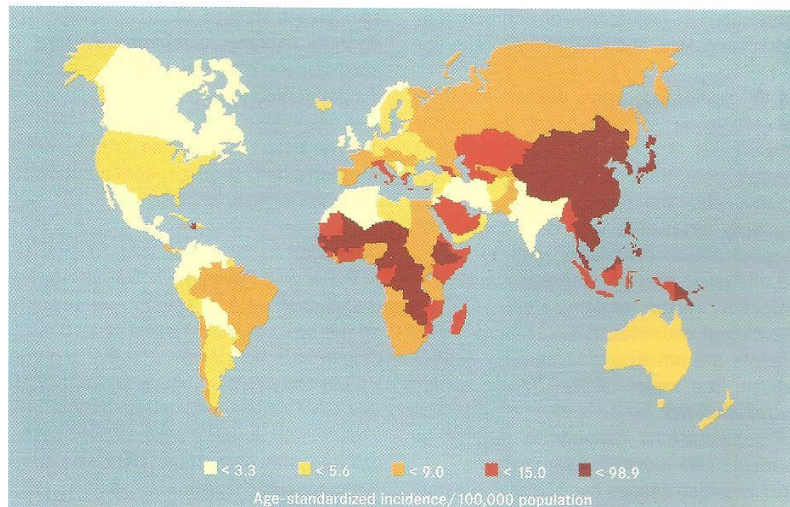
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2. (- ,)



3.

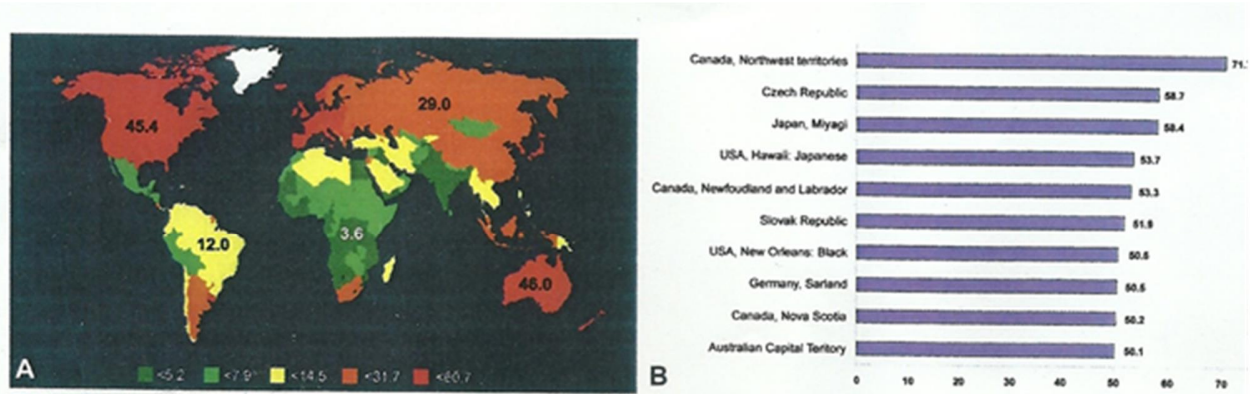
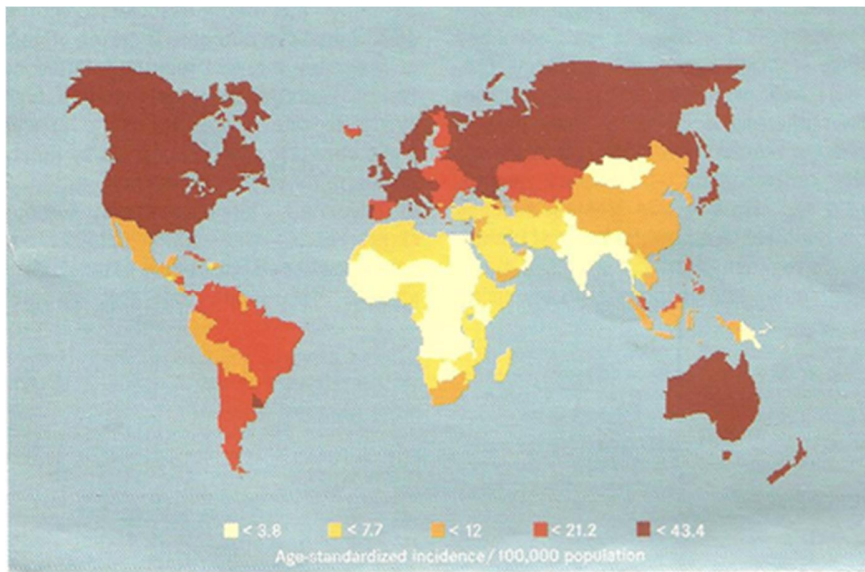


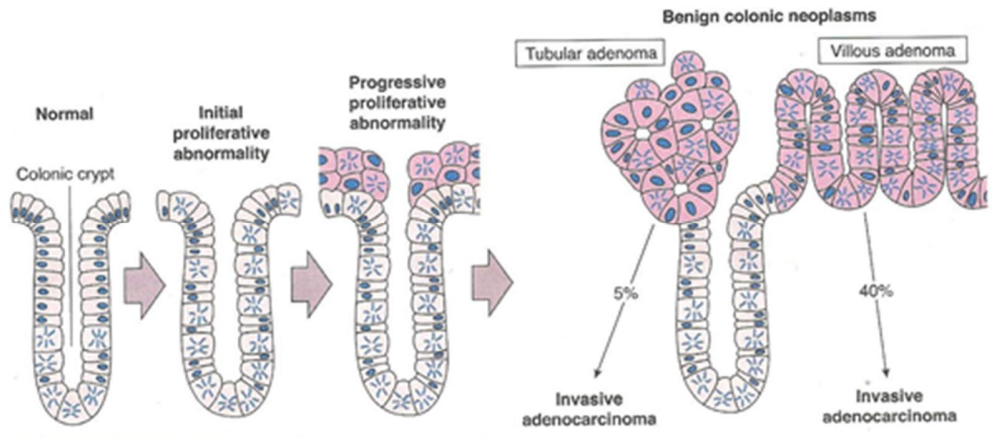
Fig. 8.01 **A** Worldwide annual incidence (per 100 000) of cancer of the colon and rectum in men (842A). Numbers on the map indicate regional average values and are higher in developed countries (842A). **B** Age-standardized incidence (per 100 000) of colorectal cancer in men in selected countries (842).

1.



2.

(- ,)



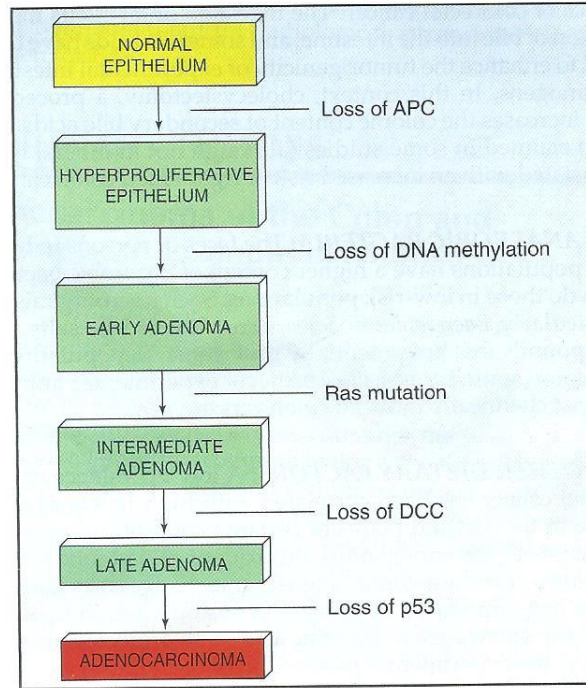
3.

ó

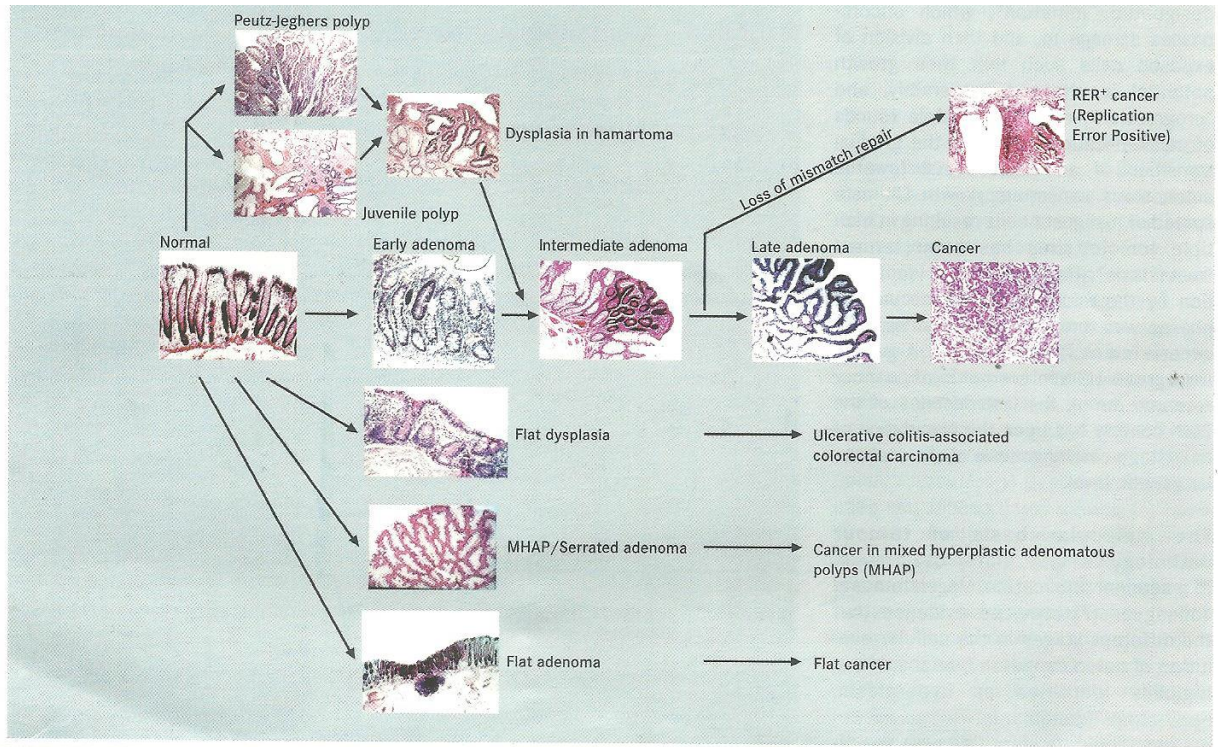
)ó

(

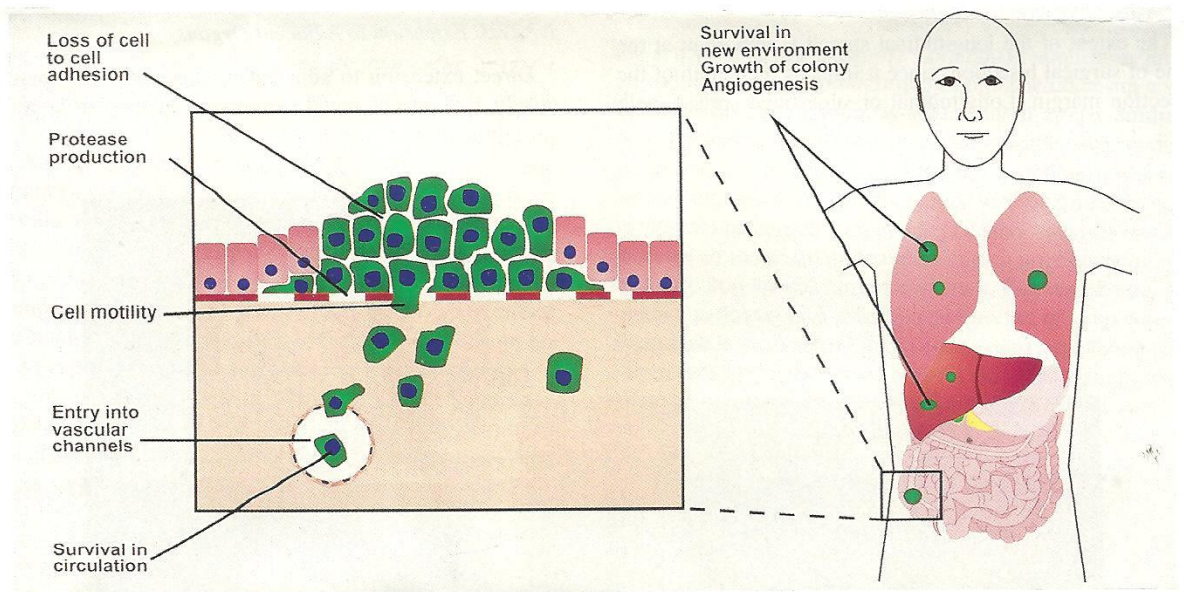
ó



3 .

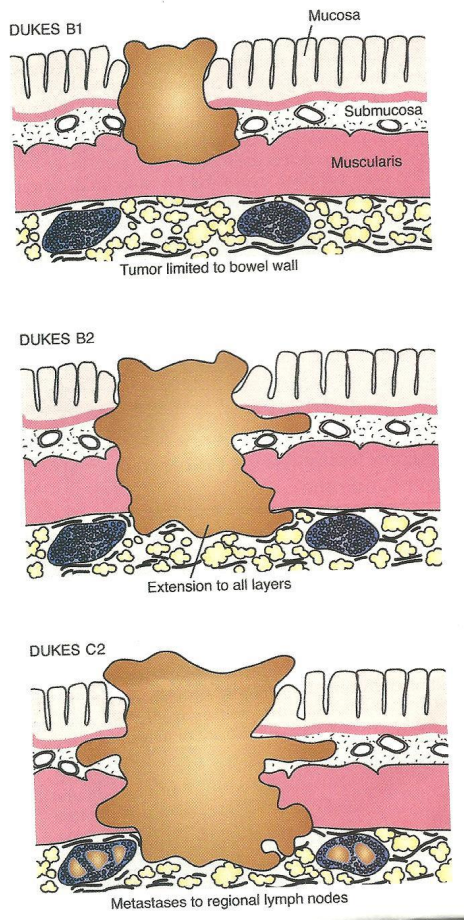


4.

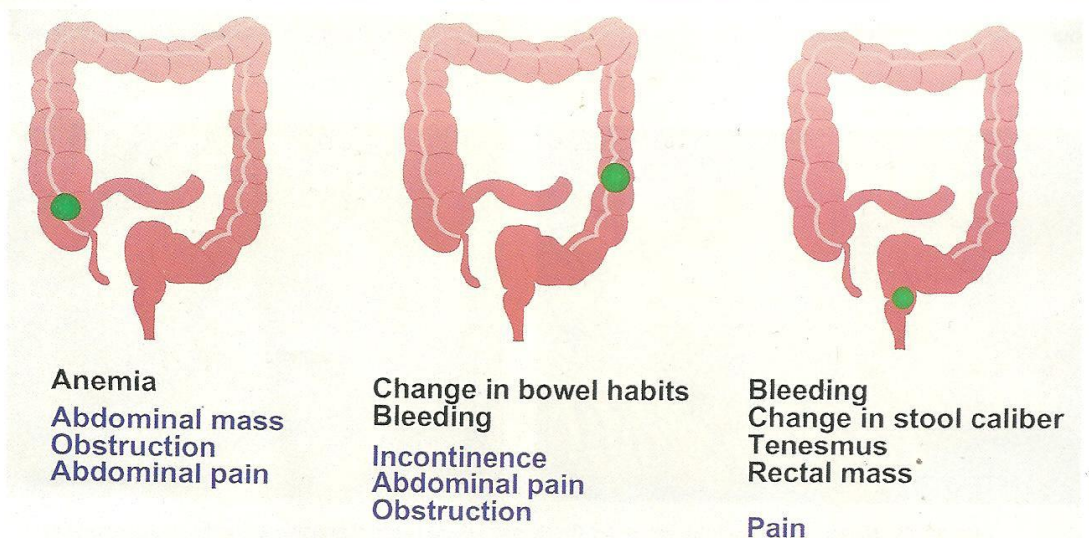


5.

6



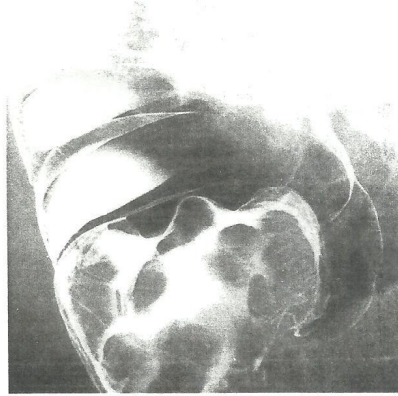
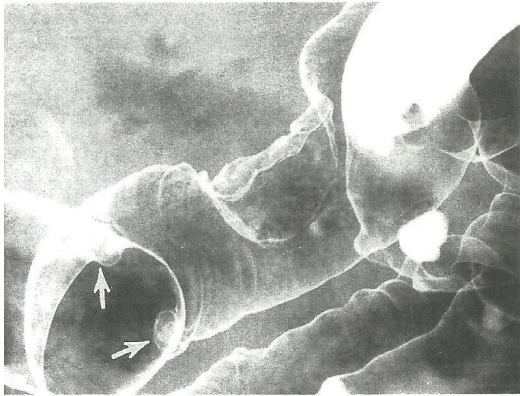
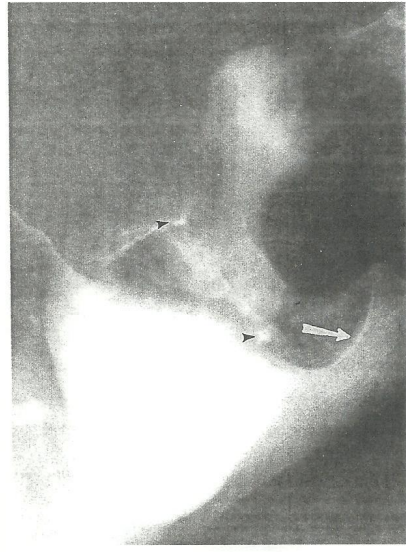
6. Dukes



7.



A



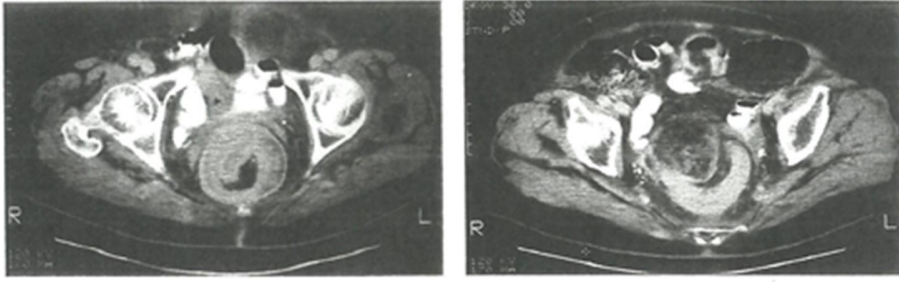
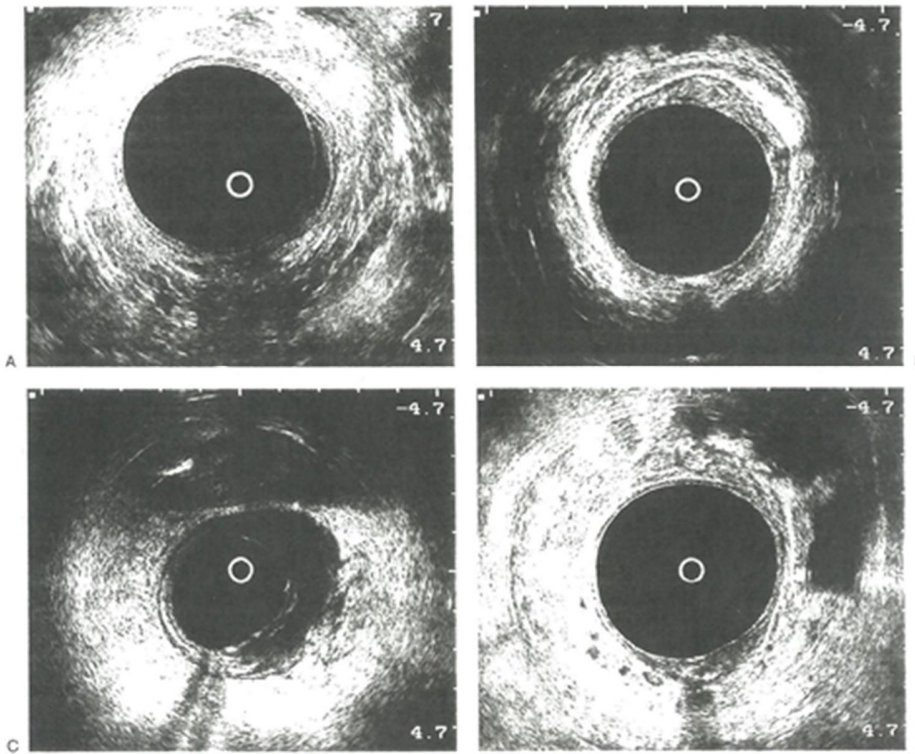


FIG. 20.140. Staging of rectal carcinoma. A: Computed tomography scan demonstrates a rectal carcinoma that has produced an intussusception. B: At another level, a mass is present that infiltrates into the perirectal soft tissue.



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25-30%

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75%.

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5-10%.

microRNA.

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microDNA

ö). - (-

, « »

(signal

100

➤ () :

sis.

(platelet-derived growth factor-PDGF).

➤ :

2

erb 2 erb B-2.

[epidermal growth factor receptor 2 (HER-2)].

HER-2



:

abl ras. abl

(. . Imatinib).

ras



:

DNA

RNA

myc.

myc

myc

t(8,14)

(q24, q32)

Burkitt



:

().

bcl-2

30
p53, BRCA1, BRCA2, APC RB1.

DNA

(), ()

(RBI)

RBI

RBI

RBI

DNA:

DNA.

DNA

DNA,

DNA

DNA,

p53

p53

DNA

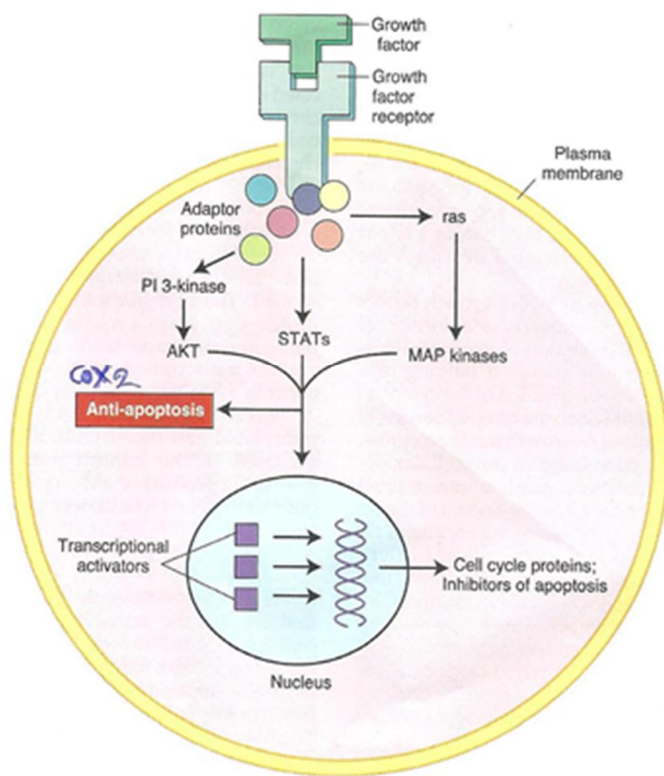
p53

Li-Fraumeni.

LFS

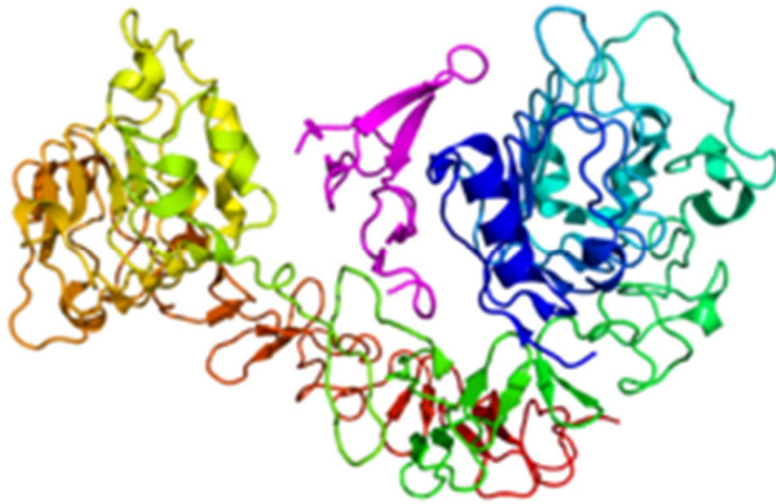
(

The Molecular Genetics of Cancer



COX2

-



EGFR

170- Da,

tyrosine kinase receptors) EGFR (ErbB
 ErbB3-Her3, ErbB4-Her4), (ErbB1, ErbB2- HER2/neu,
 c-erb B.

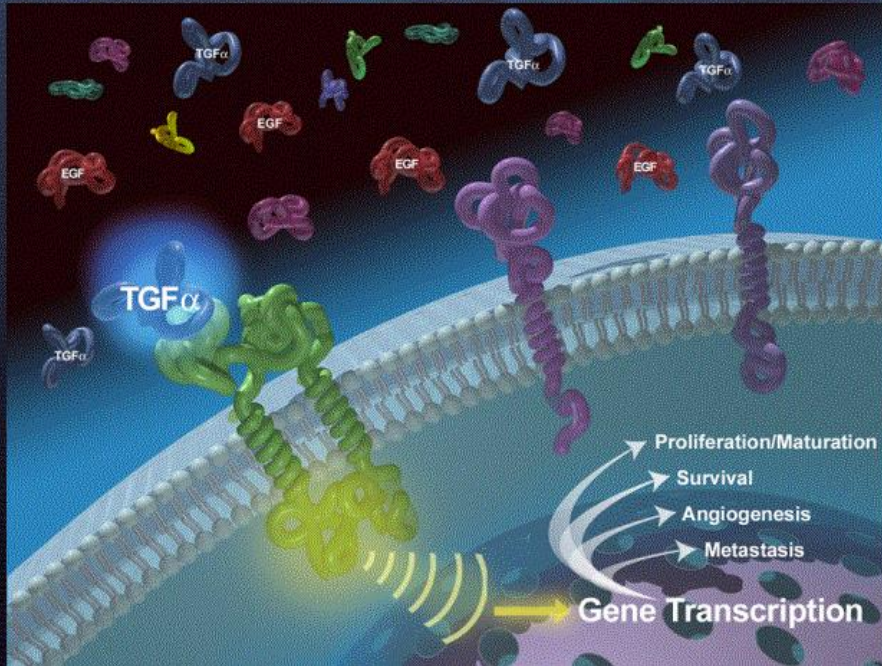
EGFR

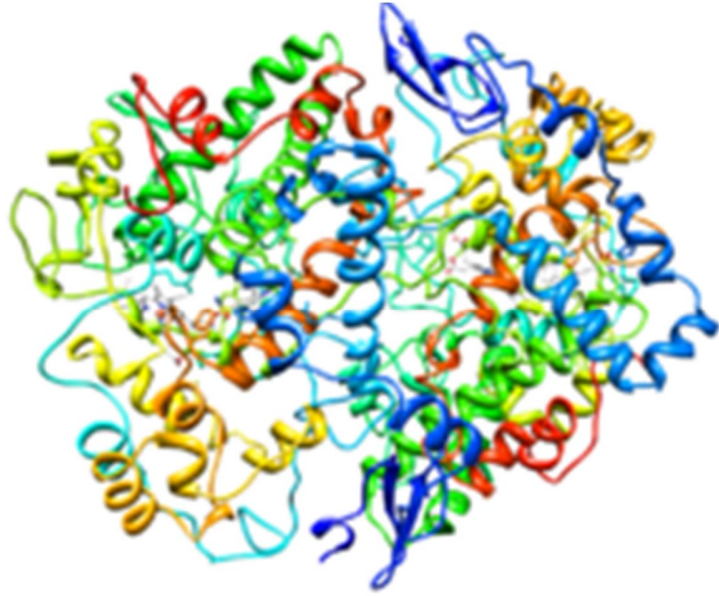
EGFR

2 (COX2)

2

EGFR Signaling Pathway: TGF- α Activation





COX-2

- (COX)

.H COX

COX1

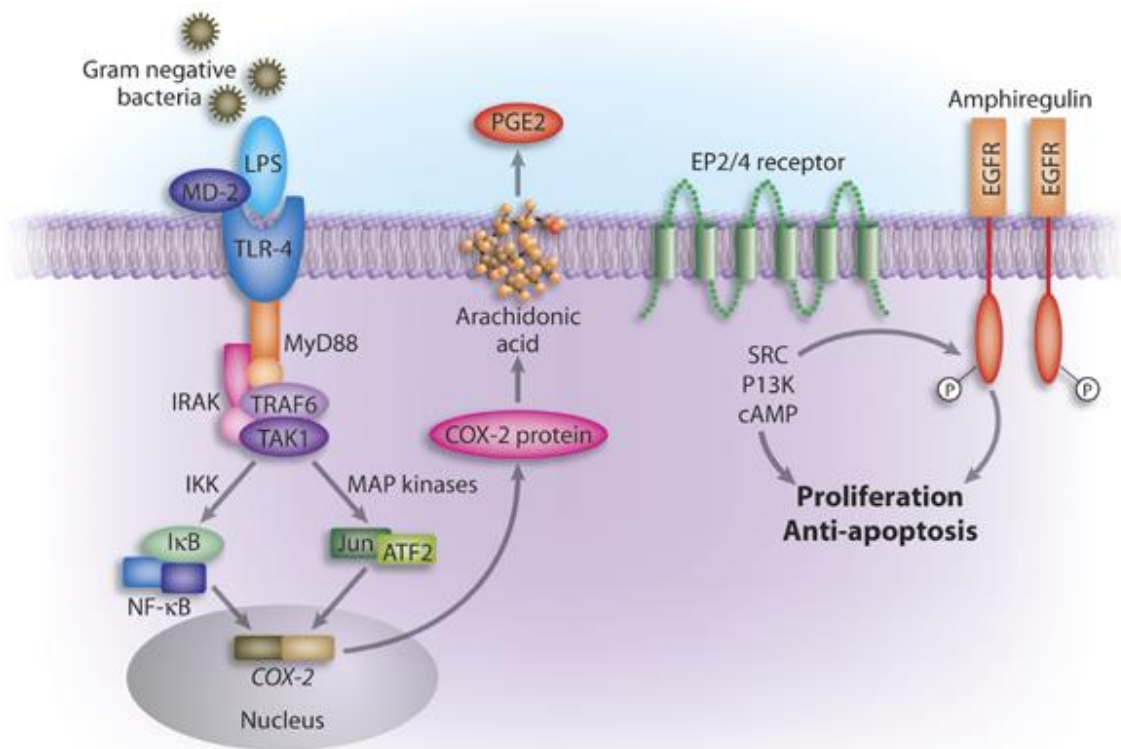
COX2

- . COX2

, , ,

,

COX2



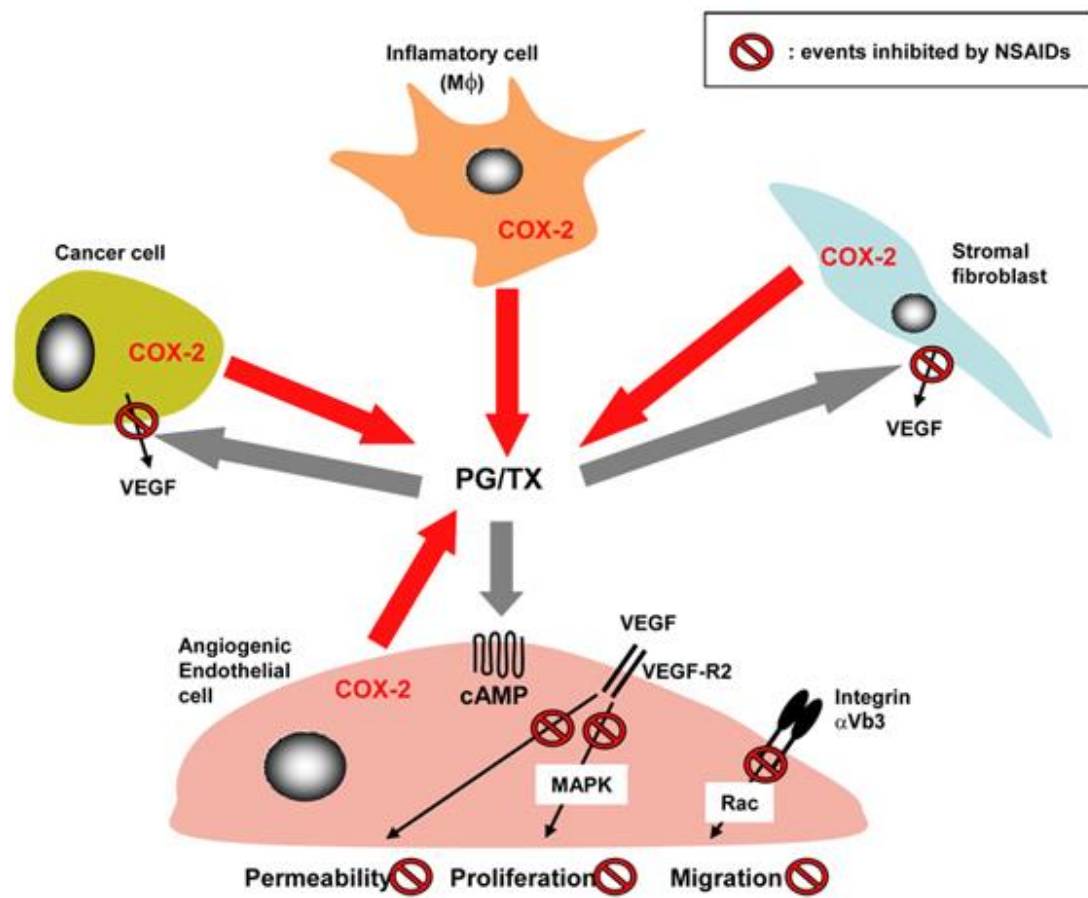
COX2

()

PGE2

«

»



COX

NSAIDs

I IK

100

∅ ()
2007.

Zeiss.

:

- 1.
2. - 15 .
- 3.
4. (3 10).
(3 5).
5. 10 20 .
6. 10 .
7. (2 2).
8. (2 3).
9. (2 2).
10. Zeiss.

∅

(.

).

1. . grade I ()
 . grade I ()
 . grade ()
2. Dukes EGFR
 COX-2 Ventana.
 Anti EGFR (clone 31G7, Zymed USA), anti-COX-2 (N-20, Somfa Gyz USA)
- (-), <10%
 (+) >10% (++)
 >30%
1. 5 60 15
 2. (10 5
) 100 , 96 , 80 70 ,
3. TBS buffer (pH: 7,6), 5
 4. 8
 2 2 3%,
5. TBS buffer (pH: 7,6), 5
 6. buffer (pH: 6),
 , buffer 5
 7. ,
 45 ,
 8. TBS buffer (pH: 7,6), 5
 9. , (
- 1.000 1 TBS 10 1) 30

10. TBS buffer (pH: 7,6), 5 .
11. streptABCComplex/HRP, Duet (DAKO),
30 (1.000 1 TBS + 10 1 A +10 1 B), .
15 .
12. TBS buffer (pH: 7,6), 5 .
13. DAB (100 1 Tris ó buffer + 2 app. DAB + 100 1 2 2
3%) 10 .
14. .
15. - 1 ,
70 , 80 , 96
100 , .

1-6:

1 -

:

/						DUKES	EGFR	COX-2
1		165/07	49		3		(+)	(+)
2		352/07	48		3,5		(+)	(+)
3		554-57/07	52		4		(-)	(+)
4		763-64/07	56		4		(+)	(+)
5		979/07	60		6		(-)	(+)
6		1078/07	51		5		(-)	(+)
7		1149-50/07	72		6		(+)	(-)
8		1226-27/07	69		7		(-)	(-)
9		1507/07, 1714/07	70		8		(-)	(+)
10		1554/07	65		8		(-)	(-)
11		1893/07	63		4,5		(-)	(-)
12		1899/07	60		5		(-)	(+)
13		1903-04/07	64		5		(-)	(+)
14		2147-48/07	62		3,5		(-)	(-)
15		2305/07	63		3		(+)	(+)
16		2470-71/07	67		6		(-)	(-)
17		2508-09/07	61		6,5		(-)	(-)
18		2674/07	59		7		(+)	(+)
19		2854/07	69		7		(+)	(+)
20		2859/07	70		7		(-)	(-)
21		2876/07	66		6		(-)	(-)
22		2918/07	49		4,5		(-)	(-)
23		3208/07	53		4		(+)	(+)
24		3260-62/07	54		5		(-)	(+)
25		3842-43/07	50		6		(-)	(+)

=

=

=

:

: 14 (56%) 11 (34%).

: 49 70 (), 59,5 , 43 72 (),
57,5.

: 4 (16%),
10 (40%) 11 (44%).

: 3 8 .(5,5).

: Dukes 15 (60%). Dukes 10 (40%).

EGFR: 17 (68%) (-). (8) (32%) .

COX-2: 10 (40%) (-) 15 (60%) (+).
36-60

.

:

EGFR: 17 (68%) (-). (8) (32%) .

COX-2: 10 (40%) (-) 15 (60%) (+).
36-60

.

2 -

:

/						DUKES	EGFR	COX-2
1		56/07	50		3		(+)	(++)
2		110-12/07	52		3,5		(-)	(+)
3		171/07	60		4	C, N1	(-)	(+)
4		391-92/07	80	A	5	C	(-)	(+)
5		435/07	69	A	6	B	(+)	(++)
6		438/07	67		4,5		(+)	(++)
7		462/07	65		3,5	C, N1	(-)	(+)
8		493/07	70		4	C, N1	(+)	(-)
9		498/07	72		4,5	C	(+)	(++)
10		501-03/07	69		7	C	(-)	(+)
11		571-72/07	68		6,5	C	(-)	(+)
12		708-09/07	80		3,5	C, N2, M	(+)	(++)
13		873/07	75		3	C	(+)	(+)
14		1017/07	73		4,5	C	(+)	(++)
15		1053/07	72		5	C	(+)	(+)
16		1086-87/07	49		5,5	C	(+)	(++)
17		1153-54/07	60	A	3	B	(+)	(++)
18		1271/07	45		6,5	B	(-)	(+)
19		1320/07	45		8	B	(-)	(+)
20		1388-90/07	42	A	7,5	B	(-)	(+)
21		1396/07	66	A	4	B	(+)	(+)
22		1577-78/07	72	A	3,5	C, N1, M	(+)	(++)
23		1582/07	79	A	3,5	B	(-)	(+)
24		1707/07	75		5,5		(-)	(+)
25		1776-77/07	63		6,5	C, N1	(+0)	(++)
26		1807-08/07	67	A	8	B	(-)	(+)
27		1819/07	72		8	B	(+)	(++)
28		1971/07, 3316/07, 3341-43/07	77		6	C, N1	(-)	(++)
29		2010/07	79	A	4	C, N2	(+)	(+)
30		2012-13/07	80		4	C,	(-)	(++)
31		2015/07	79		3,5		(+)	(++)
32		2018-19/07	57		5		(-)	(+)
33		2075/07	58		6,5		(-)	(+)
34		2280-82/07	55		7		(-)	(+)
35		2331-33/07	59		8		(+)	(+)
36		2380/07	69		5	C	(+)	(+)
37		2425/07	80		5		(-)	(++)
38		2427-28/07	79		6,5		(+)	(+)
39		2444-45/07	78		5		(+)	(+)

40	2450-51/07, 2497/07	79		5	C, N1,	(-)	(++)
41	2475-76/07	75		3	B	(+)	(++)
42	2535-38/07	64	A	3	B	(-)	(+)
43	2622/07	66		7,5	B	(-)	(+)
44	2626/07	72		6,5	B	(+)	(+)
45	2646/07	70		6	B	(-)	(+)
46	2662/07, 2687/07	70		5	C, N2, M	(+)	(++)
47	2697/07	70		4,5		(+)	(+)
48	2791-92/07	65		3,5		(+)	(+)
49	2816/07	66		4,5		(-)	(+)
50	2988/07	69		8		(-)	(+)
51	2999/07	79		3		(+)	(-)
52	3039/07, 3158-59/07	80		3,5		(-)	(-)
53	3277-78/07	44		4		(-)	(+)
54	3294-95/07	49		5		(-)	(-)
55	3360-62/07	50		6		(-)	(+)
56	3390/07	55		3	C, N2, M	(+)	(++)
57	3499/07	68		7,5	B	(-)	(-)
58	3551-52/07	69	A	8	B	(-)	(-)
59	3890/07	66		7		(+)	(+)
60	3892/07	67		5		(+)	(+)
61	3902-03/07	77		4,5		(+)	(+)
62	3905/07	73		3	C, N2, M	(+)	(++)

N1 = 1-2
2 =>4
=

:

: 32 (%) 30 (%).

: 45 80 (), , 42 80 (),

: 9 (%),
22 (%) 31 (%).

: 3 8 . (5,5).
 : Dukes - 40 (%) Dukes C ó 22 ((%). 7 1-3 . 5
 >4 .

EGFR: 30 (49,3%) (-). 32 (51,7%) (+).
COX-2: 4 (6,4%) (-), 35 (56,4%) (+) 23
 (++) (37,2%).

3 -

:

/						DUKES	EGFR	COX-2
1		19/07	65		4	C, N2, M	(+)	(++)
2		109/07	70		4,5		(+)	(++)
3		428/07	63		7		(-)	(+)
4		527-29/07	65		7		(+)	(++)
5		576-77/07	55		6,5	C, N2	(+)	(++)
6		639/07	49	A	6,5	B	(+)	(++)
7		1308/07	48		5,5		(-)	(+)
8		2494/07, 3539/07	59		6,5		(+)	(++)
9		2767/07	58		6	C, N1, M	(-)	(+)
10		2789/07	62		6	C, N2	(+)	(++)
11		3005-06/07	66		7,5		(-)	(+)
12		3183-85/07	68		7	C, N1, M	(+)	(++)
13		3651/07, 3690-91/07	70		7	C, N2	(-)	(+)

:

: 7 (53,8%) 6 (46,2%).

: 48 70 (), 59 , 55 70 (),
57,5 .

: 3 (23%),
4 (31%) 6 (46%).

: 4 7,5 . (5,7).

: Dukes : (7) (%) Dukes C: (6)
(%). 2 1-3
(4) >4

(3)

EGFR:	(4)	(30,7%)	(-)	(9)	
(69,3%)	(+)				
COX-2:	(5)	(38%)	(+)	(8)	(++)
(64%)					

4 - Dukes

/						DUKES	EGFR	COX-2
1		165/07	49		3		(+)	(+)
2		352/07	48		3,5		(+)	(+)
3		554-57/07	52		4		(-)	(+)
4		763-64/07	56		4		(+)	(+)
5		979/07	60		6		(-)	(+)
6		2470-71/07	67		6		(-)	(-)
7		2508-09/07	61		6,5		(-)	(-)
8		2674/07	59		7		(+)	(+)
9		2854/07	69		7		(+)	(+)
10		2859/07	70		7		(-)	(-)
11		2876/07	66		6		(-)	(-)
12		2918/07	49		4,5		(-)	(-)
13		3208/07	53		4		(+)	(+)
14		3260-62/07	54		5		(-)	(+)
15		3842-43/07	50		6		(-)	(+)

5 - Dukes

/						DUKES	EGFR	COX-2
1		1078/07	51		5		(-)	(+)
2		1149-50/07	72		6		(+)	(-)
3		1226-27/07	69		7		(-)	(-)
4		1507/07, 1714/07	70		8		(-)	(+)
5		1554/07	65		8		(-)	(-)
6		1893/07	63		4,5		(-)	(-)
7		1899/07	60		5		(-)	(+)
8		1903-04/07	64		5		(-)	(+)
9		2147-48/07	62		3,5		(-)	(-)
10		2305/07	63		3		(+)	(+)
11		56/07	50		3		(+)	(++)
12		110-12/07	52		3,5		(-)	(+)
13		435/07	69	A	6	B	(+)	(++)
14		438/07	67		4,5		(+)	(++)
15		1153-54/07	60	A	3	B	(+)	(++)
16		1271/07	45		6,5	B	(-)	(+)
17		1320/07	45		8	B	(-)	(+)
18		1388-90/07	42	A	7,5	B	(-)	(+)
19		1396/07	66	A	4	B	(+)	(+)
20		1582/07	79	A	3,5	B	(-)	(+)
21		1707/07	75		5,5		(-)	(+)
22		1807-08/07	67	A	8	B	(-)	(+)

23		1819/07	72		8	B	(+)	(++)
24		2015/07	79		3,5		(+)	(++)
25		2018-19/07	57		5		(-)	(+)
26		2075/07	58		6,5		(-)	(+)
27		2280-82/07	55		7		(-)	(+)
28		2331-33/07	59		8		(+)	(+)
29		2425/07	80		5		(-)	(++)
30		2427-28/07	79		6,5		(+)	(+)
31		2444-45/07	78		5		(+)	(+)
32		2475-76/07	75		3	B	(+)	(++)
33		2535-38/07	64	A	3	B	(-)	(+)
34		2622/07	66		7,5	B	(-)	(+)
35		2626/07	72		6,5	B	(+)	(+)
36		2646/07	70		6	B	(-)	(+)
37		2697/07	70		4,5		(+)	(+)
38		2791-92/07	65		3,5		(+)	(+)
39		2816/07	66		4,5		(-)	(+)
40		2988/07	69		8		(-)	(+)
41		2999/07	79		3		(+)	(-)
42		3039/07, 3158-59/07	80		3,5		(-)	(-)
43		3277-78/07	44		4		(-)	(+)
44		3294-95/07	49		5		(-)	(-)
45		3360-62/07	50		6		(-)	(+)
46		3499/07	68		7,5	B	(-)	(-)
47		3551-52/07	69	A	8	B	(-)	(-)
48		3890/07	66		7		(+)	(+)
49		3892/07	67		5		(+)	(+)
50		3902-03/07	77		4,5		(+)	(+)
51		109/07	70		4,5		(+)	(++)
52		428/07	63		7		(-)	(+)
53		527-29/07	65		7		(+)	(++)
54		639/07	49	A	6,5	B	(+)	(++)
55		1308/07	48		5,5		(-)	(+)
56		2494/07, 3539/07	59		6,5		(+)	(++)
57		3005-06/07	66		7,5		(-)	(+)

6 - Dukes C

/						DUKES	EGFR	COX-2
1		171/07	60		4	C, N1	(-)	(+)
2		391-92/07	80	A	5	C	(-)	(+)
3		462/07	65		3,5	C, N1	(-)	(+)
4		493/07	70		4	C, N1	(+)	(-)
5		498/07	72		4,5	C	(+)	(++)
6		501-03/07	69		7	C	(-)	(+)
7		571-72/07	68		6,5	C	(-)	(+)
8		708-09/07	80		3,5	C, N2, M	(+)	(++)
9		873/07	75		3	C	(+)	(+)
10		1017/07	73		4,5	C	(+)	(++)
11		1053/07	72		5	C	(+)	(+)
12		1086-87/07	49		5,5	C	(+)	(++)
13		1577-78/07	72	A	3,5	C, N1,	(+)	(++)
14		1776-77/07	63		6,5	C, N1	(+)	(++)
15		1971/07, 3316/07, 3341- 43/07	77		6	C, N1	(-)	(++)
16		2010/07	79	A	4	C, N2	(+)	(+)
17		2012-13/07	80		4	C,	(-)	(++)
18		2380/07	69		5	C	(+)	(+)
19		2450-51/07, 2497/07	79		5	C, N1,	(-)	(++)
20		2662/07, 2687/07	70		5	C, N2, M	(+)	(++)
21		3390/07	55		3	C, N2, M	(+)	(++)
22		3905/07	73		3	C, N2, M	(+)	(++)
23		19/07	65		4	C, N2, M	(+)	(++)
24		576-77/07	55		6,5	C, N2	(+)	(++)
25		2767/07	58		6	C, N1, M	(-)	(+)
26		2789/07	62		6	C, 2	(+)	(++)
27		3183-85/07	68		7	C, N1, M	(+)	(++)
28		3651/07, 3690- 91/07	70		7	C, 2	(-)	(+)

100
 2007
 7
 (48%) . 25 , 62
 13
 64,5 (7) 42 80 ,
 (54 %). Duke
 (grade)
 7. COX-2
 EGFR. 52% EGFR 48%
 (7), , 16% COX-2 84%
 COX-2. EGFR COX-2
 Duke
 1 2, 3 4 .

7-



()

(cm)

64.5	5.34
42	3.0
80	8.0

(Dukes)

46	A	54
100	B	15
100	C	57

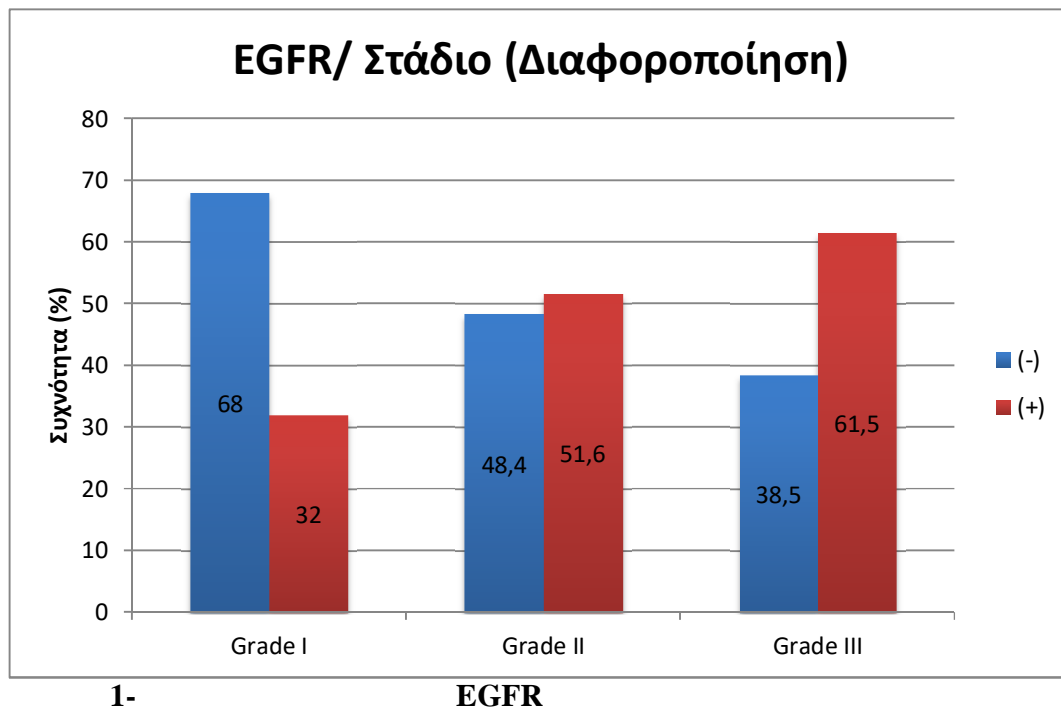
	16	grade I	25
	36	grade I	62
O	48	grade	13
EGFR		COX-2	
(-)	52	(-)	16
(+)	48	(+)	56
Total	100	(++)	28

8-

EGFR

EGFR * Grade Crosstabulation

		GRADE			
		Grade I	Grade II	Grade III	Total
EGFR (-)	Count	17	30	5	52
	% within EGFR	32.7%	57.7%	9.6%	100.0%
	% within GRADE	68.0%	48.4%	38.5%	52.0%
(+) (+)	Count	8	32	8	48
	% within EGFR	16.7%	66.7%	16.7%	100.0%
	% within GRADE	32.0%	51.6%	61.5%	48.0%
Total	Count	25	62	13	100
	% within EGFR	25.0%	62.0%	13.0%	100.0%
	% within GRADE	100.0%	100.0%	100.0%	100.0%

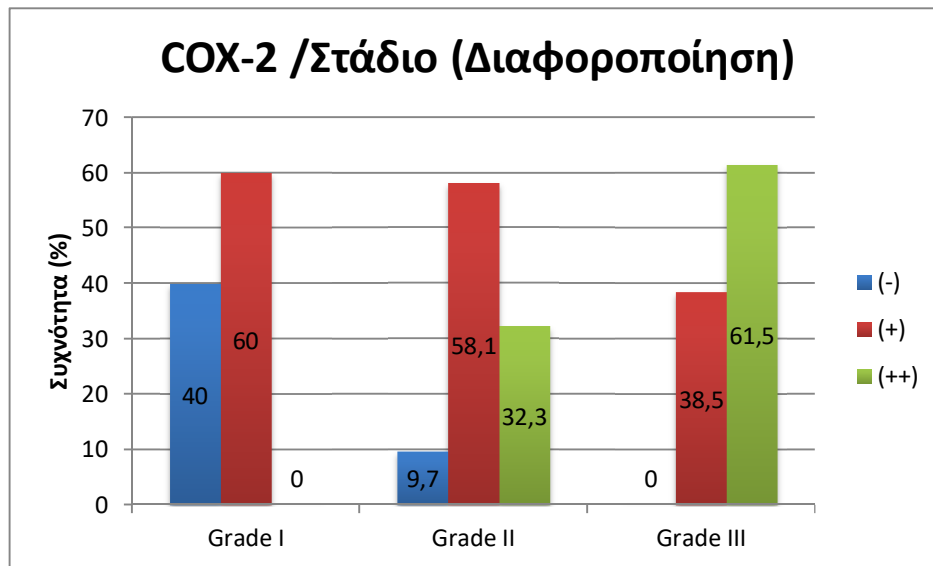


9 -

COX-2

COX-2 * GRADE Crosstabulation

		GRADE			Total
		Grade I	Grade II	Grade III	
COX-2 (-)	Count	10	6	0	16
	% within COX-2	62.5%	37.5%	0.0%	100.0%
	% within GRADE	40.0%	9.7%	0.0%	16.0%
(+) (+)	Count	15	36	5	56
	% within COX-2	26.8%	64.3%	8.9%	100.0%
	% within GRADE	60.0%	58.1%	38.5%	56.0%
(+) (+)	Count	0	20	8	28
	% within COX-2	0.0%	71.4%	28.6%	100.0%
	% within GRADE	0.0%	32.3%	61.5%	28.0%
Total	Count	25	62	13	100
	% within COX-2	25.0%	62.0%	13.0%	100.0%
	% with % within GRADE	100.0%	100.0%	100.0%	100.0%



2-

COX-2

DUKES * EGFR Crosstabulation

		EGFR		Total	
		(-)	(+)		
DUKES	A	Count	9	6	15
		% within DUKES	60.0%	40.0%	100.0%
		% within EGFR	17.3%	12.5%	15.0%
		% of Total	9.0%	6.0%	15.0%
	B	Count	33	24	57
		% within DUKES	57.9%	42.1%	100.0%
		% within EGFR	63.5%	50.0%	57.0%
		% of Total	33.0%	24.0%	57.0%
	C	Count	10	18	28
		% within DUKES	35.7%	64.3%	100.0%
		% within EGFR	19.2%	37.5%	28.0%
		% of Total	10.0%	18.0%	28.0%
Total	Count	52	48	100	
	% within DUKES	52.0%	48.0%	100.0%	
	% within EGFR	100.0%	100.0%	100.0%	
	% of Total	52.0%	48.0%	100.0%	

10 -

COX-2

Dukes

DUKES * COX-2 Crosstabulation

			COX-2			
			(-)	(+)	(++)	Total
DUKES	A	Count	5	10	0	15
		% within DUKES	33.3%	66.7%	0.0%	100.0%
	B	Count	10	35	12	57
		% within DUKES	17.5%	61.4%	21.1%	100.0%
	C	Count	1	11	16	28
		% within DUKES	3.6%	39.3%	57.1%	100.0%
Total		Count	16	56	28	100
		% within DUKES	16.0%	56.0%	28.0%	100.0%

(Pearson Chi-square test)
 (Contingency Coefficient)
 p<0.05
 (IBM SPSS
 v15.0). 1. (grade),

EGFR
 $\chi^2(2)=3.843, p=0.146,$

1. **COX-2**
COX-2 ($\chi^2(4)=26.086, p<0.0001$)

(Contingency Coefficient)
COX-2
 (Contingency Coefficient=0.455,
 p<0.0001).

Dukes 14 15.
 2. (grade),

EGFR Dukes
Dukes ($\chi^2(2)=4.153, p=0.125$)

COX-2
Dukes ($\chi^2(4)=21.279, p<0.0001$).
COX-2 Dukes
 (Contingency Coefficient=0.419,
 p<0.0001),

_____ **COX-2**
 3. **EGFR** **COX-2** .(16-18, 5 6).

Dukes

Pearson

,
(19).

Chi-Square Tests (EGFR vs Grade)

	Value	df	Asymptotic Significance (2- sided)
Pearson Chi-Square	3.843 ^a	2	.146
Likelihood Ratio	3.917	2	.141
Linear-by-Linear Association	3.599	1	.058
N of Valid Cases	100		

a. 0 cells (0.0%) have expected count less than 5. The minimum expected count is 6.24.

Chi-Square Tests (COX-2 vs Grade)

	Value	df	Asymptotic Significance (2- sided)
Pearson Chi-Square	26.086 ^a		.000
Likelihood Ratio	31.474	4	.000
Linear-by-Linear Association	24.045	1	.000
N of Valid Cases	100		

a. 3 cells (33.3%) have expected count less than 5. The minimum expected count is 2.08.

Symmetric Measures (COX-2 vsGrade)

		Value	Asymptotic Standard Error ^a	Approximate T ^b	Approximate Significance
Nominal by Nominal	Contingency Coefficient	.455			.000
Interval by Interval	Pearson's R	.493	.066	5.607	.000 ^c
Ordinal by Ordinal	Spearman Correlation	.496	.065	5.657	.000 ^c
N of Valid Cases		100			

a. Not assuming the null hypothesis.

b. Using the asymptotic standard error assuming the null hypothesis.

c. Based on normal approximation.

14 ó

-

EGFR

Duke

Chi-Square Tests EGFR-DUKES

	Value	df	Asymptotic Significance (2-sided)
Pearson Chi-Square	4.153 ^a	2	.125
Likelihood Ratio	4.189	2	.123
Linear-by-Linear Association	3.186	1	.074
N of Valid Cases	100		

15 ó

-

COX-2

Duke

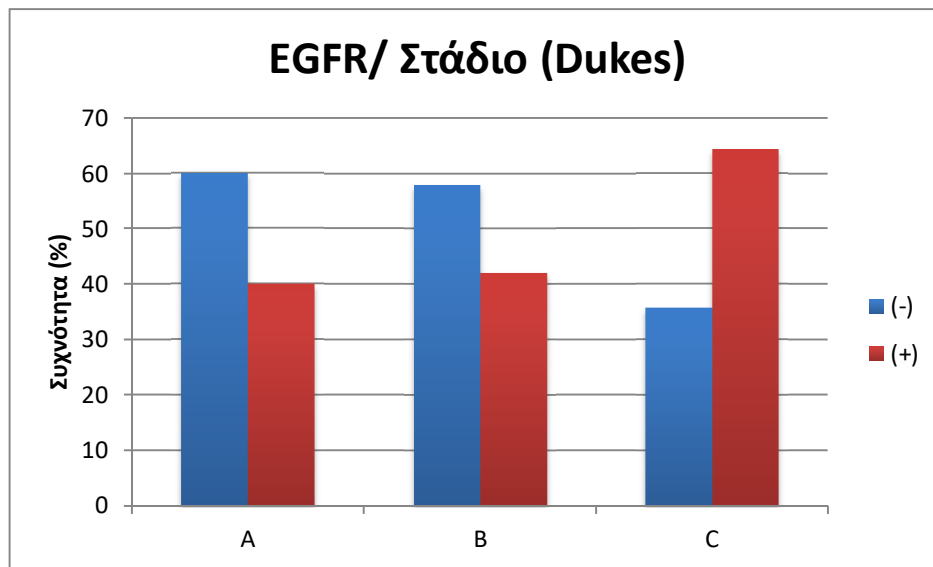
Chi-Square Tests DUKES * COX-2

	Value	df	Asymptotic Significance (2-sided)
Pearson Chi-Square	21.279 ^a	4	.000
Likelihood Ratio	24.302	4	.000
Linear-by-Linear Association	19.147	1	.000
N of Valid Cases	100		

Symmetric Measures

		Value	Asymptotic Standard Error ^a	Approximate T ^b	Approximate Significance
Nominal by Nominal	Phi	.461			.000
	Cramer's V	.326			.000
	Contingency Coefficient	.419			.000
Interval by Interval	Pearson's R	.440	.074	4.847	.000 ^c

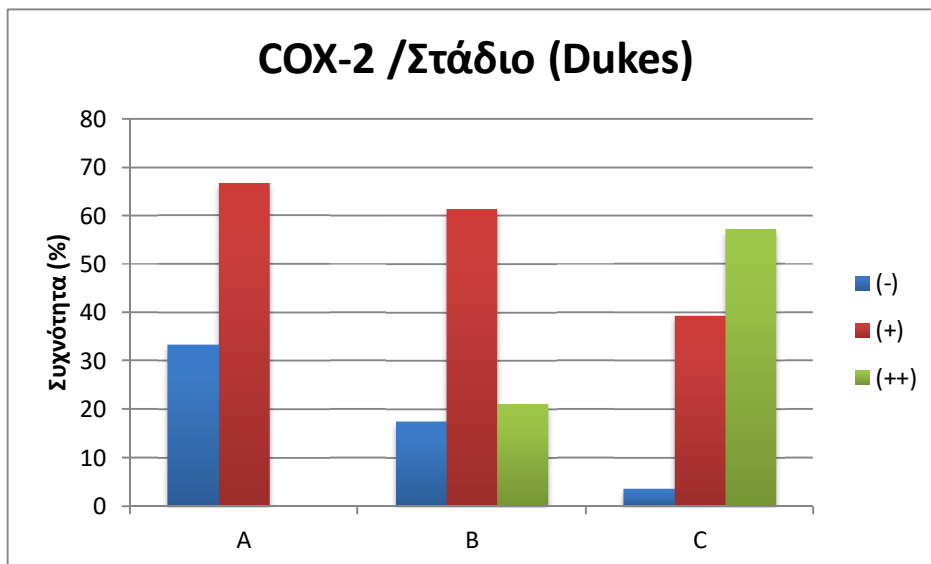
Ordinal by Ordinal	Spearman Correlation	.448	.075	4.955	.000 ^c
N of Valid Cases		100			



3.

EGFR

Duke



4.

COX-2

Duke

EGFR 16 6

EGFR * Crosstabulation

		male	female	Total
EGFR (-)	Count	27	25	52
	% within EGFR	51.9%	48.1%	100.0%
	% within	50.0%	54.3%	52.0%
EGFR (+)	Count	27	21	48
	% within EGFR	56.3%	43.8%	100.0%
	% within	50.0%	45.7%	48.0%
Total	Count	54	46	100
	% within EGFR	54.0%	46.0%	100.0%
	% within	100.0%	100.0%	100.0%

17 6 - **EGFR**
Chi-Square Tests sex vs EGFR

	Value	df	Asymptotic Significance (2-sided)	Exact Sig. (2-sided)	Exact Sig. (1-sided)
Pearson Chi-Square	.188 ^a	1	.664		
Continuity Correction ^b	.054	1	.816		
Likelihood Ratio	.188	1	.664		
Fisher's Exact Test				.692	.408
Linear-by-Linear Association	.186	1	.666		
N of Valid Cases	100				

a. 0 cells (0.0%) have expected count less than 5. The minimum expected count is 22.08.

b. Computed only for a 2x2 table

186

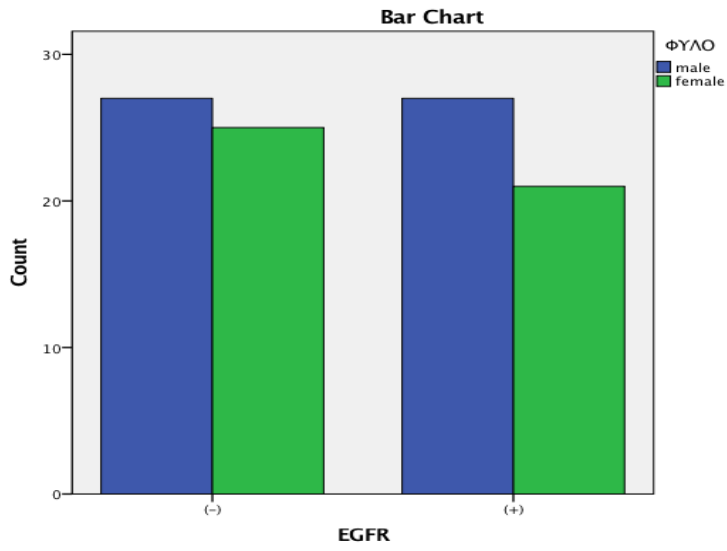
COX-2
COX-2 * **Crosstabulation**

			male	female	Total
COX-2	(-)	Count	7	9	16
		% within COX-2	43.8%	56.3%	100.0%
		% within	13.0%	19.6%	16.0%
	(+) (top)	Count	34	22	56
		% within COX-2	60.7%	39.3%	100.0%
		% within	63.0%	47.8%	56.0%
	(+) (bottom)	Count	13	15	28
		% within COX-2	46.4%	53.6%	100.0%
		% within	24.1%	32.6%	28.0%
Total	Count	54	46	100	
	% within COX-2	54.0%	46.0%	100.0%	
	% within	100.0%	100.0%	100.0%	

Chi-Square Tests sex vs COX-2

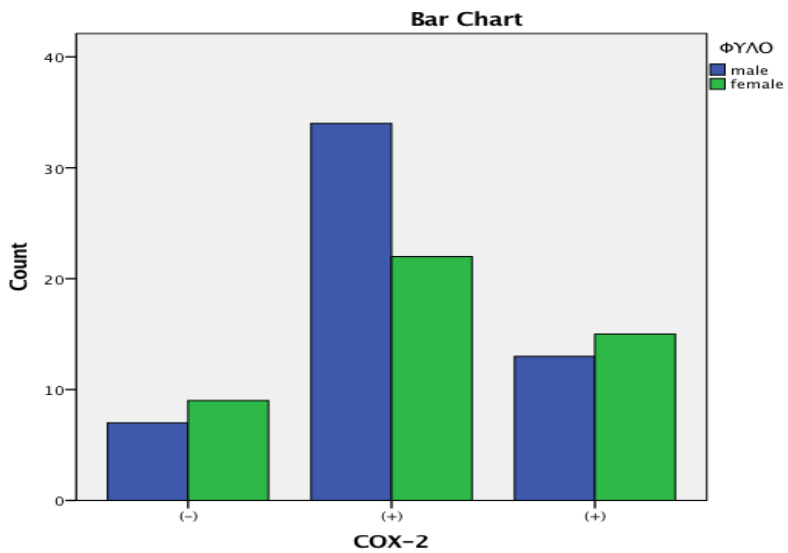
	Value	df	Asymptotic Significance (2-sided)
Pearson Chi-Square	2.339 ^a	2	.310
Likelihood Ratio	2.344	2	.310

Linear-by-Linear Association	.022	1	.883
N of Valid Cases	100		



5.

EGFR



6.

COX-2

Correlations

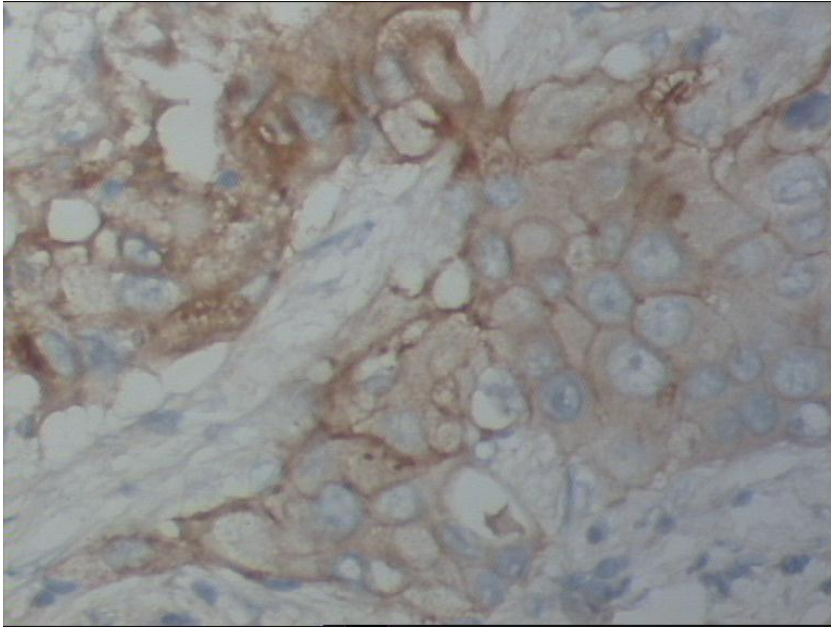
		EGFR	
	Pearson Correlation	1	.179
	Sig. (2-tailed)		.075
	N	100	100
EGFR	Pearson Correlation	.179	1
	Sig. (2-tailed)	.075	
	N	100	100
		COX-2	
	Pearson Correlation	1	.086
	Sig. (2-tailed)		.398
	N	100	100
COX-2	Pearson Correlation	.086	1
	Sig. (2-tailed)	.398	
	N	100	100
		GRADE	
	Pearson Correlation	1	.117
	Sig. (2-tailed)		.245
	N	100	100
GRADE	Pearson Correlation	.117	1
	Sig. (2-tailed)	.245	

	N	100	100
			DUKES
	Pearson Correlation	1	.357**
	Sig. (2-tailed)		.000
	N	100	100
DUKES	Pearson Correlation	.357**	1
	Sig. (2-tailed)	.000	
	N	100	100

** . Correlation is significant at the 0.01 level (2-tailed).

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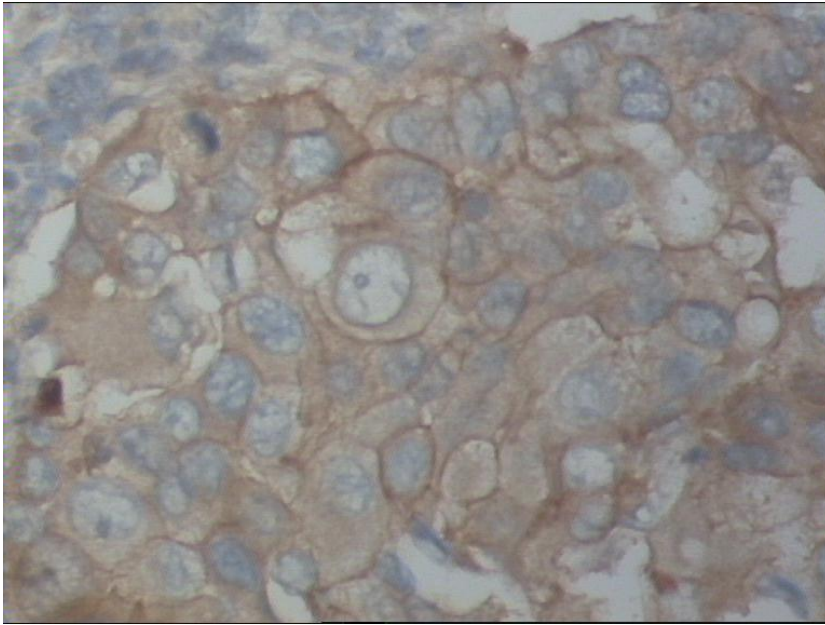
EGFR

(x 240)

(

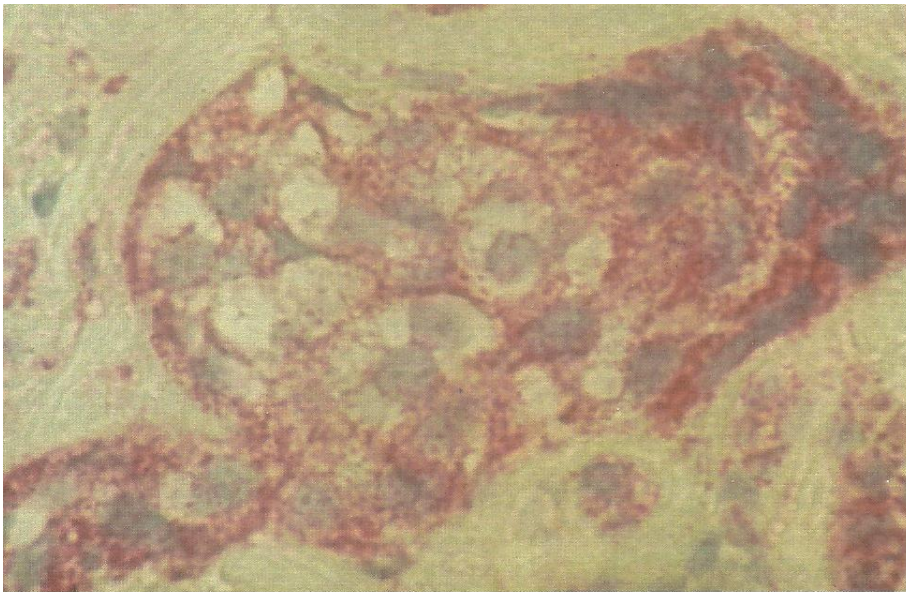
EGFR (+).

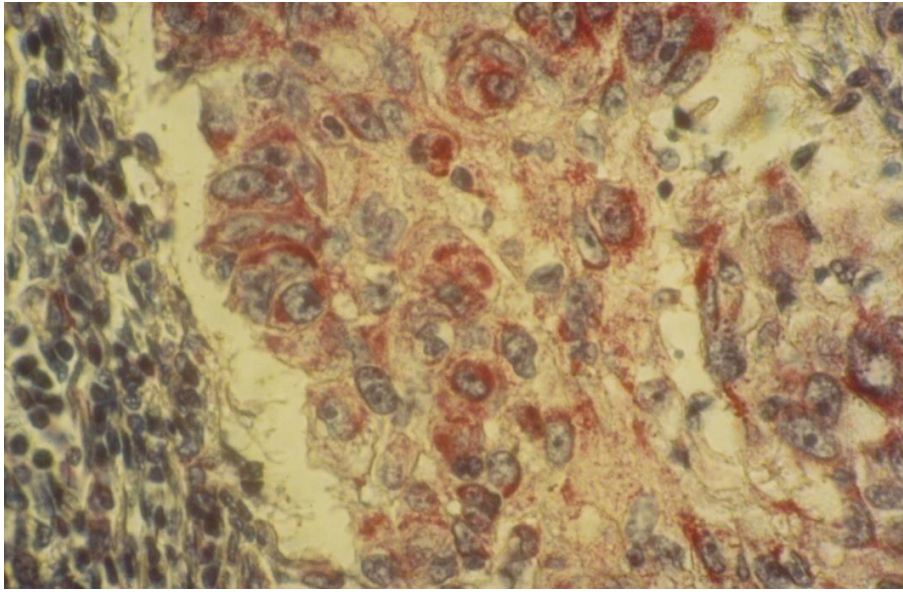
(x 240)



EGFR (+)

(x 400)





COX2
(120)

-

(1, 2).

(3).

2

50%

85%

(4-10).

(11-15).

EGFR

COX2

EGFR 100

(54%).

(48%).

64,5

42-80

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, 52%

EGFR 48% , 16%

COX2 84%

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EGFR,

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COX2

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COX2

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Se Min Jang et al,

COX2,

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COX2

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COX2

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COX

2,

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COX2

1.
COX2

EGFR

2.

COX2

3.

EGFR

COX2

- ABSTRACT

100
 , ø
 2007 . H
 (48%) . 25
 , 62
 13
 64,5 (7) 42 80 ,
 (54 %). Duke (grade)

()	(cm)
64.5	5.34
42	3.0
80	8.0

(Dukes)	
46	A 54
100	B 15
100	C 57

	16	grade I	25
	36	grade I	62
O	48	grade	13
EGFR		COX-2	
(-)	52	(-)	16
(+)	48	(+)	56
Total	100	(++)	28

COX-2 84% EGFR 48% COX-2 EGFR. 52%
 (), 16%
 COX-2. . ,

:

1. COX2 EGFR
2. COX2
3. EGFR

COX2

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