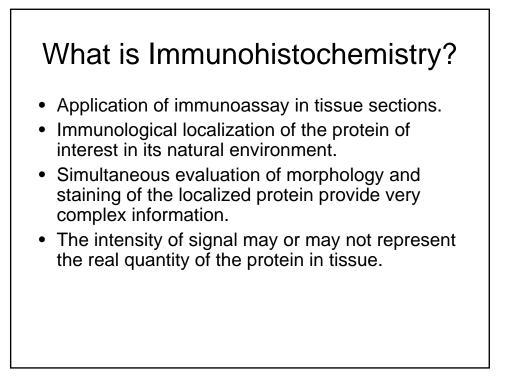


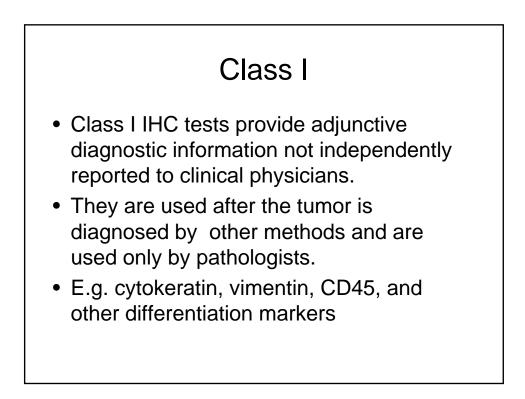


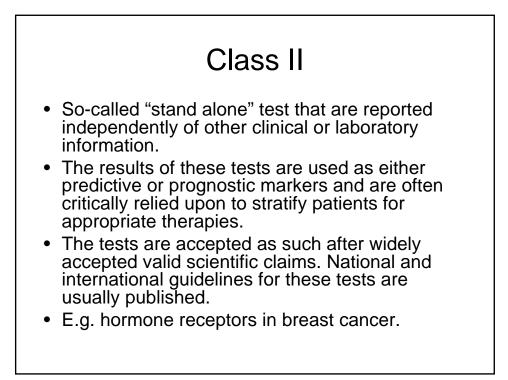
The Role of Medical Laboratories in Patients' Care

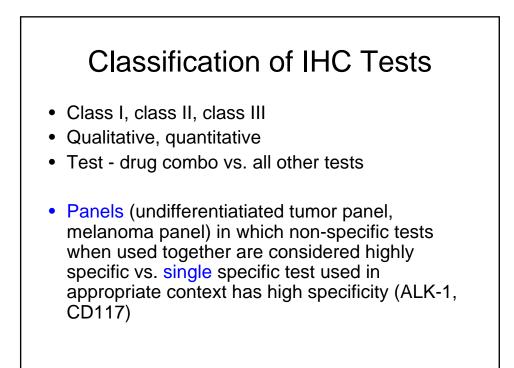
• Dr. J. Butany:

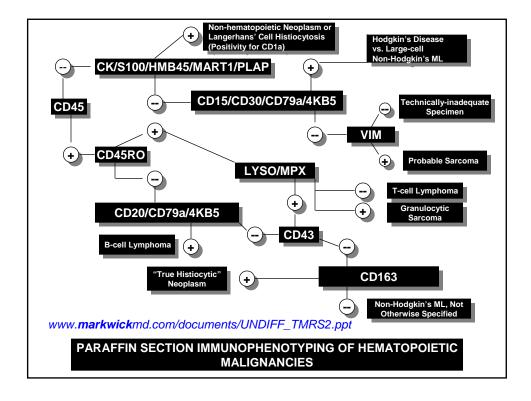
"Canada's medical laboratory system is the foundation upon which good patient care, diagnosis and treatment rest."

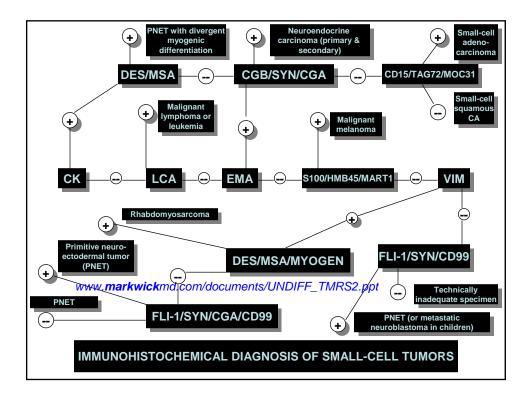


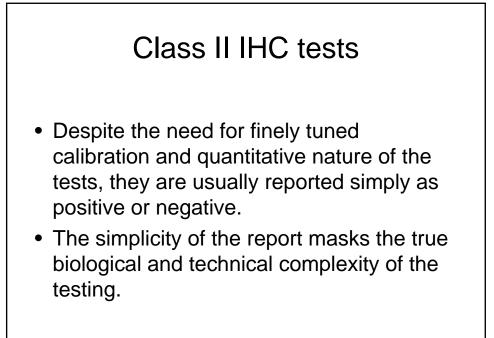


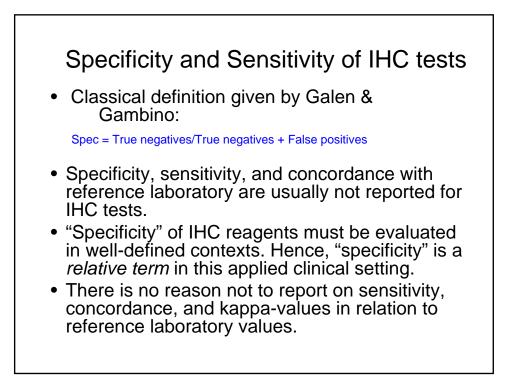


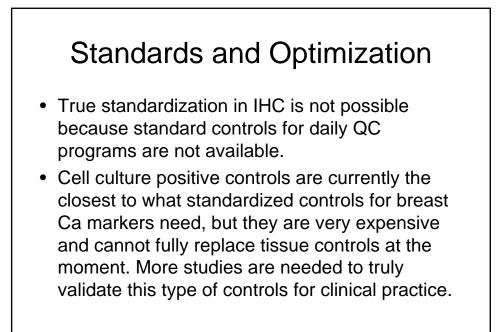












NordiQC Results with Cell Culture
Positive Controls

	Tiss	sues a	s Pos. Con	trols
Cell Lines	Optimal	Good	Borderline	Poor
Optimal	37	1	0	12
Good	4	1	0	1
Borderline	3	0	1	8
Poor	0	1	0	11

http://www.nordiqc.org/Run-23-B5/Assessment/Assessment-HER-2.htm

Use of Cell Lines as Positive Controls: Results and Conclusion

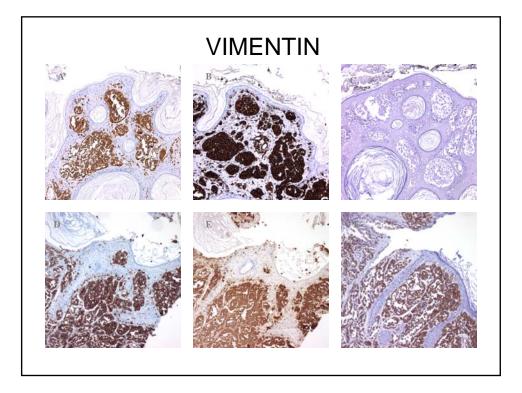
- 1. An insufficient (false negative) reaction in the breast ductal carcinoma no. 3 in combination with an optimal staining of the cell lines. This was seen in 13/17 cases.
- 2. A sufficient staining in the histological specimens in combination with an insufficient staining of the cell lines due to impaired morphology of the cell lines, probably as a results of excessive retrieval.
- 3. These data indicate that histological specimens should be preferred for EQA of HER-2. However, due to potential heterogeneity of tissue material, cell cultures may be valuable as a supplement.

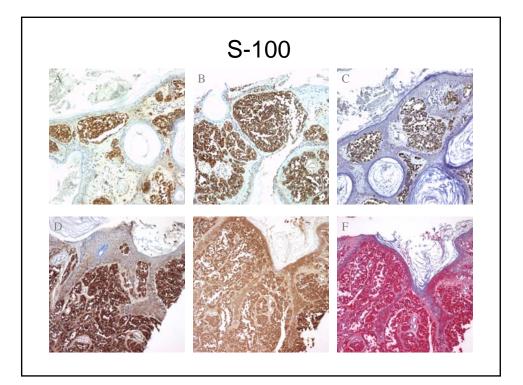
http://www.nordiqc.org/Run-23-B5/Assessment/Assessment-HER-2.htm

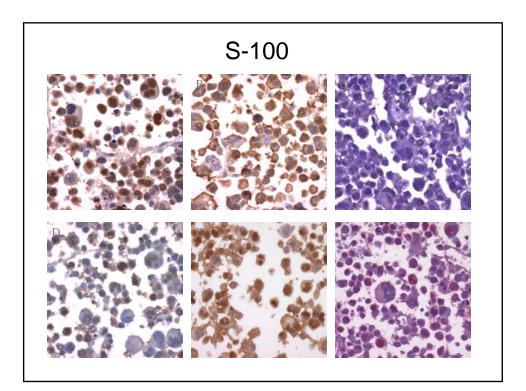
Main Conclusions Regarding Standardization

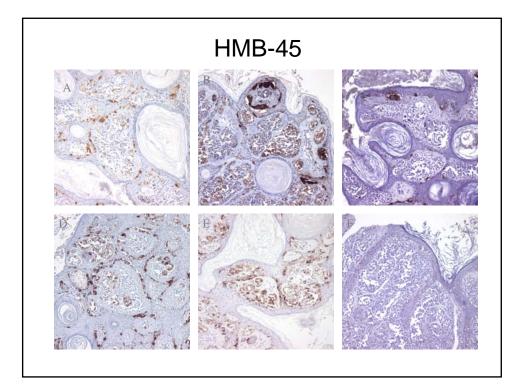
- No standardized positive controls No standardization.
- Standardization of protocols is meaningless without control standardization.
- Standardization of positive controls also includes agreement or standardization of expected results in control tissues.
- "Standardization" is greatly misused term in this context.
- Standardization is possible only if there are socalled "gold standards" for reference values.

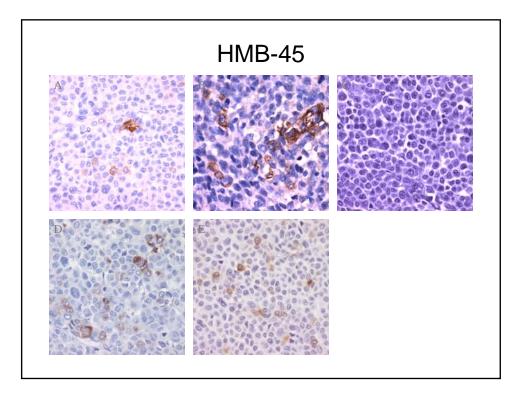
	http://www.nordiqc.org	
NordiQC	Participants (N)	Sufficient Results (%)
Run 8 2003	71	45
Run 10 2004	77	67
Run 13 2005	89	84
Run B1 2006	68	75
Run B3 2007	73	84
Run B5 2008	107	79

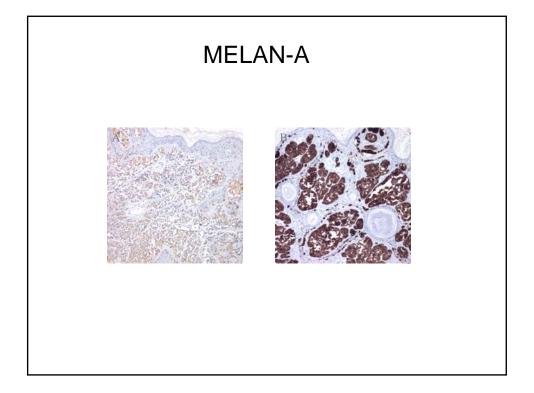


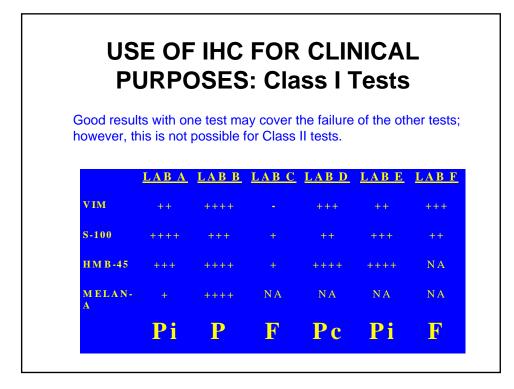




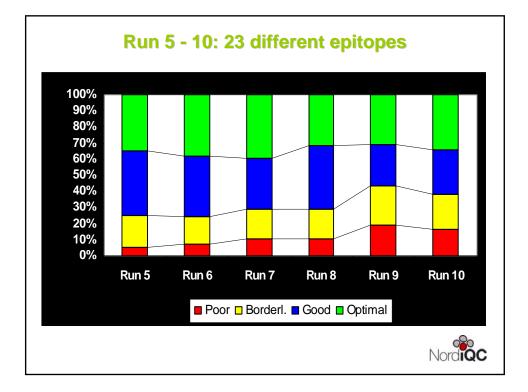




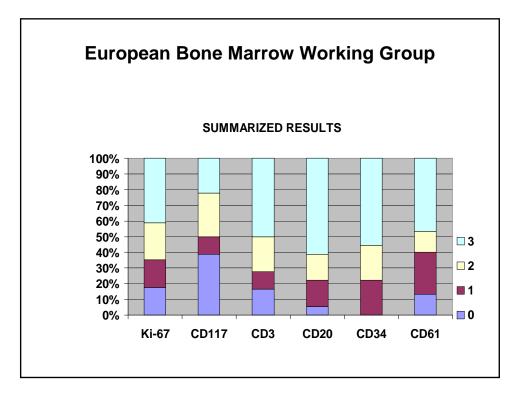




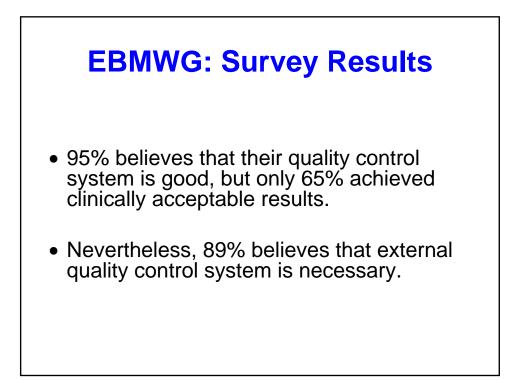




		NordiQC
	Run 8, 9 & 10 n: 382 insuffici	ent staining:
Fa	Ise negative:	
1.	Too dilute primary ab. conc.	127 (33%)
2.	Inappropriate primary ab.	51 (13%)
3.	Insufficient HIER	94 (25%)
4.	Inappropriate epitope retrieval	54 (14%)
5.	Unexplained	22 (6%)
Fa	Ise positive:	
1.	Too high primary ab. conc.	7(2%)
2.	Inappropriate primary ab.	14(4%)
3.	Excessive retrieval	1 (<1%)
4.	Unspecific reaction of the detection system	10 (3%)
5.	Unexplained	2 (<1%) 31

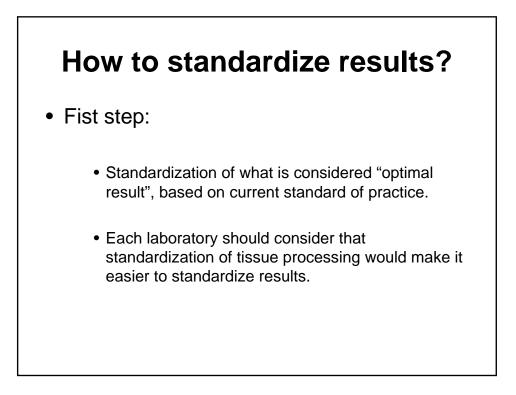


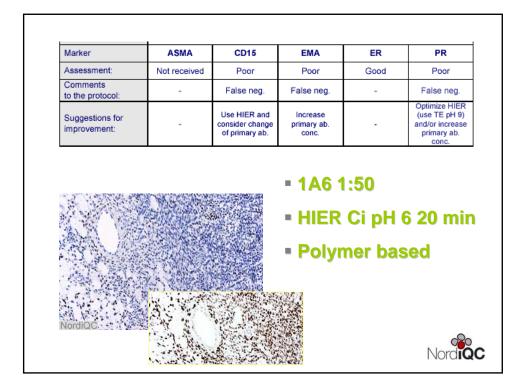
0	K: 07	00447	0.00	0504	0004	0000		Tatal
Center	Ki-67	CD117	CD3	CD61	CD34	CD20	Suboptimal/Poor	Total
1	1	1	3	1	1	1	5	6
2	1	0	2	2	1	2	6	6
3	1	3	3	3	3	3	1	6
4	3	1	3	3	2	3	1	6
5	2	2	2	2	3	2	0	6
6	2	3	2	3	3	3	0	6
7	0	0	0		3	3	3	5
8	3	2	3	3	3	3	0	6
9	3	0	3	0	2	3	2	6
10	3	2	0	1	3	3	2	6
11	2	2	2		3	3	0	5
12	0	0	1		2	2	3	5
13	3	0	3	1	1	1	4	6
14	0	3	3	0	3	3	2	6
15		0	0	1	1	0	5	5
16	2	2	3	3	3	3	0	6
17	3	3	1	3	3	1	2	6
18	3	0	3	3	2	3	1	6
Total							35.50%	104

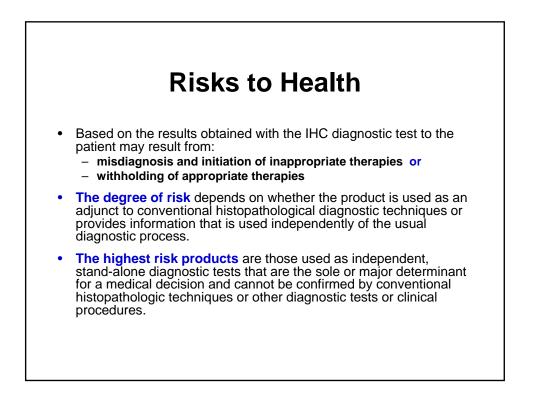


What do we want to optimize or standardize?

- METHODS Not necessarily!
- RESULTS Obligatory!







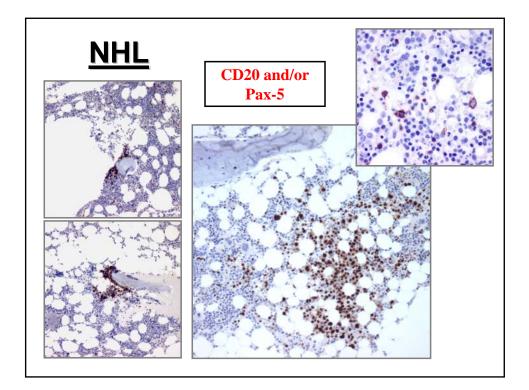
FDA is focused on whether this level of regulation is adequate for the protection of public health

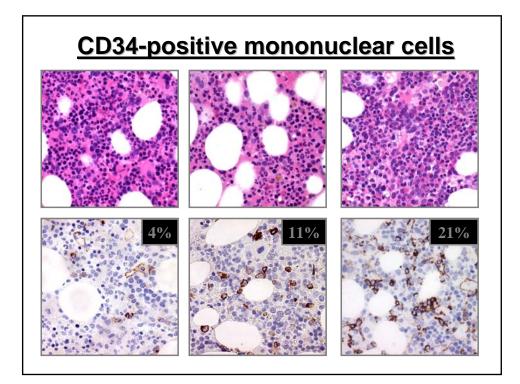
- FDA is aware that variability in IHC results may be introduced at every step:
 - Collection and fixation of the specimen,
 - Automated processing,
 - Embedding and sectioning,
 - Staining of the final slide preparation, and
 - Microscopic interpretation by the pathologist.

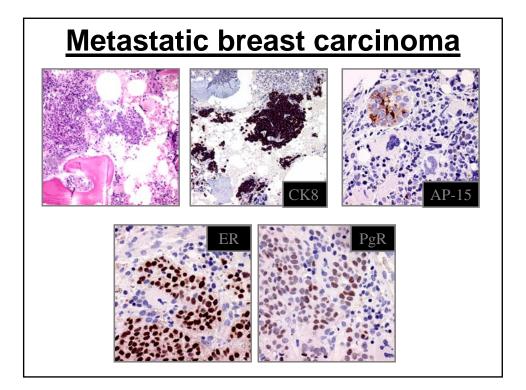
FDA counts on:

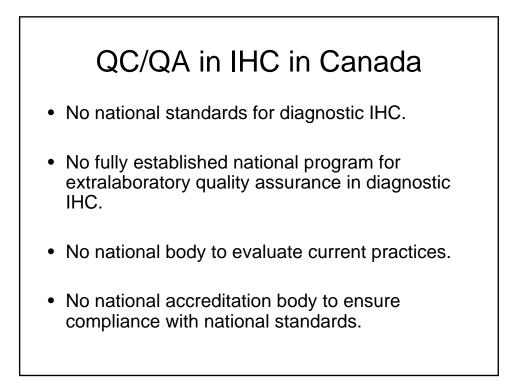
Ongoing initiatives by professional organizations and manufacturers directed at ensuring that pre- and postanalytic, as well as analytic procedures, are properly performed.

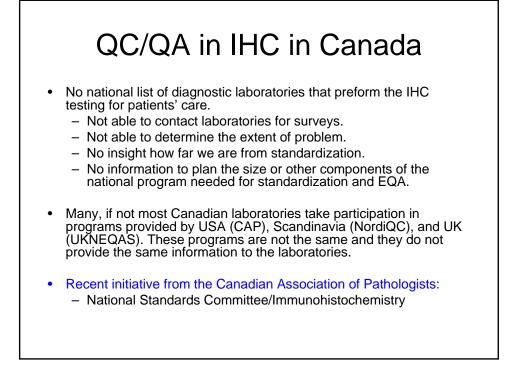
That there is clear distinction in laboratory practices regarding Class I and Class II tests in regard quality control/quality assurance measures by the laboratories.

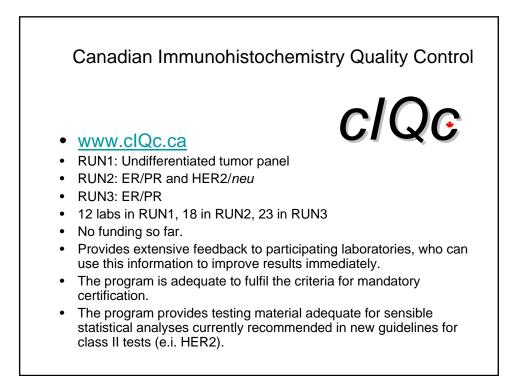


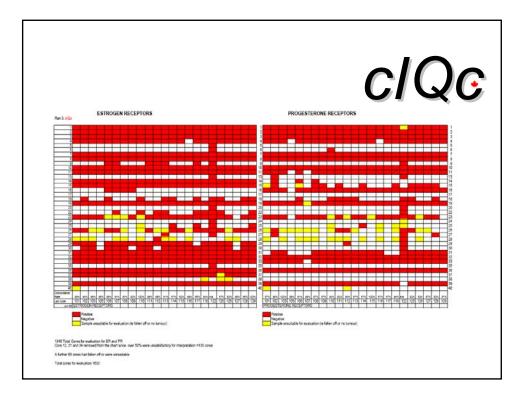


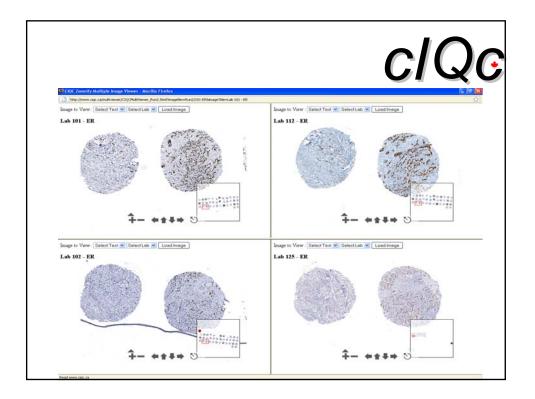












Gen Max <th< th=""><th></th><th></th><th></th><th></th><th></th><th></th><th></th><th></th><th></th><th></th><th></th><th></th><th></th><th></th><th></th><th></th><th></th></th<>																	
det Unit Description Form Description Spectro	Lib Co						Prim		Dist	Disent		Detection	Petertion		С	Enha	Type of
Unit Unit <t< th=""><th>de</th><th>val</th><th>Instrument</th><th>Temp</th><th>Time</th><th>Buffer</th><th>Clone</th><th>Supplier</th><th>ion</th><th>Туре</th><th>Time</th><th>System</th><th>System Name</th><th>Supplier</th><th>Incubation Time</th><th>ent</th><th>t</th></t<>	de	val	Instrument	Temp	Time	Buffer	Clone	Supplier	ion	Туре	Time	System	System Name	Supplier	Incubation Time	ent	t
Unit Unit <t< td=""><td>101</td><td>Yes</td><td>Ventana XT</td><td>90</td><td>30 min</td><td>CC1</td><td>SP1</td><td>Labvision</td><td>1.50</td><td>Ventana</td><td>32 min</td><td>Mew DAB</td><td></td><td>Ventana</td><td></td><td></td><td></td></t<>	101	Yes	Ventana XT	90	30 min	CC1	SP1	Labvision	1.50	Ventana	32 min	Mew DAB		Ventana			
10. 10. Number 6.2 6.2 6.9 10.<					13 min + 20	1mM EDTA				Dako Ab	30 min						
101 101 <td>102</td> <td>Yes</td> <td></td> <td>100</td> <td>min cool down</td> <td>pH 9</td> <td>SP1</td> <td>Latvision</td> <td></td> <td>DI</td> <td>RT</td> <td></td> <td>Envision +</td> <td>Dako</td> <td>30 min RT</td> <td>Yes</td> <td>Cv/80,</td>	102	Yes		100	min cool down	pH 9	SP1	Latvision		DI	RT		Envision +	Dako	30 min RT	Yes	Cv/80,
00 00 $000000000000000000000000000000000000$	103	Yes	Benchmark	42	60 min	CC1	SP1		luted		8 min	New	LSAB	Ventana	NA	No	
100 100 Description 1 0.00 Description 1 Descripion 1 <thdescription 1<="" th=""> D</thdescription>			Ventana			CC1 (Cat#		(Cat# RM		(Cate TA					Standard Ventana		
ON Part I Trange 113 3 mm (c) (c) (c) OP Vertex 17 Dest/Part 67 (E) (c) Vertex Non (c) Non	105	Yes	Discovery XT MicroMed	100	Standard	950-1240	8P1	9101-80	1.50	125-UC)	60 min		DAB MAP	Ventana	Protocol	No	
n_{11} n_{12}	106	Yes	T/Trega	115	3 min	HCI pH9	OFI	Vector	1:75	Dako Flex	RT	Eite ABC		Vector	30 min		
108 108 Model 08 8 model 09 100						Dako Target Retrieval		Labulation N	1:20	Dako Ab							
100 100 <td>108</td> <td>Yes</td> <td>Medical</td> <td>98</td> <td>8 min</td> <td>System</td> <td>SP1</td> <td>eo Markers</td> <td>0</td> <td>diluent</td> <td>30 min</td> <td>HRP-DAB</td> <td>HRP system</td> <td>Dako</td> <td>30 min</td> <td>No</td> <td></td>	108	Yes	Medical	98	8 min	System	SP1	eo Markers	0	diluent	30 min	HRP-DAB	HRP system	Dako	30 min	No	
Open Deckoder Open Dec			Biccare			Reveal-									linke 10 min, label		
100 No. Vertex U No. 198 All Al	109	Yes			1 min	Biocare	GFI	Vector		Zymed	1 hour		Histo +	Zymed	= 10 min	No	
111 No. AT 60 30 mm Operating Operating Operating Vestman Annual Mark 112 Area Area<	110	Yes	Ventana ES		19.45 min	Citrate Buffer	SP1	Ventana	Predi luted		28 min		AEC detection	Ventana	Unknown	No	
$ \begin{array}{ c c c c c c c c c c c c c c c c c c c$		¥	Benchmark	66	10 min	001 (m)4	601	Mastana	Predi		22 min		1 Bran (and	Masters		Ver	Copper
131 Feat. Benchman Ben	111	Tes	AT	90	30 min	CC1 TRIS	39/1		L(Nex)	New	32 min					785	Sulfate
120 Main			Benchmark			buffer oH 8		Ventana (Cat#760-	Pred			based detection	Ultraview universal DAB	Ventana (Caté 760-			
131 Ten Monometer 000 min conduction Origin (min party of min part	112	Yes	from Ventana	95-100			SP1		luted		20 min			500)			
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	113	Yes	Microweve	100	min cool down	Citrate pH 6	GFI	Ventana	NA	NA	32 min		New	Ventana		Yes	Sulfate
114 Max Max <td></td> <td></td> <td>Ventana</td> <td></td>			Ventana														
$ \begin{array}{ c c c c c c c c c c c c c c c c c c c$	114	Yes	XT	90	4 min	2x880	8P1	Labvision	1.50	Ventana	32 min		XT, Mew, DAB	Ventana		Yes	Copper
19 Ver. XT 00 30mm Contract Non- Vertice			Benchmark						Predi				Wew DAS				
Image: Notice of the sector of the	115	Yes	XT	95	30 min	CC1 Solution	SP1	Ventana			32 min			Ventana		Yes	Copper
19 Vertices 0 + vertices 0 - vertices <			Benchmark					for for		Antibody	48		Ultraview				
Image: Processing of the state of	***	Yes		15.100	30 min	001/60740	501					Phosphatase Red		Ventere	12 min (1/14)	No	
117 Participal Description Description Description Optimized Optimized <thoptimized< th=""> Optimized <t< td=""><td></td><td></td><td>10.02.0</td><td>Accordin</td><td>According to</td><td></td><td></td><td>500 p</td><td>-</td><td>(recound)</td><td>0.10</td><td></td><td></td><td></td><td></td><td>.40</td><td></td></t<></thoptimized<>			10.02.0	Accordin	According to			500 p	-	(recound)	0.10					.40	
Image: Normal State (Normal State (Norma State (Norma State (Norma State (Norma State (Norma State (Norma	117	Yes	Benchmark	g to Protocol		CC1 Ventana	SP1	Ventana	No	No	32 min	Mew DAB	Wew DAB detection kit	Ventana	According to Protocol		
19 Ym Discovery XT Marcial Discovery XT Marcial Para Discovery XT Marcial Ym Buffer 19 Ym Discovery XT Marcial Discovery XT Marcial Pres Discovery XT Marcial Pres Discovery XT Marcial Pres Discovery XT <																	
Discussion (2) Discussion (2) Discusision (2) Discussion (2) Discus						Cet Conditioning		medical	y-to-		min			Medical			Sulfate
U2 Vers Obstanting (main constraints) Distanting (main constraints) <thdistanti< td=""><td>118</td><td>Yes</td><td></td><td>Max 100</td><td>Standard</td><td>1 Dato Tarnet</td><td>SP1</td><td>Systems</td><td>use</td><td></td><td>37°C</td><td></td><td>DAB-MAP</td><td>Systems</td><td>10/10/5 (Mourie</td><td>Yes</td><td>Solution</td></thdistanti<>	118	Yes		Max 100	Standard	1 Dato Tarnet	SP1	Systems	use		37°C		DAB-MAP	Systems	10/10/5 (Mourie	Yes	Solution
150 Verticity 27 95:100 20:nm C/S 97 Verticity 20			Decloaking			Retrieval					30 min				probe/HRP		
USE View View District Office Office <thoffice< th=""> <thoffice< th=""></thoffice<></thoffice<>	122	Yes	Chamber	120	30 sec	SolA pH 9	OFIL	Biocare			RT	Polymer	MAch4	Biocare	polymer(DAB)	-	
Type Multi-Answer Dirac Dirac <thdirac< th=""> <thdirac< th=""> <thdirac< th=""></thdirac<></thdirac<></thdirac<>	125	Yes	Ventana XT	95-100	30 min	CC1	SP1	Ventana			16 min		Ultraview	Ventena		No	
V27 VP V2021 VP Pare Cond Code Code Pare Code Values			Mcroweve		min, 20 min	EDTA pH 8 #			Predi			vidin /View			kit (approx 1 1/2		
128 Yes XT 95 30 min CC1 SP1 Vertana 0 37* 0.6 ///wer Vertana Appr Vertana Copper 128 Yes XT 95 30 min CC1 SP1 Vertana 0.0 Vertana	127	Yes	1000 Watts	Power	cooi down	0.05	SP1	Ventana	lute		32 min	Det Kit	Wew Det	Ventana	hours] total	No	
128 Yes XT 95 30 min CC1 5P1 Vertana lute use C DAB New Vertana protocols Yes Copper Vision Boordiners Boord Max			Benchmark								Q 37*						
Biosystems Bond Max	128	Yes		95	30 min	CC1	SP1	Ventana	lute	u50	C	DAB	New	Ventana	protocols	Yes	Copper
			Biosystems			Bond Max	0.01	1004									

