



# In-situ hybridization using GeneDetect™ oligonucleotide probes

## DIG-labeled probe, paraffin tissue sections, Detection by AP (alkaline phosphatase), FITC or rhodamine.

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### Introduction:

Of the common non-radioactive methods used with in-situ hybridization, digoxigenin (DIG)-based detection remains one of the most sensitive. Because digoxigenin is synthesized solely in plants, non-specific background staining in animal and human tissues is usually very low. The method described below allows for reproducible detection of low abundance target mRNAs in paraffin sections. Tissue type and the nature of prior fixation determine which permeabilization steps may be required or left out of the protocol.

### Other Materials/Kits required:

For best results with GeneDetect™ oligonucleotide probes and in-situ hybridization we recommend using reagents supplied by Roche Molecular Biochemicals, <http://biochem.roche.com>.

1. DIG Oligonucleotide Tailing Kit, Cat#1417231
2. Anti-DIG-AP Fab fragments, Cat#1093274

3. Anti-DIG-fluorescein (FITC) Fab fragments, Cat#1207741
4. Anti-DIG-rhodamine Fab fragments, Cat#1207750
5. NBT/BCIP Ready-to-use tablets, Cat#1697471 or NBT/BCIP *FAST*™ tablets from Sigma.

### Abbreviations/Codes:

RT = room temp  
4°C, standard refrigerator  
-20°C, non-cycling standard freezer (i.e. should not be auto-defrost type)  
-70°C, ultra low temperature freezer for long term storage of biological materials  
2.5pmol of a 48mer oligonucleotide ~ 35ng  
x = times, as in wash 2 times  
X = times (concentration), as in 2 X SSC

**Simplifying the procedure: Optional steps.** We have tried various combinations of this procedure. Not performing the optional steps can “fast track” and simplify the procedure. You should first assess the effects of leaving out any ONE of the optional steps by performing parallel in-situ hybridization experiments comparing mRNA signal intensity in the standard and simplified procedures.

### Protocol Summary (4 steps)

- A. **Preparation of your DIG-labeled GeneDetect™ oligonucleotide probe** (2-4 hours)
- B. **Tissue Preparation** (1.5 hours)  
Dewax and rehydrate the tissue specimens.  
Post Fix.
- C. **In-situ hybridization of DIG-labeled GeneDetect™ oligonucleotide probe to tissue**  
Acetylation (optional\*, 25 mins).  
Permeabilize (optional\*).
  1. None (10 mins)
  2. Proteinase K (40 mins)

3. Pepsin (40 mins)
  4. Triton X-100 (30 mins)
- Prehybridization (optional\*, 2.5 hours).  
Hybridization with probe (18-40 hours).  
Post-hybridization washes (2 hours).

**D. Immunological detection of DIG** (8 hours to overnight)

1. Indirect detection of DIG-labeled oligonucleotides using an anti-DIG antibody conjugated to alkaline phosphatase.
2. Direct fluorescent detection of DIG-labeled oligonucleotides using anti-DIG antibodies conjugated to either fluorescein (FITC) or rhodamine.

*The protocol can be completed over 2-3 days when solutions are pre-made. \*Optional Steps, see previous page for explanation.*

**A. Preparation of your DIG-labeled GeneDetect™ oligonucleotide probe:**

Various methods can be used to DIG-label your probe. The aim of this step is to tail your probe at the 3' end with multiple nucleotides (~10-100, with an average tail of 50 nucleotides leading to the incorporation of ~5 DIG-labeled dUTP nucleotides into the tail). Note that tailing is quite different from 3'-end labeling which only adds a single DIG-labeled nucleotide to the probe. For in-situ hybridization 3' tailing is preferred. This tail which includes multiple DIG-labeled nucleotides can be detected with an anti-DIG antibody after in-situ hybridization is complete thus localizing the bound probe and therefore the target mRNA within the tissue section. We suggest that a commercially available kit be used to perform this reaction. For example Roche Molecular Biochemicals (<http://biochem.roche.com>, DIG Oligonucleotide Tailing Kit). Follow the manufacturers specific instructions.

Briefly, the core of the 3' tailing kit involves adding the following together:

- GeneDetect™ oligonucleotide probe
- + DIG-labeled dUTP
- + unlabeled dATP
- + Terminal transferase (TdT)
- + TdT reaction buffer.
- + BSA

Total reaction volume ~ 20µl. If you choose to use the DIG Oligonucleotide Tailing Kit from Roche Diagnostics, it is necessary to add 100pmol of probe in 10µl to the tailing reaction.

Hint: Therefore when you receive your lyophilized GeneDetect™ probe (2500pmoles) reconstitute using

250µl sterile, dH<sub>2</sub>O (distilled water). While reconstitution in TE buffer is probably safer for the probe it may also unnecessarily interfere with the DIG-labeling procedure.

The tailing reaction is allowed to proceed for 15-30 mins at 37°C and is stopped by adding 2µl of 0.2M EDTA/glycogen solution (to prepare add 1 µl of 20mg glycogen/ml dH<sub>2</sub>O to 200µl 0.2M EDTA pH 8.0). A typical labeling reaction should provide 100pmol or 1400ng of DIG-labeled GeneDetect™ probe (48mer). **With in-situ hybridization it will be necessary to purify your DIG-labeled probe.** A standard ethanol/LiCl precipitation is suggested.

Precipitate the labeled GeneDetect™ oligonucleotide. To the 22-23µl reaction mixture add:

2.5µl 4M LiCl and 75µl pre-chilled (-20°C) 100% ETOH. Mix well.

Let the precipitate form for at least 30 mins at -70°C or 2 hrs at -20°C

Centrifuge tubes at 13,000 x g, for 15 mins at 4°C

Discard the supernatant

Wash pellet with 50µl ice-cold 70% (v/v) ETOH

Centrifuge tubes at 13,000 x g, for 5 min at 4°C

Discard the supernatant

Dry the pellet under vacuum (we use an Eppendorf 5301 concentrator).

Reconstitute the pellet in sterile dH<sub>2</sub>O

Commercial kits provide methods for yield determination allowing for non-variable, reproducibility of labeling results between experiments. An advantage of DIG-versus <sup>35</sup>S-labeled probes beyond the obvious, that they are non-radioactive, is that they are stable for up to 1 year after labeling when stored at -20°C and are equally as sensitive as <sup>35</sup>S-labeled probes in detecting target mRNAs in tissue sections.

**B. Tissue Preparation:**

**ONCE TISSUE SECTIONS ARE REHYDRATED DO NOT LET SECTIONS DRY OUT AT ANY STEP IN THE ENTIRE IN-SITU HYBRIDIZATION PROCEDURE.**

Fresh tissue should be removed and immediately fixed overnight, preferably in 4% paraformaldehyde (4%

PFA in 0.1M PB, phosphate buffer) or fixed as soon as possible to prevent RNA degradation. Prolonged fixation beyond 24 hrs should be avoided. Once the tissue is fixed, standard laboratory techniques for dehydrating and embedding tissue in paraffin can be followed. Paraffin-embedded tissue is best-kept long term at 4°C in a dry environment. For in-situ hybridization, 7-14µm sections should be cut and mounted onto poly-L-lysine coated slides (or equivalent) and air dried (overnight if necessary).

All of the following steps in the protocol are performed by incubating slides containing tissue specimens (held in a slide rack or Coplin jar) in the indicated solutions. All solutions are made up fresh and are used only once unless indicated otherwise. All solutions are presumed to be at RT unless otherwise indicated.

Dewax and rehydrate the tissue specimens:

Remove your slides (containing your tissue sections) from 4°C storage (dry atmosphere).

Allow to warm to RT

Dewax sections by 3 x 2 mins washes using fresh xylene each time.

Rehydrate by performing the following steps with your slides

100% ethanol (ETOH) 2 x 2 mins washes  
95% ETOH 1 x 5 mins  
70% ETOH 1x 5 mins  
50% ETOH 1x 5 mins  
2 quick washes in DEPC-treated dH<sub>2</sub>O  
2 x 5 mins DEPC-PBS

Post-Fix:

Post-fix sections for 10-15 mins with 4% PFA in 0.1M PB solution (made up within the last 7 days). At this point your tissue RNA is “reasonably” safe from RNases and since GeneDetect™ oligonucleotide probes are resistant to degradation by RNases we have found that the following steps only require that sterile, autoclaved solutions be used.

Wash slides 2 x 5 mins with PBS.

### **C. In-situ hybridization of DIG-labeled GeneDetect™ oligonucleotide probe to tissue:**

Acetylation:

Optional but strongly suggested. This step blocks polar and charged groups on tissue sections which will cause non-specific binding of some probes. If this step is used it is important for its effectiveness that the acetic anhydride only be added to the triethanolamine buffer (0.1M TEA, pH 8.0) immediately before slides are incubated as the half-life of acetic anhydride in solution is very short.

Transfer slides to fresh 0.1 M TEA buffer. Add acetic anhydride to a concentration of 0.25% (v/v). Mix quickly and incubate slides for 5 mins. Add additional acetic anhydride to reach a final concentration of 0.5% (v/v) and incubate for another 5 mins.

Wash slides 1 x 3 mins in 2 X SSC

Permeabilize:

We suggest trying all four alternatives “individually” on different sections in one particular experiment to determine what works best with your particular situation/tissue/fixation method. Hint: Use our control GeneDetect™ polyd(T) probe to optimize your protocol.

1. No permeabilization.

2. Permeabilize sections for 30 mins at 37°C with DEPC-treated TE buffer containing 5-20µg/ml RNase-free Proteinase K. Follow by washing for 30 sec to 1 min in PBS + 2mg/ml glycine. Titration of the enzyme concentration is usually required here to achieve the best results for any given tissue. In general the aim is to use the highest concentration that still gives good cellular morphology and doesn't lead to the tissue section falling off the slide! It is a good idea to stick to one source of Proteinase K as different products can differ markedly in their enzymatic activity. Glycine stops proteolysis.

3. An alternative permeabilization protocol that we have found to give excellent results for formalin-fixed, paraffin-embedded tissue sections involves incubating slides for 20-30 mins at 37°C with 100-500µg/ml pepsin in 0.2M HCl.

4. Another alternative permeabilization step involves incubating sections with Triton X-100 detergent. 1 x 15 mins wash in PBS containing 0.3% Triton X-100.

At the end of all permeabilization options,

Wash slides 2 x 5 mins PBS.

Prehybridization:

Optional. Get excess buffer solutions off the tissue sections. You will want to wipe around the sections with a cloth. This keeps the prehybridization buffer from “leaving” the tissue section and running off the slide. Do not wipe the sections. DO NOT let the tissue sections dry out. Carefully overlay each section with well mixed prehybridization buffer (recipe below).

Hint: Plastic tupperware containers can be used as good in-situ hybridization chambers. We balance slides on an overturned plastic 1.5ml microfuge tube rack placed into a tupperware container which is then partially filled

with dH<sub>2</sub>O to keep the chamber humid and the sections from drying out.

Cover each section with a piece of Parafilm about the same size as the tissue. Make sure you don't get any air bubbles trapped between the section and the Parafilm. Note: if prehybridization buffer has been made earlier (and therefore is being stored at -20°C) you should remove it from the -20°C freezer and allow it to heat to 37°C in your incubator/oven for about 30 mins before adding to sections.

The volume of pre-hybridization buffer added to each tissue section will depend upon the size of the tissue. 50µl is a normal volume for a tissue section 2cm x 2cm. With a large block of human tissue you may require a higher volume.

Pre-Hybridization Buffer (prepare in 50 ml Falcon tube with screw cap, store @ -20°C)

To make 20 mls.

20 X SSC	4 ml
Dextran sulphate	4 g
Formamide (deionized)	10 ml

Add these all together then sonicate (with cap on) for about 3-4 hrs.

Then add aliquots of the following (stored at -20°C)

PolyA (10mg/ml)	0.5 ml
ssDNA (10mg/ml)	0.5 ml
tRNA (10mg/ml)	0.5 ml
DTT (of 1M solution)	2 ml
50 x Denhardt's	0.2 ml

Mix VERY well before use. Pre-hybridization buffer can be pre-made and stored long-term at -20°C. Bring to 37°C before use.

Incubate your slides in the sealed humid chamber (i.e. water filled tupperware container) for 2 hrs at 37°C by putting the lid on the container and carefully placing it in an oven set at 37°C.

While your sections are undergoing the pre-hybridization step you can add your DIG-labeled GeneDetect™ oligonucleotide probe to the pre-hybridization buffer (brought to 37°C) to make hybridization buffer. The optimal amount of probe you will need to add will require a bit of trial-and-error, but is usually within the range 100-1000ng/ml of hybridization buffer with 200ng/ml a good starting concentration. Mix well BY HAND to ensure even probe dispersal. Since a typical 3' labeling reaction produces ~ 100 pmoles (1400ng of a 48mer DIG-labeled probe) you should be able to make 7mls of hybridization buffer if you probe is diluted to 200ng/ml.

Hybridization with probe:

At the end of 2 hrs remove the Parafilm from tissue sections using forceps/tweezers before tipping off the pre-hybridization buffer and putting the slides into 2 X SSC for 5 mins.

Get excess buffer solutions off the tissue sections. You will want to wipe around the sections with a cloth. This keeps the hybridization buffer from "leaving" the tissue section and running off the slide and this becomes very important in an overnight incubation step. It is important not to let tissue sections dry out. Do not wipe the sections off the slides though! Now overlay each section with hybridization buffer. Cover each section with a piece of Parafilm about the same size as the tissue as before. Make sure you don't get any air bubbles trapped between the section and the Parafilm. Carefully put your slides into a humid chamber for overnight incubation at 37°C (approx 18 hrs). Hybridization can be left for up to 40 hrs to increase signal intensity as long as tissue sections DO NOT dry out.

Hint: While hybridization buffer is generally quite viscous at RT, when it is heated to 37°C it loses some of its viscosity and becomes "runny". Therefore another step you should take to stop hybridization buffer running off the section with the overnight hybridization is to carefully ensure all of your slides are kept level. If your slides dry out they have a higher chance of having high background staining. Further, not having your slides level could lead to uneven hybridization of the probe across the tissue section.

Post-hybridization washes:

For washes, prepare 0.5 and 1 X SSC solutions from 20 X SSC and add DTT on the day of use. (Note: make sure stock 0.5 and 1 x SSC solution are at 55°C before incubating slides.) Note: 1.2g DTT into 800mls SSC = 10mM

At the end of the overnight hybridization remove the parafilm from sections by using forceps/tweezers before tipping off the hybridization buffer and putting the slides into wash solutions. Using a shaking water bath at 55°C, give slides the following washes

Quick wash	1 X SSC (10mM DTT) RT
2 x 15 mins	1 X SSC (10mM DTT) 55C
2 x 15 mins	0.5XSSC(10mM DTT) 55C
1 x 10 mins	0.5XSSC (10mM DTT) RT

#### D. Immunological detection of DIG:

DIG-labeled oligonucleotides can be detected after hybridization by an anti-DIG antibody conjugated to either (1) the enzyme alkaline phosphatase which catalyzes a chemical reaction for indirect probe detection, OR (2) either of the fluorescent molecules

fluorescein (FITC) or rhodamine for direct fluorescence detection.

1. Indirect detection of DIG-labeled oligonucleotides using an anti-DIG antibody conjugated to alkaline phosphatase:

An antibody (anti-Digoxigenin-AP, Fab fragments, made in sheep) for the detection of DIG-labeled compounds is available from Roche Molecular Biochemicals.

Hint: Small antibody aggregates in the Anti-DIG-AP may lead to higher background staining. It is therefore suggested that the vial be centrifuged for 5 min at 13,000 rpm before its first use. Thereafter, it is sufficient to centrifuge for 1 min directly before dilution.

Transfer slides to Tris-buffered saline (TBS, 100mM Tris HCl, 150mM NaCl, pH 7.5)

Wash sections 3 x 5 mins in TBS.

Optional. Cover sections for 30 mins with blocking solution (TBS + 0.1% Triton X-100 + 1% normal sheep serum from Sigma or 1% proprietary blocking agent from Roche).

Hint: The use of fetal calf serum as a blocking agent is not advised due to its endogenous alkaline phosphatase activity.

Pour off blocking solution and incubate sections for 4 hrs minimum at RT with anti-DIG antibody diluted 1:100 to 1:1000 (1:200 initial recommended dilution) in TBS + 0.1% Triton X-100 + 1% normal sheep serum from Sigma or 1% proprietary blocking agent from Roche).

Wash sections 3 x 5 mins in TBS.

Dissolve one NBT/BCIP ready to use tablet from Roche in 10mls of dH<sub>2</sub>O to make 10 mls of staining solution or alternatively prepare yourself (below).

10mls of NBT/BCIP staining solution:

0.4 mg/ml NBT (Nitro blue tetrazolium chloride)

0.19 mg/ml BCIP (5-Bromo-4-chloro-3-indolyl-phosphate, toluidine salt)

100mM Tris buffer, pH 9.5

50mM MgSO<sub>4</sub>

Optional: Add 10µl of stock 1M levamisole solution per 10 mls staining solution (1mM levamisole). Although

unlikely, some endogenous phosphatase activity may remain at this stage of the protocol and cause increased non-specific binding. The option is to add levamisole (Sigma). For convenience prepare a 1M stock solution in dH<sub>2</sub>O which will be stable at 4°C for several weeks. If endogenous phosphatase activity is particularly high (found in certain tissue types) increase the amount of levamisole added to the detection solution.

Incubate sections in a Coplin jar or similar with the staining solution. A blue precipitate will form. The development time will depend on numerous factors but is usually between several minutes for high abundance to several hours or even overnight for low level mRNAs.

Stop the reaction by rinsing the slides several times in tap water.

Finally rinse in dH<sub>2</sub>O and mount slides with any water-soluble mounting medium (do not use xylene-based mounting medium) or optionally counter stain the sections.

2. Direct fluorescent detection of DIG-labeled oligonucleotides using anti-DIG antibodies conjugated to either fluorescein (FITC) or rhodamine:

Antibodies (anti-Digoxigenin-fluorescein (FITC) Fab fragments and anti-Digoxigenin-rhodamine Fab fragments both made in sheep) can be used for the direct fluorescent detection of DIG-labeled compounds. Both are available from Roche Molecular Biochemicals.

Hint: Small antibody aggregates in the Anti-DIG-FITC/rhodamine solutions may lead to higher background staining. It is therefore suggested that the vial(s) be centrifuged for 5 mins at 13,000 rpm before first use. Thereafter, it is sufficient to centrifuge each for 1 min directly before dilution.

Transfer slides to Tris-buffered saline (TBS, 100mM Tris HCl, 150mM NaCl, pH 7.5)

Wash sections 3 x 5 mins in TBS.

Optional. Cover sections for 30 mins with blocking solution (TBS + 0.1% Triton X-100 + 1% normal sheep serum from Sigma or 1% proprietary blocking agent from Roche).

Pour off blocking solution and incubate sections for 4 hrs minimum at RT with anti-DIG (FITC or rhodamine) diluted 1:50 to 1:500 (1:100 initial recommended dilution) in TBS + 0.1% Triton X-100 + 1% normal sheep serum from Sigma or 1% proprietary blocking agent from Roche).

Wash sections 3 x 5 mins in TBS.

Finally rinse sections several times in dH<sub>2</sub>O to remove salts and then mount sections using a medium containing an anti-fading agent (for example, Vectashield, Vector Labs) and evaluate staining using a fluorescent microscope.

FITC, Green: Excitation  $\lambda_{max}$ [nm]: 494, Emission  $\lambda_{max}$ [nm]: 523 (pH 8.0)

Rhodamine, Red: Excitation  $\lambda_{max}$ [nm]: 555, Emission  $\lambda_{max}$ [nm]: 580 (pH 8.0)

### Controls:

Of course the most important part of any experimental procedure is the inclusion of controls. However often with insitu hybridization experiments controls not used properly, if at all. In carrying out an insitu hybridization experiment one has to be confident that the hybridization reaction is specific and that the probe is in fact binding selectively to the target mRNA sequence and not to other components of the cell or other closely related mRNA sequences. In addition if no staining is observed with the probe does this mean that there really is no expression of that mRNA in the tissue or does it mean that there may be a problem with tissue preparation or the tissue itself or your technique?

If the correct controls are included in the experiment we can, with high certainty, answer these questions. Note that the polyd(T) probe is included with all orders and that the nonsense probe and pan-species actin probe are contained within our Control Probes product. Both sense and antisense probes are sent when you order a probe from us in amounts that allow for 10X competition studies to be performed as mentioned below. RNase enzyme should be purchased from a trusted supplier.

### *Controls for tissue mRNA quality and the efficacy of your protocol.*

If the quality of your tissue is poor and/or your RNA is degraded it will be very hard to get good results with in situ hybridization. There are however a number of controls you can add to your experiment to verify the status of your tissue and mRNA within the tissue. If you are using fresh tissue and these controls are negative, then this suggests a problem with your technique or protocol.

### *Poly d(T) probe.*

The poly d(T) probe we supply will detect total mRNA polyA tails. If a very weak signal is obtained using this probe then it is likely your

tissue RNA is degraded. The chance of detecting a specific mRNA in this tissue is therefore unlikely.

### *Probes against house keeping sequences.*

Some genes are always expressed constitutively such as Actin or beta-tubulin. We offer probes to detect these mRNAs. A low signal once again suggests tissue RNA degradation.

### *Positive control.*

Perform in situ hybridization using the correct oligonucleotide probe on a fresh, positive control tissue known to have the sequence of interest (not always possible). If you detect no signal then this suggests the problem exists within your technique or protocol.

### *Specificity controls.*

#### *Determine that your probe is only binding to RNA.*

When probing for mRNAs one can determine that the binding is specific to RNA by digesting the tissue with RNases prior to hybridization with the oligonucleotide probe. The absence of binding after RNase treatment indicates that binding was indeed to RNA within the tissue.

RNase solution (200ml) can be made up as follows.

0.8ml 10mg/ml RNase (Sigma)  
4ml 1M Tris buffer (pH 7.5)  
0.4ml 0.5M EDTA  
Add dH<sub>2</sub>O to 200ml (i.e. add 194.8ml).

Sections should be incubated in RNase solution for 1 hour at 37°C immediately after the 2 x 5 mins washes with PBS that follow post fixation. These RNase-treated sections should be compared with sections also incubated with RNase solution (minus the RNase) for 1 hr at 37°C. Following RNase treatment, sections should be washed 2 x 5 min in PBS before entering the main protocol again.

### *Specific versus non-specific binding.*

The first control involves hybridization of the tissue with both labeled sense and antisense probes in parallel. The antisense probe in theory detects both the target mRNA and any non-specific targets it can bind to due to the chemical properties of the probe (but not due to the probe sequence). The sense control

probe gives a measure of non-specific probe binding only due to the chemical properties of the probe. In essence if your sense probe detects nothing, then you can be sure that any signal detected by your antisense probe is due to sequence-specific binding to mRNA and not due to binding to other targets within the cell.

Competition studies with labeled and excess unlabeled probes can also help distinguish between specific versus non-specific binding. This is because by definition specific binding is saturable (i.e. there are finite target mRNA molecules to which the probe can bind) while non-specific binding is not (there are infinite non-specific targets). Therefore excess unlabeled probe can displace (by competition for binding sites) the specific binding of the labeled probe (i.e. to the target mRNA) but not non-specific binding of the labeled probe.

We recommend co-hybridizing tissue with:

1. Excess 10 X molar concentration of unlabeled antisense probe plus the usual concentration of labeled antisense probe
2. Excess 10 X molar concentration of unlabeled nonsense probe plus the usual concentration of labeled antisense probe

The nonsense probe should preferably have a similar CG content, a similar length and have no homology to the sequence of interest.

It is important to note however that competition studies do not verify the identity of the mRNA to which the labeled probe is binding since both the labeled and unlabeled probes have the same sequence.

*Determine that your probe is binding to the correct target sequence.*

The best way to ensure that your probe is binding to the correct target sequence is by choosing a correct probe sequence from the start and having high stringency hybridization and wash conditions in your experiment.

In summary we recommend that the following controls are performed in parallel with your in situ experiments.

1. polyd(T) probe is hybridized to sections. What is the quality of the mRNA in your tissue sample?
2. RNase treatment of sections before labeled antisense probe hybridized. Is probe binding to mRNA?

3. Hybridize in parallel labeled sense and labeled antisense probes. Is the probe binding to the tissue in a sequence-specific fashion?
4. Can sequence-specific binding be displaced? Hybridize labeled antisense probe in presence of
  - a) 10X unlabeled antisense probe and separately in presence of
  - b) 10X unlabeled nonsense probe.

#### **Solutions and chemicals required:**

If the purchases of salts, buffers and the like are made from Sigma or another similar supplier we suggest you use Molecular Biology grade products.

All glassware, plasticware, pipette tips etc should be autoclaved.

dH<sub>2</sub>O (distilled water). Must also be autoclaved.

Standard buffers. SSC, PBS, TE, PB (Any protocols book will have the recipes for these).

Diethylpyrocarbonate (DEPC)-treated dH<sub>2</sub>O, TE buffer and PBS. To DEPC treat solutions, add 0.2 ml DEPC to 100 ml of the solution to be treated. Shake vigorously to get the DEPC into solution. Autoclave. Prepare in a fume hood and wear gloves.

0.1 M triethanolamine buffer (0.1M TEA in dH<sub>2</sub>O, pH 8.0). TEA purchased from Sigma. Add 18.57g triethanolamine.Cl to 900ml dH<sub>2</sub>O. Dissolve and adjust pH to 8.0 using NaOH. Adjust to 1 litre with dH<sub>2</sub>O. Use the same day.

0.2M EDTA + glycogen solution. Add 1µl of 20mg glycogen/ml dH<sub>2</sub>O to 200µl 0.2M EDTA (pH 8.0)

0.1 and 0.2M phosphate buffer (PB, store on lab shelves @ RT)

20 x SSC (stock, store on lab shelves @ RT)

100% Ethanol

Parafilm

Hybridization chambers (fill with dH<sub>2</sub>O before overnight hybridization)

Oven @ 37°C, for overnight hybridizations.

Shaking water bath @ 55°C

Wash containers that will hold slides and SSC wash solutions (Plastic Coplin Jars work well)

Dextran sulphate (Sigma, 50g D8906, store @ 4°C)

Deionized formamide (use as bought, Sigma, 100ml F9037, store 10ml aliquots @ -20°C)

PolyA (Sigma, 25mg, P9403, dissolve to 10 mg/ml in dH<sub>2</sub>O, store 0.5ml aliquots @ -20°C)

ssDNA (Life Technologies, 10 mg/ml, use as bought, store 0.5ml aliquots @ -20°C)

tRNA (use as bought Sigma, 10 mg/ml, store 0.5ml aliquots @ -20°C)

DTT (various sources, 154mg/ml in dH<sub>2</sub>O sterile = 1M solution, store as 1ml aliquots @ -20°C)

50XDenhardtts (use as bought, if required dissolve with dH<sub>2</sub>O, store 0.2ml aliquots @ -20°C)

BSA (dilute stock 20mg/ml Molecular Biological grade from Sigma Cat#B8894 to 2 mg/ml using dH<sub>2</sub>O, store @ -20°C)

4% PFA (in 0.1M PB, phosphate buffer): 8g PFA dissolved in 100ml of H<sub>2</sub>O. Heat to partially dissolve then add 1M NaOH dropwise until solution clears. Add 100ml of 0.2M PB, mix. Filter. Cool to 4°C before use. Store at 4°C. Prolonged storage (>1 week) may require that pH is checked. pH should be around 7.4, DO NOT autoclave.

0.5XSSC and 1XSSC buffers + 10mM DTT for washes. (Make 0.5XSSC and 1XSSC buffers by diluting 20XSSC with dH<sub>2</sub>O. Into 200ml SSC (0.5X or 1X) buffer add 0.31g DTT powder and mix well. We usually make up 200ml 1XSSC (0.31g DTT) and 200 ml 0.5XSSC (0.31g DTT) on the day of washes because DTT inactivates in solution.

Questions? For our customers we guarantee a response within 24 hrs.

Email us at [Scientific@GeneDetect.com](mailto:Scientific@GeneDetect.com)

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