

HCR v3.0 protocol for whole-mount nematode larvae (Caenorhabditis elegans)

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

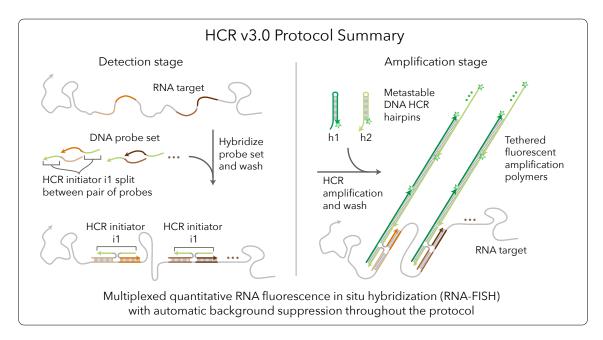
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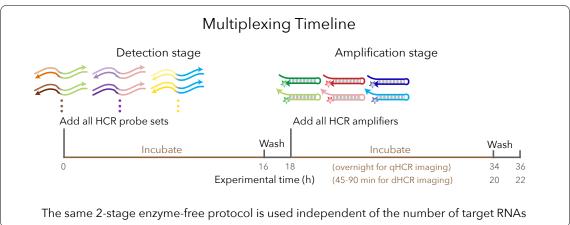
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Storage conditions

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.





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Preparation of fixed whole-mount nematode larvae

- 1. Wash nematode larvae off plate with 1 mL of M9 buffer and transfer to a 1.5 mL tube.
- 2. Centrifuge at $200 \times g$ for 2 min to bring larvae to the bottom of the tube and remove the M9 buffer.
- 3. Wash larvae 3 times with 1 mL of M9 buffer each. Centrifuge at $200 \times g$ for 2 min between washes.
- 4. Centrifuge and remove $\approx 800 \ \mu L$ of M9 buffer.
- 5. Aliquot larvae sufficient for a reaction into 1.5 mL tubes.
- 6. Add 1 mL of 4% paraformaldehyde (PFA).

CAUTION: use PFA with extreme care as it is a hazardous material.

NOTE: use fresh PFA to avoid increased autofluorescence.

7. Immediately freeze sample at -80 °C overnight before use.

Note: Larvae could stay in -80 °C freezer for long-term storage.

- 8. Fix larvae by thawing at room temperature for 45 min.
- 9. Wash larvae 2 times with 1 mL of PBST each. Bring larvae to the bottom of the tube with centrifugation at $200 \times g$ for 2 min in between washes.
- 10. Treat larvae with 1 mL of proteinase K (100 μ g/mL) for 10 min at 37 °C.

NOTE: Proteinase K concentration and treatment time should be reoptimized for each batch of proteinase K, or for samples at a different developmental stage.

- 11. Wash larvae 2 times with 1 mL of PBST each.
- 12. Incubate in 1 mL of 2 mg/mL of glycine solution for 15 min on ice.
- 13. Wash larvae 2 times with 1 mL of PBST each.

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Buffer recipes for sample preparation

M9 bufferFor 1 L of solution22 mM KH2PO43 g of KH2PO442 mM Na2HPO46 g of Na2HPO4

20.5 mM NaCl 5 g of NaCl

 1 mM MgSO_4 $1 \text{ mL of } 1 \text{ M MgSO}_4$

Fill up to 1 L with ultrapure H₂O

Sterilize by autoclaving

Store buffer at 4 °C before use

4% Paraformaldehyde (PFA) For 40 mL of solution

4% PFA 10 mL of 16% PFA solution

 $1 \times PBS$ 4 mL of $10 \times PBS$

Fill up to 40 mL with ultrapure H₂O

PBSTFor 50 mL of solution $1 \times PBS$ $5 \text{ mL of } 10 \times PBS$

0.1% Tween 20 500 μ L of 10% Tween 20

Fill up to 50 mL with ultrapure H₂O

Proteinase K solution For 1 mL of solution

100 μ g/mL proteinase K 5 μ L of 20 mg/mL proteinase K

Fill up to 1 mL with PBST

Glycine solution For 50 mL of solution

2 mg/mL glycine 100 mg of glycine
PBST Fill up to 50 mL with PBST

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

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Multiplexed HCR v3.0 protocol

Detection stage

- 1. Incubate larvae in 1 mL of 50% PBST / 50% probe hybridization buffer for 5 min at room temperature. CAUTION: *probe hybridization buffer contains formamide, a hazardous material*.
- 2. Centrifuge at $200 \times g$ for 2 min to remove solution.
- 3. Pre-hybridize larvae in 300 μ L of probe hybridization buffer at 37 °C for 1 h.
- 4. Prepare probe solution by adding 2 pmol of each probe set (e.g. 2 μ L of 1 μ M stock) to 200 μ L of probe hybridization buffer at 37 °C.

NOTE: For dHCR imaging, use higher concentration of probe to improve probe hybridization efficiency.

- 5. Add the probe solution to reach a final hybridization volume of 500 μ L.
- 6. Incubate larvae overnight (12–16 h) at 37 °C.
- 7. Remove excess probes by washing larvae 4×15 min with 1 mL 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.*

NOTE: bring larvae to the bottom of the tube with centrifugation at $500 \times g$ for 2 min for each wash.

8. Wash larvae 2×5 min with 1 mL of $5 \times$ SSCT at room temperature.

Amplification stage

1. Pre-amplify larvae with 300 μ L 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 μ L of 3 μ M 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 200 μ L of amplification buffer at room temperature.
- 4. Add the hairpin solution to reach a final amplification volume of 500 μ L.
- 5. Incubate the 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 1 mL of $5 \times$ SSCT at room temperature:
 - (a) $2 \times 5 \min$
 - (b) $2 \times 30 \text{ min}$
 - (c) $1 \times 5 \min$

NOTE: bring larvae to the bottom of the tube with centrifugation at $500 \times g$ for 2 min for each wash.

7. Samples can be stored at 4 °C protected from light before microscopy.

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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 framwork (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 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: $10 \times$ increase in signal, $10 \times$ 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).

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