

HCR v3.0 protocol for whole-mount zebrafish embryos and larvae (*Danio rerio*)

This protocol has not been optimized for all stages and should only be used as a template.

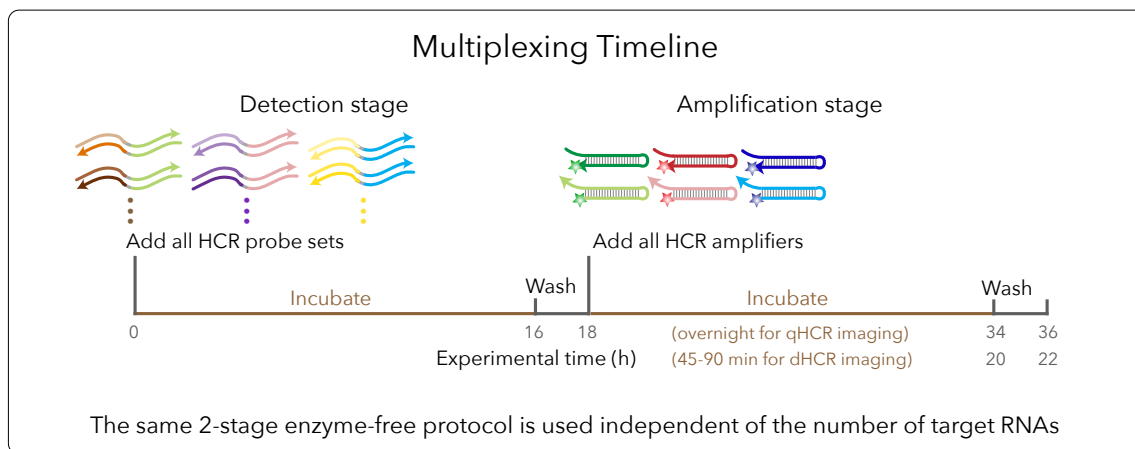
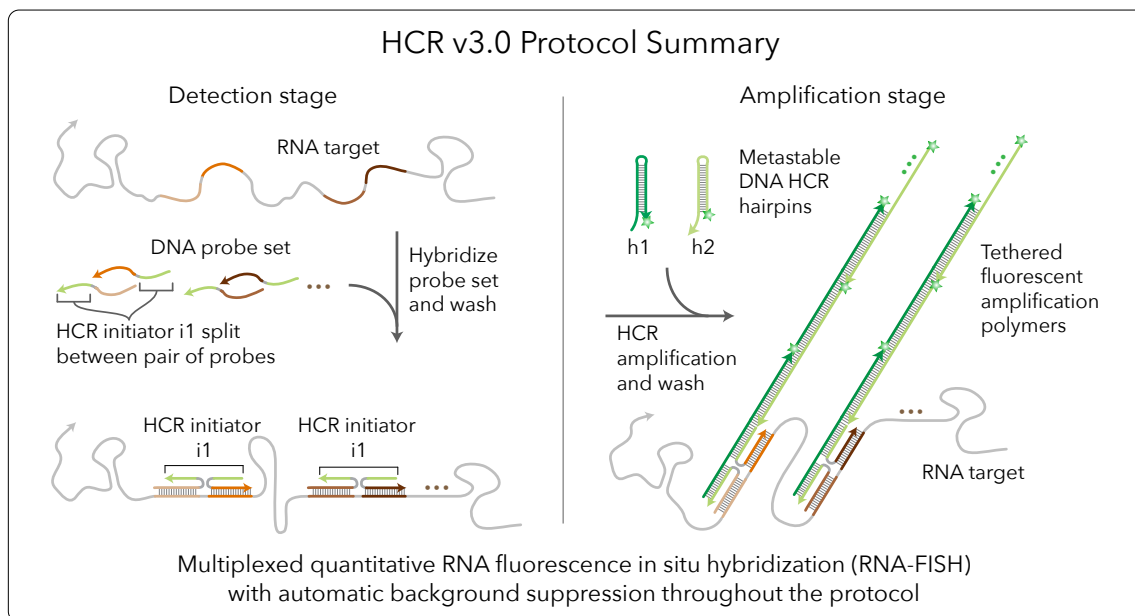
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Storage conditions

Store HCR probe sets, HCR amplifiers, HCR probe hybridization buffer, and HCR probe wash buffer at -20 °C. Store HCR amplification buffer at 4 °C. On the bench top, keep stock solutions on ice. Make sure all solutions are well mixed before use.



Preparation of fixed whole-mount zebrafish embryos and larvae

1. Collect zebrafish embryos and incubate at 28 °C in a petri dish with egg H₂O.
2. Exchange egg H₂O with egg H₂O containing 0.003% of 1-phenyl 2-thiourea (PTU) when embryos reach 12 hpf.
NOTE: PTU treatment is not necessary if working with embryos younger than 30 hpf.
NOTE: PTU inhibits melanogenesis but can be toxic at high concentrations. PTU treatment must start before the initial pigmentation occurs as PTU does not remove pigment that has already formed. PTU treatment is not necessary for nacre embryos used in this paper.
3. Replace with fresh egg H₂O containing 0.003% of PTU everyday until the larvae reach 5 dpf (days post-fertilization).
4. Transfer ~40 embryos/larvae (5 dpf) to a 2 mL eppendorf tube and remove excess egg H₂O.
5. Fix embryos/larvae in 2 mL of 4% paraformaldehyde (PFA) for 24 h at 4 °C.
CAUTION: use PFA with extreme care as it is a hazardous material.
NOTE: use fresh PFA and cool to 4 °C before use to avoid increased autofluorescence.
6. Wash embryos/larvae 3 × 5 min with 1 mL of 1× phosphate-buffered saline (PBS) to stop the fixation.
NOTE: avoid using calcium chloride and magnesium chloride in PBS as this leads to increased autofluorescence in the embryos/larvae.
7. Dehydrate and permeabilize with a series of methanol (MeOH) washes (1 mL each):
 - (a) 100% MeOH for 4 × 10 min
 - (b) 100% MeOH for 1 × 50 min.
8. Store embryos/larvae at -20 °C overnight before use.
NOTE: Embryos/larvae can be stored for six months at -20 °C.
9. Transfer the required number of embryos/larvae for an experiment to a 2 mL eppendorf tube.
10. Rehydrate with a series of graded 1 mL MeOH/PBST washes for 5 min each at room temperature:
 - (a) 75% MeOH / 25% PBST
 - (b) 50% MeOH / 50% PBST
 - (c) 25% MeOH / 75% PBST
 - (d) 5 × 100% PBST.
11. Treat 5 dpf embryos/larvae with 1 mL of proteinase K (30 µg/mL) for 45 min at room temperature.
NOTE: Proteinase K concentration and treatment time should be reoptimized for each batch of proteinase K, or for samples at a different developmental stage. Skip proteinase K treatment and postfixation (steps 11–14) for embryos 30 hpf and younger.

12. Wash embryos/larvae two times with PBST (1 mL each) without incubation.
13. Postfix with 1 mL of 4% PFA for 20 min at room temperature.
14. Wash embryos/larvae 5 × 5 min with 1 mL of PBST.

Buffer recipes for sample preparation

6% PTU stock solution

6% PTU

For 100 mL of solution

6 g of 1-phenyl 2-thiourea powder

Fill up to 100 mL with egg H₂O

Heat solution at 50–60 °C overnight to dissolve powder

0.3% PTU in egg H₂O

0.3% PTU

For 50 mL of solution

2.5 mL of 6% PTU

Fill up to 50 mL with with egg H₂O

4% paraformaldehyde (PFA)

4% PFA

1 × PBS

For 25 mL of solution

1 g of PFA powder

25 mL of 1 × PBS

Heat solution at 50–60 °C to dissolve powder

PBST

1 × PBS

0.1% Tween 20

For 50 mL of solution

5 mL of 10 × PBS

500 μL of 10% Tween 20

Fill up to 50 mL with ultrapure H₂O

Proteinase K solution

30 μg/mL proteinase K

For 1 mL of solution

1.5 μL of 20 mg/mL proteinase K

Fill up to 1 mL with PBST

NOTE: avoid using calcium chloride and magnesium chloride in PBS as this leads to increased autofluorescence in the samples.

Multiplexed HCR v3.0 protocol

Detection stage

1. For each sample, transfer 8 embryos/larvae to a 1.5 mL tube.
2. Pre-hybridize with 500 μL of probe hybridization buffer for 30 min at 37 °C.
CAUTION: probe hybridization buffer contains formamide, a hazardous material.
3. Prepare probe solution by adding 2 pmol of each probe set (e.g. 2 μL of 1 μM stock) to 500 μL of probe hybridization buffer at 37 °C.
NOTE: For dHCR imaging, use higher concentration of probe to improve probe hybridization efficiency.
4. Remove the pre-hybridization solution and add the probe solution.
5. Incubate embryos/larvae overnight (12–16 h) at 37 °C.
6. Remove excess probes by washing embryos/larvae 4 \times 15 min with 500 μL of probe wash buffer at 37 °C.
CAUTION: probe wash buffer contains formamide, a hazardous material.
NOTE: pre-heat probe wash buffer to 37 °C before use.
7. Wash embryos/larvae 2 \times 5 min with 5 \times SSCT at room temperature.

Amplification stage

1. Pre-amplify embryos/larvae with 500 μL of amplification buffer for 30 min at room temperature.
NOTE: equilibrate amplification buffer to room temperature before use.
2. Separately prepare 30 pmol of hairpin h1 and 30 pmol of hairpin h2 by snap cooling 10 μL of 3 μM stock (heat at 95 °C for 90 seconds and cool to room temperature in a dark drawer for 30 min).
NOTE: HCR hairpins h1 and h2 are provided in hairpin storage buffer ready for snap cooling. h1 and h2 should be snap cooled in separate tubes.
3. Prepare hairpin solution by adding snap-cooled h1 hairpins and snap-cooled h2 hairpins to 500 μL of amplification buffer at room temperature.
4. Remove the pre-amplification solution and add the hairpin solution.
5. Incubate the embryos/larvae overnight (12–16 h) in the dark at room temperature.
NOTE: For dHCR imaging, amplify for a shorter period of time to ensure single-molecule dots are diffraction-limited.
6. Remove excess hairpins by washing with 500 μL of 5 \times SSCT at room temperature:
 - (a) 2 \times 5 min
 - (b) 2 \times 30 min
 - (c) 1 \times 5 min
7. Samples can be stored at 4 °C protected from light before microscopy.

Citation Notes

For citation, please select from the list below as appropriate for your application:

- **HCR v3.0**

Automatic background suppression for dramatically enhanced performance (signal-to-background, qHCR precision, dHCR fidelity) and ease-of-use (no probe set optimization for new targets and organisms) (Choi *et al.*, 2018). Quantitative analysis modes:

qHCR imaging: analog mRNA relative quantitation with subcellular resolution in the anatomical context of thick autofluorescent samples.

qHCR flow cytometry: analog mRNA relative quantitation for high-throughput expression profiling of mammalian and bacterial cells.

dHCR imaging: digital mRNA absolute quantitation with single-molecule resolution in the anatomical context of thick autofluorescent samples.

Protocols for v3.0 in diverse organisms are adapted from the Zoo paper.

- **qHCR imaging**

mRNA relative quantitation with subcellular resolution in the anatomical context of whole-mount vertebrate embryos; read-out/read-in analysis framework (Trivedi *et al.*, 2018).

- **Zoo paper**

Protocols for multiplexed mRNA imaging in diverse sample types (Choi *et al.*, 2016):

bacteria in suspension	FFPE human tissue sections
generic sample in suspension	generic sample on slide
whole-mount chicken embryos	whole-mount fruit fly embryos
whole-mount mouse embryos	whole-mount sea urchin embryos
whole-mount worm larvae	whole-mount zebrafish embryos and larvae

- **dHCR imaging**

Single-molecule mRNA imaging in thick autofluorescent samples (e.g., 0.5 mm adult mouse brain sections) (Shah *et al.*, 2016).

- **qHCR northern blot**

Simultaneous quantification of RNA target size and abundance for up to 5 target RNAs (Schwarzkopf & Pierce, 2016).

- **HCR v2.0**

2nd generation in situ HCR technology (v2.0) using DNA HCR probes and DNA HCR amplifiers: 10× increase in signal, 10× reduction in cost, dramatic increase in reagent durability (Choi *et al.*, 2014).

- **HCR v1.0**

1st generation in situ HCR technology (v1.0) using RNA HCR probes and RNA HCR amplifiers: multiplexed mRNA imaging in whole-mount vertebrate embryos with simultaneous signal amplification for up to 5 target mRNAs (Choi *et al.*, 2010).

- **HCR mechanism**

Isothermal enzyme-free molecular signal amplification (Dirks & Pierce, 2004).

References

- Choi, H. M. T., Chang, J. Y., Trinh, L. A., Padilla, J. E., Fraser, S. E., & Pierce, N. A. (2010). Programmable in situ amplification for multiplexed imaging of mRNA expression. *Nat Biotechnol*, **28**(11), 1208–12.
- Choi, H. M. T., Beck, V. A., & Pierce, N. A. (2014). Next-generation in situ hybridization chain reaction: higher gain, lower cost, greater durability. *ACS Nano*, **8**(5), 4284–4294.
- Choi, H. M. T., Calvert, C. R., Husain, N., Huss, D., Barsi, J. C., Deverman, B. E., Hunter, R. C., Kato, M., Lee, S. M., Abelin, A. C. T., Rosenthal, A. Z., Akbari, O. S., Li, Y., Hay, B. A., Sternberg, P. W., Patterson, P. H., Davidson, E. H., Mazmanian, S. K., Prober, D. A., van de Rijn, M., Leadbetter, J. R., Newman, D. K., Readhead, C., Bronner, M. E., Wold, B., Lansford, R., Sauka-Spengler, T., Fraser, S. E., & Pierce, N. A. (2016). Mapping a multiplexed zoo of mRNA expression. *Development*, **143**, 3632–3637.
- Choi, H. M. T., Schwarzkopf, M., Fornace, M. E., Acharya, A., Artavanis, G., Stegmaier, J., Cunha, A., & Pierce, N. A. (2018). Third-generation in situ hybridization chain reaction: multiplexed, quantitative, sensitive, versatile, robust. *Development*, **145**, dev165753.
- Dirks, R. M., & Pierce, N. A. (2004). Triggered amplification by hybridization chain reaction. *Proc Natl Acad Sci USA*, **101**(43), 15275–15278.
- Schwarzkopf, M., & Pierce, N. A. (2016). Multiplexed miRNA northern blots via hybridization chain reaction. *Nucleic Acids Res*, **44**(15), e129.
- Shah, S., Lubeck, E., Schwarzkopf, M., He, T.-F., Greenbaum, A., Sohn, C. H., Lignell, A., Choi, H. M. T., Gradi-naru, V., Pierce, N. A., & Cai, L. (2016). Single-molecule RNA detection at depth via hybridization chain reaction and tissue hydrogel embedding and clearing. *Development*, **143**, 2862–2867.
- Trivedi, V., Choi, H. M. T., Fraser, S. E., & Pierce, N. A. (2018). Multidimensional quantitative analysis of mRNA expression within intact vertebrate embryos. *Development*, **145**, dev.156869.