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Colorimetric in situ hybridisation

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ABSTRACT

The protocol for performing colorimetric *in situ* hybridisation in zebrafish embryos and larvae in the Wilson lab.

IMAGE ATTRIBUTION

mab2112 expression at 1 dpf, taken from ZFIN.

MATERIALS

Buffers and solutions

Hybridization buffer (50ml):
Formamide 25 ml
20x SSC 12.5 ml
Torula RNA 500 µl
100mg/ml Heparin 25 µl
20% Tween-20 250 µl
1M citric acid 460 µl

Store at -20°C

MiliQ H₂0

MaBl (100ml):

Boehringer Blocking Reagent (Roche) 2 g 100mM Maleic Acid buffer (pH7.5) 100 ml Autoclave and store at -20°C

11.27 ml

20x SSC:

Final concentration

3M NaCl

0.3M Na₃C₆H₅O₇

Autoclave and store at room temperature

AP (alkaline phosphatase) buffer:

Final concentration

5mM MgCl₂

100 mM NaCl

100mM tris-HCL (pH 9.5)

0.1% Tween-20

Make fresh every time

Bleaching solution:

Final concentration

3% H₂O₂

0.5% KOH

Make fresh every time

Probe design

This *in situ* uses Digoxigenin-labelled antisense RNA probes. Probes are transcribed from DNA templates, which are themselves generally amplified by RT-PCR of mRNA extracted from zebrafish embryos, larvae or tissues. Antisense transcription is achieved by the addition of a promoter sequence to the reverse primer used in the RT-PCR. Add the following sequence to the beginning (5' end) of your reverse primer, depending on the polymerase being used:

T7 - GGATCCTAATACGACTCACTATAG
T3 - GGATCCATTAACCCTCACTAAAGG
SP6 - TATTTAGGTGACACTATAG

Generally, T7 is the best of the three polymerases.

Alternatively, the PCR product can be cloned into a expression vector such that one of these promoters is at the 3' end of the template. This allows for long term storage and re-use of the probe, as the plasmid can be amplified by bacterial transformation.

Probe length can vary from <200 to >1000 nucleotides. Somewhere in the middle of this range is usually considered ideal. The longer the probe, the more difficult it is for it to penetrate the sample. Shorter probes meanwhile are more prone to off-target binding, especially to similar mRNAs (e.g. paralogues).

Probe synthesis

2h 15m

2 If the template DNA is a plasmid, it must first be linearised by restriction enzyme digestion. If it is a PCR product, it can be used directly.

Prepare probe synthesis reaction mix

Mix and incubate for 02:00:00 at 37-40 °C (temperature depending on the polymerase you are using-see polymerase sheet). Longer incubations, up to overnight, can be performed if yield is low with 2 hrs incubation.

3 OPTIONAL: add Δ 2 μL of DNase1 to remove the template DNA; ৩ 00:15:00 at \$37 °C We tend to skip this step. If you are doing a precipitation step it's better if you have the DNA template because it helps the precipitation of your RNA. The DNA is not DIG labelled so it does not affect your in situ reaction.

4 Probe purification (option 1)

30m

15m

Add \bot 115 μ L of water and \bot 15 μ L of 5M ammonium acetate (if you have the message machine kit for the capped RNA transcription you will find the ammonium acetate there).

Centrifuge at 4 °C for 00:15:00

Dispose of the supernatant and resuspend the RNA in $\boxed{1}$ 25 μ L of water.

Probe purification (option 2)

Alternatively, use an RNA purification kit (for example, ZYMO RNA clean-up kit). Add \pm 30 μ L of water before starting, as it's better if the minimum starting volume is \pm 50 μ L.

Elute in \blacksquare 25 μ L of water.

You can run $\boxed{1 \mu}$ of the purified probe on a gel to check yield/integrity.

- We find it best practice to at this point, **check probe yield and integrity**. Measure concentration using a Nanodrop; there should be no less than 100ng/uL. Run on a 1% agarose gel, ideally there should be crisp bands (smearing on a gel indicates RNA degradation)
- 6 Store probe

Store the probe at -20 °C diluted at least 1:1 in hybridization buffer (the formamide protects the probe from degradation). The working concentration of a given probe is variable and requires optimization. Generally, 1:100, 1:250 or 1:500 are good starting points.

Sample collection

4h 15m

7 Perform in Eppendorf tubes or well plates, depending on number of samples.

4h 25m

Fix embryos/larvae at desired stage in 4% PFA in PBS for 04:00:00 at room temperature or Overnight at 4 °C.

Wash x4 with 1x PBS, (5) 00:05:00 per wash

Note

If you haven't used PTU and wish to bleach your larvae, do so at this point. Incubate in bleaching solution (3% H₂0₂ and 0.5% KOH in water) for up to 1 hr.

Peroxide decomposition releases O₂ gas. DO NOT perform bleaching in a closed tube.

Observe the larvae every 5-10 mins and stop bleaching by washing with PBS x2 once pigmentation has almost completely been removed.

8 Wash samples into methanol. Start with 50% for 00:10:00 at room temperature, then 100%.

Store samples Overnight at -20°C in methanol.

Embryos or larvae can be stored in methanol at -20°C for several months.

In situ: Day 1

4h 45m

9 Rehydrate samples: 20m

75% methanol/PBST (5) 00:05:00



50% methanol/PBST (5) 00:05:00



25% methanol/PBST (5) 00:05:00





Note

Bleaching can also be performed here.

10 Permeabilize samples with proteinase K digestion at room temperature. Dilute proteinase K stock (10mg/ml, 1000x) in PBST. Concentration and duration are stage dependent.

3-9 somites - 1x for 30 s-1 min

10-18 somites - 1x for 2 mins

24 hrs - 1x for 10 mins

2-3 dpf - 2x for 20-30 mins

4-6 dpf - 3x for 45mins - 1 hr

Larval brains - 1x for 1 min (or not at all)

Juvenile brains - 2x for 20 mins

Quickly wash twice with PBST (no incubation) to stop digestion. 11 25m Post-fix with 4% PFA for 00:20:00 at room temperature. Wash x4 with PBST for 00:05:00 each. 12 Pre-hybridize samples by incubation in hybridization buffer for (5) 02:00:00 at (8 68 °C) Remove hybridization buffer and add probe diluted in the same buffer. Incubate 🚫 Overnight 5h In situ: Day 2 13 Remove probe and save for reuse (working probe dilutions can be used many times). Wash once with 2x SSC for 00:30:00 (perform at 6 68 °C Wash twice with 0.2x SSC for 00:30:00 each (perform at 68 °C). Note SSC washes determine the stringency of probe binding. Low salt, high temperature conditions cause partial probe/mRNA duplexes to unravel, thereby reducing off-target staining of low complimentarity mRNAs. 4h 14 Wash with PBST for 1 hr with several changes. Note The times given for washes are the minimum times. In the experience of some lab members, elongating the PBST washes (especially those directly post-probe above) can significantly reduce background staining. These washes can even be extended overnight, adding an extra day to the protocol.

Incubate samples in anti-DIG-AP Fab fragments (Roche) diluted 1:6000 in MaBI Overnight

Block in MaBl at room temperature for at least 02:00:00

6

at 4 °C . Leave samples on a plate rocker if possible.

In situ: Day 3

20m

15 Wash x3 with PBST.

20m

Wash multiple times in fresh AP buffer over 1-2 hrs.

Prepare NBT/BCIP (Roche) solution (🗷 1 µL NBT, 🗷 3.5 µL BCIP, 🗷 1 mL AP buffer) just prior to use.

Develop in situs in NBT/BCIP solution at room temperature. Protect samples from light at this point (e.g. cover with aluminium foil). Check development under a microscope every 5-10 minutes initially.

It is easier to do this if the samples are in a plate or dish. Depending on the probe/target mRNA, development can take minutes to hours (even overnight).

Note

Concentration of probe, anti-DIG fragments and NBT/BCIP solution all affect the final staining intensity, development time and the amount of background staining. All can be freely varied, however you risk getting no staining or excessive background by doing so. Sometimes it is necessary to perform an *in situ* mutliple times to optimise these parameters, especially when using a new probe.

Stop reaction with two quick PBST washes.

Post-fix with 4% PFA for 00:20:00 at room temperature.

Wash x3 with PBS.

Transfer samples to glycerol through a graded series of glycerol/PBS washes (25%, 50% and 75%). Store samples in 100% glycerol at $\$ 4 \degree C$, in the dark.

Image samples under a compound microscope mounted in either glycerol or low-melting agarose.