

phosphoELISA™ Kit Catalog # KHR9131 (96 tests)

IR [pYpY1162/1163]

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Contents and Storage

Storage Store at 2 to 8°C.

Contents

Reagent	96 Test Kit
<i>IR</i> [<i>pYpY1162/1163</i>] <i>Standard</i> : Contains 0.1% sodium azide. Refer to vial label for quantity and reconstitution volume.	2 vials
Standard Diluent Buffer. Contains 0.1% sodium azide; 25 mL per bottle.	1 bottle
<i>IR (β-subunit) Antibody Coated Well</i> s. 12 x 8 Well Strips.	1 plate
<i>IR [pYpY1162/1163] Detection Antibody.</i> Contains 0.1% sodium azide; 11 mL per bottle.	1 bottle
Anti-Rabbit IgG HRP (100X). Contains 3.3 mM thymol; 0.125 mL per vial.	1 vial
HRP Diluent. Contains 3.3 mM thymol; 25 mL per bottle.	1 bottle
Wash Buffer Concentrate (25X). 100 mL per bottle.	1 bottle
Stabilized Chromogen, Tetramethylbenzidine (TMB). 25 mL per bottle.	1 bottle
Stop Solution; 25 mL per bottle.	1 bottle
Plate Covers, adhesive strips.	3

Disposal Note This kit contains materials with small quantities of sodium azide. Sodium azide reacts with lead and copper plumbing to form explosive metal azides. Upon disposal, flush drains with a large volume of water to prevent azide accumulation. Avoid ingestion and contact with eyes, skin and mucous membranes. In case of contact, rinse affected area with plenty of water. Observe all federal, state and local regulations for disposal.

Safety All blood components and biological materials should be handled as potentially hazardous. Follow universal precautions as established by the Centers for Disease Control and Prevention and by the Occupational Safety and Health Administration when handling and disposing of infectious agents.

Introduction

Purpose The Invitrogen IR [pYpY1162/1163] ELISA is designed to detect and quantify the levels of insulin receptor that are phosphorylated at tyrosine residues 1162 and 1163 of insulin receptor. Both natural (heterotetrameric) and recombinant IR react in this assay. This ELISA is not cross-reactive with IGF-1R and allows a differentiation of the phosphorylation and activation of IR from that of IGF-1R. Although this ELISA kit is developed using human cells, cross-reactivity with mouse and rat insulin receptor is documented. This assay is intended for detection of IR [pYpY1162/1163] from lysates of cells. For normalizing the IR content of the samples, an IR (β -subunit) ELISA kit which is independent of phosphorylation status is available from Invitrogen (Cat. #KHR9111).

For Research Use Only. CAUTION: Not for human or animal therapeutic or diagnostic use

Principle of the Method

The Invitrogen IR [pYpY1162/1163] kit is a solid phase sandwich Enzyme Linked-Immuno-Sorbent Assay (ELISA). A monoclonal antibody specific for IR $(\beta$ -subunit) (regardless of phosphorylation state) has been coated onto the wells of the microtiter strips provided. Samples, including a standard containing IR [pYpY1162/1163], control specimens and unknowns, are pipetted into these wells. During the first incubation, the IR antigen binds to the immobilized (capture) antibody. After washing, an rabbit monoclonal antibody specific for IR phosphorylated at tyrosines 1162 and 1163 is added to the wells. During the second incubation, this antibody serves as a detection antibody by binding to the immobilized IR protein captured during the first incubation. After removal of excess detection antibody, a horseradish peroxidase-labeled Anti-Rabbit IgG (Anti-Rabbit IgG HRP) is added. This binds to the detection antibody to complete the four-member sandwich. After a third incubation and washing to remove all the excess Anti-Rabbit IgG HRP, a substrate solution is added, which is acted upon by the bound enzyme to produce color. The intensity of this colored product is directly proportional to the concentration of IR [pYpY1162/1163] present in the original specimen.

Background Information

Insulin receptor (IR), a cell surface receptor, binds insulin and mediates its action on target cells. Insulin receptor belongs to the superfamily of the growth factor receptor tyrosine kinases that regulate multiple signaling pathways through activation of a series of phosphorylation cascades. The insulin receptor is a heterotetrameric membrane glycoprotein consisting of disulfide-linked subunits in a $\beta-\alpha-\alpha-\beta$ configuration. The α -subunit (135 kDa) is completely extracellular, whereas the β -subunit (95 kDa) possesses a single transmembrane domain with tyrosine kinase activity. Insulin binding to the extracellular domain leads to autophosphorylation of the receptor and activation of the intrinsic tyrosine kinase activity, which allows appropriate substrates to be phosphorylated.

Once activated, the IR initiates a variety of metabolic functions including glucose transport, glycogen synthesis, protein synthesis, translational control and mitogenesis. Defects in IR signaling pathway result in insulin resistance and thus high blood glucose associated with type II/non-insulin-dependent diabetes. The IR also plays an important role in neurological function and in hypertension. Important down-stream proteins associated with IR signaling include IRS-1, JAK1, JAK2, STAT1, STAT3, PI3 kinase, GSK-3b, PKC, SHC, ERK1/2 and many others.

Insulin receptor and insulin-like growth factor 1 receptor (IGF-1R), which are structurally related, share conserved tyrosine residues that are phosphorylated in IR in response to insulin and in IGF-1R in response to IGF-1. Functionally, the insulin receptor regulates metabolism and IGF-1R mediates growth and differentiation. The catalytic loops within the tyrosine kinase domains of the IR/IGF-1R share the same residue sequence with a three-tyrosine motif corresponding to Tyr1158, 1162 and 1163 (for the IR) and Tyr1131, 1135 and 1136 (for the IGF-1R). It is generally believed that autophosphorylation within the activation loop proceeds in a progressive manner initiating at the second tyrosine (1162 or 1135), followed by phosphorylation at the first tyrosine (1158 or 1131), then the last (1163 or 1136), upon which the IR or IGF-1R becomes fully active.

Methods

Materials Needed But Not Provided	 Microtiter plate reader (at or near 450 nm) with software Calibrated adjustable precision pipettes Distilled or deionized water Plate washer: automated or manual (squirt bottle, manifold dispenser, etc.) Glass or plastic tubes for diluting solutions Absorbent paper towels Calibrated beakers and graduated cylinders
Procedural Notes	 When not in use, kit components should be refrigerated. All reagents should be warmed to room temperature before use. Microtiter plates should be allowed to come to room temperature before opening the foil bags. Once the desired number of strips has been removed, immediately reseal the bag and store at 2 to 8°C to maintain plate integrity. Samples should be collected in pyrogen/endotoxin-free tubes. Samples should be frozen if not analyzed shortly after collection. Avoid multiple freeze-thaw cycles of frozen samples. Thaw completely and mix well prior to analysis. When possible, avoid use of badly hemolyzed or lipemic sera. If large amounts of particulate matter are present, centrifuge or filter prior to analysis. It is recommended that all standards, controls and samples be run in duplicate. When pipetting reagents, maintain a consistent order of addition from well-to-well. This ensures equal incubation times for all wells. Do not mix or interchange different reagent lots from various kit lots. Absorbances should be read immediately, but can be read up to 2 hours after assay completion. For best results, keep plate covered in the dark. In-house controls should be run with every assay. If control values fall outside pre-established ranges, the accuracy of the assay is suspect. All residual wash liquid must be drained from the wells by efficient aspiration or by decantation followed by tapping the plate forcefully on absorbent paper. <i>Never</i> insert absorbent paper directly into the wells. Because Stabilized <i>Chromogen</i> is light sensitive, avoid prolonged exposure to light. Avoid contact between chromogen and metal, or color may develop.
Directions for Washing	 Incomplete washing will adversely affect the test outcome. All washing must be performed with the <i>Wash Buffer Concentrate (25X)</i> provided. Washing can be performed manually as follows: completely aspirate the liquid from all wells by gently lowering an aspiration tip into the bottom of each well. Take care not to scratch the inside of the well. After aspiration, fill the wells with at least 0.4 mL of diluted <i>Wash Buffer</i>. Let soak for 15 to 30 seconds, and then aspirate the liquid. Repeat as directed under Assay Procedure. After the washing procedure, the plate is inverted and tapped dry on absorbent tissue. Alternatively, the diluted <i>Wash Buffer</i> may be put into a squirt bottle. If a squirt bottle is used, flood the plate with the diluted <i>Wash Buffer</i>, completely filling all wells. After the washing procedure, the plate is inverted and tapped dry on absorbent tissue. If using an automated washer, follow the washing instructions carefully.

Recommended Formulation of Cell and Tissue Extraction Buffer: Procedure For **Extraction Of** 10 mM Tris, pH 7.4 2 mM Na₃VO₄ • Proteins 100 mM NaCl 1% Triton X-100 From Cells Or • Tissues 1 mM EDTA 10% glycerol •

- 1 mM EGTA
- 1 mM NaF
- 20 mM Na₄P₂O₇

- 0.1% SDS
- 0.5% deoxycholate

This Cell Extraction Buffer (Invitrogen, Cat. # FNN0011) needs the following reagents to be added:

- 1 mM PMSF (stock is 0.3 M in DMSO)
- Protease inhibitor cocktail (e.g., Sigma Cat. # P-2714) (reconstituted according to manufacturer's guideline). Add 500 µl per 5 mL Cell Extraction Buffer.

The Cell Extraction Buffer is stable for 2 to 3 weeks at 4°C or for up to 6 months when aliquoted (without protease inhibitors and PMSF added) and stored at -20°C. When stored frozen, the buffer should be thawed on ice. **Important:** add the protease inhibitors just before using. The stability of protease inhibitor supplemented Cell Extraction Buffer is 24 hours at 4°C. PMSF is very unstable and must be added prior to use, even if added previously.

The following protocol has been applied to several cell lines using this Cell Extraction Buffer. Researchers may optimize the cell extraction procedures that work best in their hands.

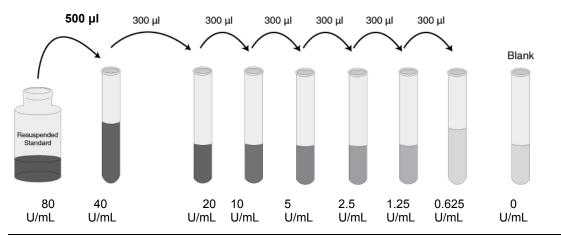
- 1. Collect cells in PBS by centrifugation (non-adherent) or scraping from culture flasks (adherent).
- 2. Wash cells twice with cold PBS.
- 3. Remove and discard the supernatant and collect the cell pellet. (At this point the cell pellet can be frozen at -80°C and lysed at a later date).
- Lyse the cell pellet in Cell Extraction Buffer for 30 minutes, on ice, with vortexing at 10 minute intervals. The volume of Cell Extraction Buffer depends on the cell number in cell pellet and expression of IR [pYpY1162/1163].
- 5. Transfer lysates to microcentrifuge tubes and centrifuge at 13,000 rpm for 10 minutes at 4°C.
- 6. Aliquot the clear lysate to clean microcentrifuge tubes. These samples are ready for assay. Lysates can be stored at -80°C. Avoid multiple freeze-thaw cycles.

Preparation of Reagents

Dilution of Standard Note: The IR [pYpY1162/1163] standard was prepared from autophosphorylated recombinant protein. One Unit of standard is equivalent to the amount of IR [pYpY1162/1163] autophosphorylated from 0.6 ng of full-length IR (β-subunit) protein.

- 1. Reconstitute *IR* [*pYpY1162/1163*] *Standard* to 80 Units/mL with *Standard Diluent Buffer*. Refer to standard vial label for instructions. Swirl or mix gently and allow to sit for 10 minutes to ensure complete reconstitution. Use standard within 1 hour of reconstitution.
- 2. Add 0.500 mL of the reconstituted standard to a tube containing 0.500 mL *Standard Diluent Buffer*. Label as 40 Units/mL IR [pYpY1162/1163].
- 3. Add 0.300 mL of *Standard Diluent Buffer* to each of 6 tubes labeled 20, 10, 5.0, 2.5, 1.25 and 0.625 Units/mL IR [pYpY1162/1163].
- 4. Make serial dilutions of the standard as described in the following dilution diagram. Mix thoroughly between steps.

Remaining reconstituted standard should be discarded or frozen in aliquots at -80°C for further use. Standard can be frozen and thawed one time only without loss of immunoreactivity.



Preparing IgG HRP

Note: Prepare within 15 minutes of usage. The *Anti-Rabbit IgG -HRP (100X)* is in 50% glycerol, which is viscous. To ensure accurate dilution, allow the *Anti-Rabbit IgG -HRP (100X)* to reach room temperature. Gently mix. Pipette the *Anti-Rabbit IgG -HRP (100X)* slowly. Remove excess concentrate solution from pipette tip by gently wiping with clean absorbent paper.

- Dilute 10 µL of this 100X concentrated solution with 1 mL of HRP Diluent for each 8-well strip used in the assay. Label as Anti-Rabbit IgG HRP Working Solution.
- 2. Return the unused Anti-Rabbit IgG HRP (100X) to the refrigerator.

# of 8-Well Strips	Volume of Anti-Rabbit IgG HRP (100X)	Volume of Diluent
2	20 µL solution	2 mL
4	40 µL solution	4 mL
6	60 µL solution	6 mL
8	80 µL solution	8 mL
10	100 µL solution	10 mL
12	120 µL solution	12 mL

Note

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Dilution of Wash Buffer 1. Allow the *Wash Buffer Concentrate (25X)* to reach room temperature and mix to ensure that any precipitated salts have redissolved. Dilute 1 volume of the *Wash Buffer Concentrate (25X)* with 24 volumes of deionized water (e.g., 50 mL may be diluted up to 1.25 liters, 100 mL may be diluted up to 2.5 liters). Label as Working Wash Buffer.

2. Store both the concentrate and the Working Wash Buffer in the refrigerator. The diluted buffer should be used within 14 days.

Assay Procedure Be sure to read the *Procedural Notes* section before carrying out the assay.

Allow all reagents to reach room temperature before use. Gently mix all liquid reagents prior to use.

Note: A standard curve must be run with each assay.

- 1. Determine the number of 8-well strips needed for the assay. Insert these in the frame(s) for current use. (Re-bag extra strips and frame. Store these in the refrigerator for future use.)
- 2. Add 100 µL of the *Standard Diluent Buffer* to the zero standard wells. Well(s) reserved for chromogen blank should be left empty.
- Add 100 μL of standards and diluted samples or controls to the appropriate microtiter wells. Samples prepared in Cell Extraction Buffer must be diluted 1:10 or greater in *Standard Diluent Buffer* (for example, 10 μL sample into 90 μL buffer). The dilution chosen should be optimized for each experimental system. Tap gently on side of plate to mix.
- 4. Cover wells with *plate cover* and incubate for **2 hours at room temperature.**
- 5. Thoroughly aspirate or decant solution from wells and discard the liquid. Wash wells 4 times. See **Directions for Washing**.
- 6. Pipette 100 μL of *IR* [*pYpY1162/1163*] *Detection Antibody* solution into each well except the chromogen blank(s). Tap gently on the side of the plate to mix.
- 7. Cover plate with *plate cover* and incubate for **1 hour at room temperature**.
- 8. Thoroughly aspirate or decant solution from wells and discard the liquid. Wash wells 4 times. See **Directions for Washing**.
- Add 100 µL Anti-Rabbit IgG HRP Working Solution to each well except the chromogen blank(s). (Prepare the working dilution as described in **Preparing** IgG-HRP).
- 10. Cover wells with the *plate cover* and incubate for **30 minutes at room temperature**.
- 11. Thoroughly aspirate or decant solution from wells and discard the liquid. Wash wells 4 times. See **Directions for Washing**.
- 12. Add 100 μL of *Stabilized Chromogen* to each well. The liquid in the wells will begin to turn blue.
- 13. Incubate for **30 minutes at room temperature and in the dark.** *Note:* **Do not cover the plate with aluminum foil or metalized mylar**. The incubation time for chromogen substrate is often determined by the microtiter plate reader used. Many plate readers have the capacity to record a maximum optical density (O.D.) of 2.0. The O.D. values should be monitored and the substrate reaction stopped before the O.D. of the positive wells exceeds the limits of the instrument. The O.D. values at 450 nm can only be read after the *Stop Solution* has been added to each well. If using a reader that records only to 2.0 O.D., stopping the assay after 20 to 25 minutes is suggested.
- 14. Add 100 μL of *Stop Solution* to each well. Tap side of plate gently to mix. The solution in the wells should change from blue to yellow.
- 15. Read the absorbance of each well at 450 nm having blanked the plate reader against a chromogen blank composed of 100 μL each of *Stabilized Chromogen* and *Stop Solution*. Read the plate within 2 hours after adding the *Stop Solution*.
- 16. Use a curve fitting software to generate the standard curve. A four parameter or five parameter algorithm provides the best standard curve fit.
- 17. Read the concentrations for unknown samples and controls from the standard curve. (Samples producing signals greater than that of the highest standard should be diluted in *Standard Diluent Buffer* and reanalyzed. Multiply value(s) obtained for sample(s) by the appropriate dilution factor to correct for the dilution in step 3.

Typical DataThe following data were obtained for the various standards over the range of 0 to
40 Units/mL IR [pYpY1162/1163].

Standard IR [pYpY1162/1163] (Units/mL)	Optical Density (450 nm)
40	2.89
20	1.48
10	0.80
5.0	0.49
2.5	0.34
1.25	0.22
0.63	0.17
0	0.15

Sensitivity The analytical sensitivity of this assay is <0.4 Units/mL of IR [pYpY1162/1163]. This was determined by adding two standard deviations to the mean O.D. obtained when the zero standard was assayed 30 times.

The sensitivity of this ELISA was compared to Western blotting using cell lysate with known quantities of IR [pYpY1162/1163]. The data presented below show that the sensitivity of the ELISA is about 4x greater than that of Western blotting. The bands shown in the Western blot data were developed using rabbit anti-IR [pYpY1162/1163], an alkaline phosphatase conjugated anti-rabbit IgG followed by chemiluminescent substrate and autoradiography.

Detection of IR [pYpY1162/1163] by ELISA vs Western Blot:

Western Blotting (95 kDa)							-	-
ELISA: OD 450 nm	0.16	0.23	0.30	0.44	0.71	1.35	2.28	3.47
Cell lysate (µg/test)	0	0.31	0.63	1.25	2.5	5	10	20

Precision 1. Intra-Assay Precision

Samples of known IR [pYpY1162/1163] concentration were assayed in replicates of 16 to determine precision within an assay.

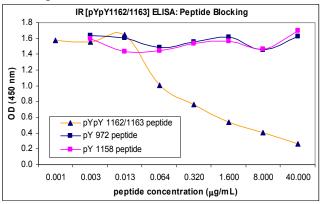
	Sample 1	Sample 2	Sample 3
Mean (U/mL)	20.56	4.64	1.42
SD	0.89	0.28	0.11
%CV 4.33 6.03 7.75			
SD = Standard Deviation CV = Coefficient of Variation			

2. Inter-Assay Precision

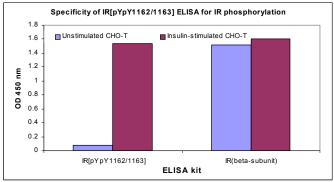
Samples were assayed 48 times in multiple assays to determine precision between assays

	Sample 1	Sample 2	Sample 3	
Mean (U/mL)	19.73	4.77	1.42	
SD	1.01	0.41	0.13	
%CV 5.12 8.60 9.15				
SD = Standard Deviation CV = Coefficient of Variation				

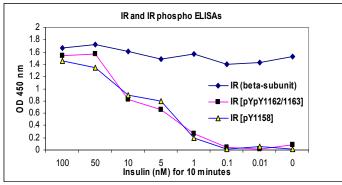
Specificity The specificity of this assay for IR phosphorylated at tyrosines 1162/1163 was confirmed by peptide competition. Phosphorylated IR was quantitated in the assay as usual except that the detection antibody was preincubated with IR-derived peptides at a concentration of 0.001- 40 μ g/mL. The data presented below show that only the peptide corresponding to the region surrounding tyrosines 1162/1163, containing the phospho-tyrosines, could block the ELISA signal. The peptides containing phosphorylated tyrosine at positions 972 and 1158 did not block the signal.



IR of natural or recombinant (β -subunit) origin is reactive in this assay. This kit detects phosphorylated IR in insulin-stimulated IR transfected CHO-T cells (CHO-T/IR) and does not detect non-phosphorylated IR in unstimulated cells.



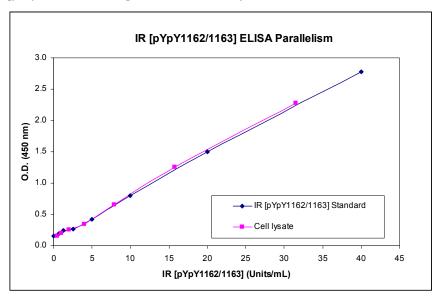
IR [pYpY1162/1163] phosphorylation in CHO-T/IR cells is dependent on levels of insulin stimulation. Cells (approximately 90% confluent) were treated with insulin at varying concentrations (0-100 nM) for 10 minutes, lysed and quantitated in parallel for IR content (both β -subunit and pYpY1162/1163). The amount of IR (β -subunit) remains constant, while the level of phosphorylation at tyrosines 1162/1163 decreases with diminishing insulin dosage. Phosphorylation at IR [pY1158], as measured in another Invitrogen ELISA (Cat. # KHR9121), shows a similar pattern.



Linearity of Extract Buffer was spiked with IR [pYpY1162/1163] and serially diluted in Standard Diluent Buffer over the range of the assay. Linear regression analysis of samples versus the expected concentration yielded a correlation coefficient of 0.99.

	Cell Lysate			
Dilution	Measured (Units/mL)	Expected (Units/mL)	% Expected	
Neat	31.5	-	-	
1/2	16.8	15.8	106%	
1/4	8.2	7.9	104%	
1/8	4.1	3.9	105%	
1/16	2.3	2.0	115%	

- **Recovery** The recovery of IR [pYpY1162/1163] added to 100 μg/mL of a Jurkat cell lysate in Cell Extract Buffer (followed by a 1:10 dilution in *Standard Diluent Buffer*) averaged 104%.
- **Parallelism** Natural IR [pYpY1162/1163] from lysate of insulin-stimulated CHO-T/IR cells was serially diluted in *Standard Diluent Buffer*. The optical density of each dilution was plotted against the human IR [pYpY1162/1163] standard curve. Parallelism was demonstrated by the figure below and indicated that the Standard accurately reflects IR [pYpY1162/1163] content in samples.



Limitations of the Procedure

Do not extrapolate the standard curve beyond top standard point; the dose-response is non-linear in this region and accuracy is difficult to obtain. Dilute all samples above the highest standard point with *Standard Diluent Buffer*, reanalyze these and multiply results by the appropriate dilution factor.

The influence of various extraction buffers has not been thoroughly investigated. The rate of degradation of native IR [pYpY1162/1163] in various matrices has not been investigated.

Troubleshooting Guide

Standard curve wells develop, but sample wells	<i>Cause:</i> Improper sample preparation. <i>Solution 1:</i> Make sure to prepare cell extracts in the protease inhibitor- supplemented Cell Extraction Buffer recommended in the protocol booklet. Other buffer formulations have not been evaluated.
produce weak or no signal.	<i>Cause:</i> Samples contain materials that interfere with the assay. <i>Solution 1:</i> The Cell Extraction Buffer recommended in the protocol booklet contains SDS. This detergent can potentially interfere with the immunoassay. For this reason, we recommend that all samples be diluted by a factor of at least 1:10 using the Standard Diluent Buffer provided in the kit. This buffer also contains blocking proteins that will reduce background signal.
	<i>Cause:</i> The concentration of the target analyte is too dilute. <i>Solution 1:</i> When preparing the cell extracts, increase the number of cells per volume of protease inhibitor supplemented Cell Extraction Buffer. Ideally, the concentration of protein in each cell extract, as determined by the Quant-iT ^{m} protein assay kit, will be between 1 and 10 mg/mL (Method 1) or 1 and 5 mg/mL (Method 2). It is recommended that 5-10 µg of total cellular protein as a starting point be loaded into each well. <i>Solution 2:</i> Optimize the stimulation procedure and time.
	Cause: A sample treatment step was not performed. Solution 1: Certain analytes may require a sample treatment step to improve performance with Invitrogen phosphoELISA [™] kits. Please see the analyte- specific protocol booklet for information on sample treatment procedures.
	<i>Cause:</i> Samples deteriorated during storage. <i>Solution 1:</i> Make sure that the Cell Extraction Buffer is supplemented with phosphatase inhibitors and protease inhibitors just prior to use. <i>Solution 2:</i> All samples should be stored frozen at –80°C. <i>Solution 3:</i> Samples should be subjected to only one freeze-thaw cycle. <i>Solution 4:</i> Some proteins can be lost by adsorption when stored in containers made of polystyrene or certain kinds of glass. Polypropylene tubes are best for storing samples.
Sample wells develop, but standard wells produce weak or no	<i>Cause:</i> Improper dilution of standards. <i>Solution 1:</i> Check reconstitution volume of standard. <i>Solution 2:</i> Standard curves are generated by serially diluting the reconstituted standard. Check the serial dilution method. <i>Solution 3:</i> Standards should be used within an hour of reconstitution and serial dilution.
signal.	<i>Cause:</i> Improper storage of standards. <i>Solution 1:</i> Standards are provided as lyophilized powders that should be stored at 2–8°C. Once reconstituted, standard should be stored at –80°C.

Neither the standard curve wells nor the sample wells develop.	<i>Cause:</i> Insufficient horseradish peroxidase (HRP)-conjugated secondary antibody activity. <i>Solution 1:</i> Check the dilution of the HRP secondary antibody. <i>Solution 2:</i> The HRP secondary antibody must be freshly diluted for each assay. <i>Solution 3:</i> The HRP secondary antibody must be stored at 2–8°C. <i>Solution 4:</i> Sodium azide is an irreversible inhibitor of horseradish peroxidase enzyme activity. Make sure to dilute the HRP secondary antibody in the correct buffer. A quick test can be performed to determine if the HRP secondary antibody is active. Into a clean test tube, dispense 200 µl of the TMB substrate solution provided in the kit. This TMB substrate solution should be clear to slightly blue-green tinted. Next, pipette 2 µl of the HRP secondary antibody. The color of the TMB will change to an intense aqua blue instantaneously if the HRP has retained its enzyme activity.
	 <i>Cause:</i> Insufficient Detector Antibody. <i>Solution 1:</i> The Detector Antibody must be stored at 2–8°C. <i>Solution 2:</i> Improper dilution of Detector Antibody. <i>Cause:</i> TMB solution lost activity. <i>Solution 1:</i> The TMB solution should be clear before it is dispensed into the wells of the microtiter plate. An intense aqua blue color indicates that the product is contaminated. Please contact Technical Support if this problem is noted. To avoid contamination, we recommend that the quantity required for an assay be dispensed into a disposable trough for pipetting. Any TMB solution left in the trough should be discarded. <i>Solution 2:</i> Avoid contact of the TMB solution with items containing metal ions.
Standard curves are not developing consistently in between different runs.	<i>Cause:</i> Improper dilution of Secondary antibody. <i>Solution 1:</i> The Anti-Rabbit IgG HRP (100X) is in 50% glycerol. This solution is viscous. To ensure accurate dilution, allow Anti-Rabbit IgG HRP (100X) to reach room temperature. Gently mix. Pipette concentrate slowly. Remove excess concentrate solution from pipette tip by gently wiping with clean absorbent paper. <i>Solution 2:</i> Check plate washing technique. Results can be effected if some liquid remains after aspiration.

Technical Support



E-mail: eurotech@invitrogen.com

Reference	s
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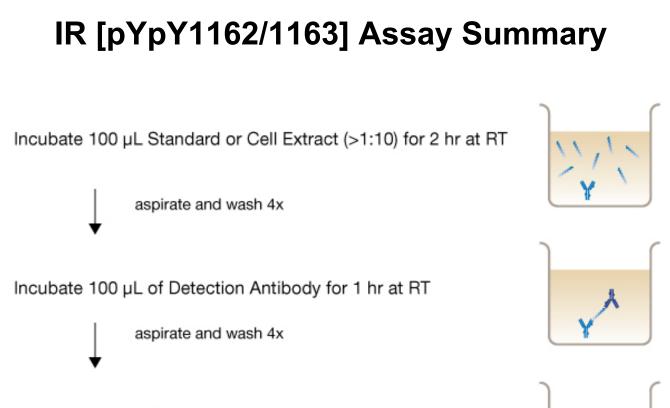
Explanation of symbols			
Symbol	Description	Symbol	Description
REF	Catalogue Number	LOT	Batch code
RUO	Research Use Only	IVD	In vitro diagnostic medical device
\mathbf{X}	Use by		Temperature limitation
***	Manufacturer	EC REP	European Community authorised representative
[-]	Without, does not contain	[+]	With, contains
from Light	Protect from light	Â	Consult accompanying documents
[]i	Directs the user to consult instructions for use (IFU), accompanying the product.		

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aspirate and wash 4x

Add 100 µL Stop Solution and read at 450 nm

Total time: 4 hr

Incubate 100 µL of Stabilized Chromogen for 30 min at RT

