

## HCR RNA-FISH 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.

### Technical support

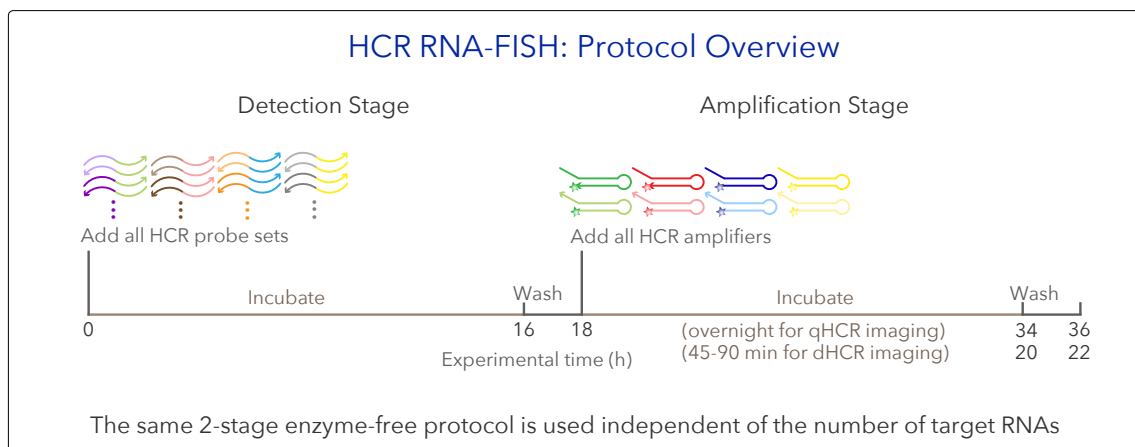
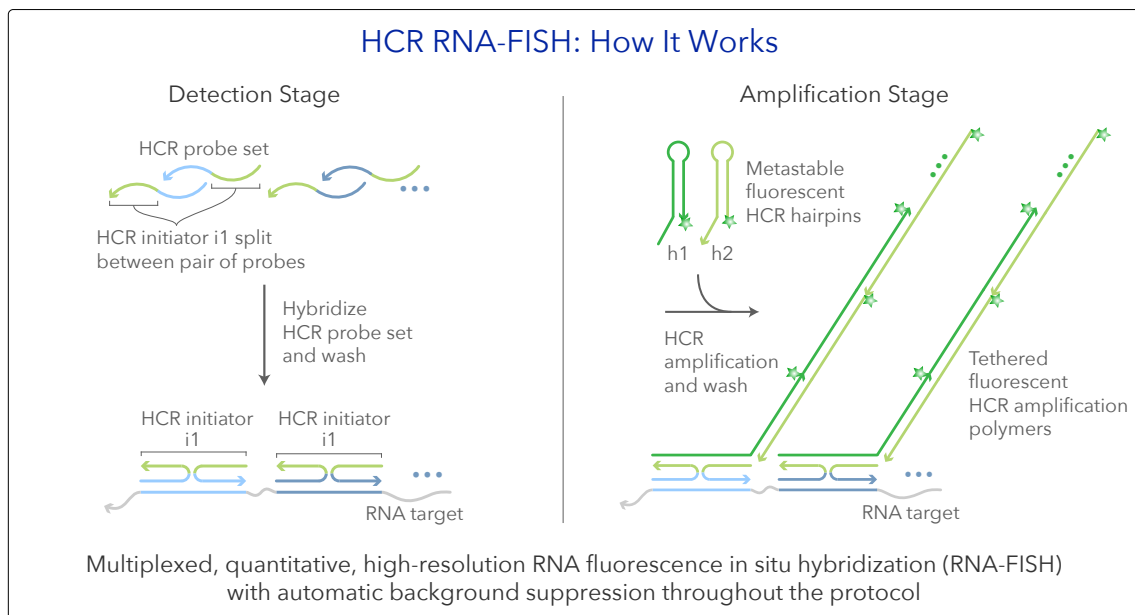
[support@molecularinstruments.com](mailto:support@molecularinstruments.com)

### Safety data sheets (SDS)

[www.molecularinstruments.com/safety](http://www.molecularinstruments.com/safety)

### Patents

Molecular Instruments® products are protected by and for use under patents: [www.molecularinstruments.com/patents](http://www.molecularinstruments.com/patents)



## **HCR RNA-FISH**

*Multiplexed, quantitative, high-resolution RNA imaging*

### **Multiplexed Experiment**

- Order one HCR RNA-FISH bundle per target RNA

### **Example 2-Plex Experiment**

- HCR bundle for target mRNA1
  - HCR split-initiator probe set: target mRNA1 for use with amplifier B1
  - HCR amplifier: B1-647
  - HCR RNA-FISH buffers: probe hybridization buffer, probe wash buffer, amplification buffer (for use with all bundles)
- HCR bundle for target mRNA2
  - HCR split-initiator probe set: target mRNA2 for use with amplifier B2
  - HCR amplifier: B2-488

### **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 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  $\approx 800 \mu\text{L}$  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  $-80^\circ\text{C}$  overnight before use.  
*NOTE: Larvae could stay in  $-80^\circ\text{C}$  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 ( $100 \mu\text{g}/\text{mL}$ ) for 10 min at  $37^\circ\text{C}$ .  
*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.

## Multiplexed HCR RNA-FISH 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  $\mu\text{L}$  of probe hybridization buffer at  $37^\circ\text{C}$  for 1 h.
4. Prepare probe solution by adding 2 pmol of each probe set (e.g. 2  $\mu\text{L}$  of 1  $\mu\text{M}$  stock) to 200  $\mu\text{L}$  of probe hybridization buffer at  $37^\circ\text{C}$ .  
*NOTE: For dHCR imaging, use higher probe concentration (e.g., 16 nM) to increase probe hybridization yield. If desired, this approach can also be used to increase signal for qHCR imaging.*
5. Add the probe solution to reach a final hybridization volume of 500  $\mu\text{L}$ .
6. Incubate larvae overnight (12–16 h) at  $37^\circ\text{C}$ .
7. Remove excess probes by washing larvae  $4 \times 15$  min with 1 mL of probe wash buffer at  $37^\circ\text{C}$ .  
*CAUTION: probe wash buffer contains formamide, a hazardous material.*  
*NOTE: pre-heat probe wash buffer to  $37^\circ\text{C}$  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  $2 \times 5$  min with 1 mL of  $5\times$  SSCT at room temperature.

### Amplification stage

1. Pre-amplify larvae with 300  $\mu\text{L}$  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  $\mu\text{L}$  of 3  $\mu\text{M}$  stock (heat at  $95^\circ\text{C}$  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  $\mu\text{L}$  of amplification buffer at room temperature.
4. Add the hairpin solution to reach a final amplification volume of 500  $\mu\text{L}$ .
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)  $2 \times 30$  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  $4^\circ\text{C}$  protected from light before microscopy.

## Buffer recipes

### M9 buffer

22 mM  $\text{KH}_2\text{PO}_4$   
42 mM  $\text{Na}_2\text{HPO}_4$   
20.5 mM NaCl  
1 mM  $\text{MgSO}_4$

### For 1 L of solution

3 g of  $\text{KH}_2\text{PO}_4$   
6 g of  $\text{Na}_2\text{HPO}_4$   
5 g of NaCl  
1 mL of 1 M  $\text{MgSO}_4$   
Fill up to 1 L with ultrapure  $\text{H}_2\text{O}$   
Sterilize by autoclaving  
Store buffer at 4 °C before use

### 4% Paraformaldehyde (PFA)

4% PFA  
1× PBS

### For 40 mL of solution

10 mL of 16% PFA solution  
4 mL of 10× PBS  
Fill up to 40 mL with ultrapure  $\text{H}_2\text{O}$

### PBST

1× PBS  
0.1% Tween 20

### For 50 mL of solution

5 mL of 10× PBS  
500  $\mu\text{L}$  of 10% Tween 20  
Fill up to 50 mL with ultrapure  $\text{H}_2\text{O}$

### Proteinase K solution

100  $\mu\text{g}/\text{mL}$  proteinase K

### For 1 mL of solution

5  $\mu\text{L}$  of 20 mg/mL proteinase K  
Fill up to 1 mL with PBST

### Glycine solution

2 mg/mL glycine  
PBST

### For 50 mL of solution

100 mg of glycine  
Fill up to 50 mL with PBST

### 5× SSCT

5× sodium chloride sodium citrate (SSC)  
0.1% Tween 20

### For 40 mL of solution

10 mL of 20× SSC  
400  $\mu\text{L}$  of 10% Tween 20  
Fill up to 40 mL with ultrapure  $\text{H}_2\text{O}$

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

## HCR Technology Citation Notes

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

- **HCR IHC + HCR RNA-FISH**

[HCR IHC + HCR RNA-FISH](#) enables a unified approach to multiplexed, quantitative, high-resolution protein immunohistochemistry (IHC) and RNA fluorescence in situ hybridization (RNA-FISH), with quantitative 1-step enzyme-free HCR signal amplification performed for all protein and RNA targets simultaneously ([Schwarzkopf et al., 2021](#)).

- **HCR IHC**

[HCR IHC](#) enables multiplexed, quantitative, high-resolution protein immunohistochemistry (IHC) in highly autofluorescent samples (e.g., FFPE brain tissue sections) ([Schwarzkopf et al., 2021](#)).

- **HCR RNA-FISH (v3.0)**

Third-generation [HCR RNA-FISH \(v3.0\)](#) enables multiplexed, quantitative, high-resolution RNA fluorescence in situ hybridization (RNA-FISH) with [automatic background suppression throughout the protocol](#) 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 RNA imaging](#): analog mRNA relative quantitation with subcellular resolution in the anatomical context of thick autofluorescent samples.
- [dHCR RNA imaging](#): digital mRNA absolute quantitation with single-molecule resolution in the anatomical context of thick autofluorescent samples.
- [qHCR RNA flow cytometry](#): analog mRNA relative quantitation for high-throughput expression profiling of mammalian cells and bacteria.

[Protocols for HCR RNA-FISH \(v3.0\)](#) in diverse organisms are adapted from the Zoo paper.

- **qHCR RNA imaging**

[qHCR RNA imaging](#) enables mRNA relative quantitation with subcellular resolution in the anatomical context of thick autofluorescent samples (e.g., whole-mount vertebrate embryos). The [read-out/read-in analysis framework](#) enables bidirectional quantitative discovery in an anatomical context ([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 solution
- generic sample on a slide
- mammalian cells on a slide
- mammalian cells in suspension
- whole-mount chicken embryos
- whole-mount fruit fly embryos
- whole-mount mouse embryos
- whole-mount nematode larvae
- whole-mount sea urchin embryos
- whole-mount zebrafish embryos and larvae

- **dHCR imaging**

[dHCR RNA imaging](#) enables RNA absolute quantitation with single-molecule resolution in the anatomical context of thick autofluorescent samples (e.g., 0.5 mm adult mouse brain sections) ([Shah et al., 2016](#)).

- **qHCR northern blots**

[qHCR northern blots](#) enable multiplexed quantification of RNA target size and abundance for up to 5 target RNAs ([Schwarzkopf & Pierce, 2016](#)).

- **HCR RNA-FISH (v2.0)**

Second-generation in situ HCR RNA-FISH technology (v2.0) using DNA HCR probes and 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 RNA-FISH (v1.0)**

First-generation HCR RNA-FISH 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**

The [hybridization chain reaction \(HCR\) mechanism](#) enables multiplexed, quantitative, 1-step, isothermal, enzyme-free signal amplification in diverse technological settings ([Dirks & Pierce, 2004](#)).

## HCR Technology 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.
- Schwarzkopf, M., Liu, M.C., Schulte, S.J., Ives, R., Husain, N., Choi, H.M.T., & Pierce, N. A. (2021). Hybridization chain reaction enables a unified approach to multiplexed, quantitative, high-resolution immunohistochemistry and in situ hybridization. *bioRxiv*, 2021.06.02.446311.
- 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.