

## In situ HCR v2.0 protocol for tissue section on slide

This protocol has not been validated for all sample types yet and should only be used as a template.  
Technical support: [support@moleculartechnologies.org](mailto:support@moleculartechnologies.org)

### Sample pre-treatment

Samples should be prepared in the same manner as for a traditional in situ hybridization, up to the probe hybridization step. This may include permeabilization and protease digestion. Then proceed with the protocol described below.

## Multiplexed in situ HCR v2.0 protocol

### Detection stage

1. Pre-warm a humidified chamber to 45 °C.
2. Dry slide by blotting edges on a Kimwipe.
3. Add 200  $\mu\text{L}$  of probe hybridization buffer on top of the tissue sample.  
*CAUTION: probe hybridization buffer contains formamide, a hazardous material.*
4. Pre-hybridize for 10 min inside the humidified chamber.
5. Prepare probe solution by adding 0.2 pmol of each probe (0.2  $\mu\text{L}$  of 1  $\mu\text{M}$  stock per probe) to 100  $\mu\text{L}$  of probe hybridization buffer at 45 °C.
6. Remove the pre-hybridization solution and drain excess buffer on slide by blotting edges on a Kimwipe.
7. Add 50–100  $\mu\text{L}$  of the probe solution on top of the tissue sample.  
*NOTE: Amount of probe solution depends on the size of the coverslip.*
8. Place a coverslip on the tissue sample and incubate overnight (12–16 h) in the 45 °C humidified chamber.
9. Immerse slide in probe wash buffer at 45 °C to float off coverslip.  
*CAUTION: probe wash buffer contains formamide, a hazardous material.*
10. Remove excess probes by incubating slide at 45 °C in:
  - (a) 75% of probe wash buffer / 25% 5 $\times$  SSCT for 15 min
  - (b) 50% of probe wash buffer / 50% 5 $\times$  SSCT for 15 min
  - (c) 25% of probe wash buffer / 75% 5 $\times$  SSCT for 15 min
  - (d) 100% 5 $\times$  SSCT for 15 min  
*NOTE: Wash solutions should be pre-heated to 45 °C before use.*
11. Immerse slide in 5 $\times$  SSCT for 5 min at room temperature.

### Amplification stage

1. Dry slide by blotting edges on a Kimwipe.
2. Add 200  $\mu\text{L}$  of amplification buffer on top of the tissue sample and pre-amplify in a humidified chamber for 30 min at room temperature.

3. Separately prepare 6 pmol of hairpin H1 and 6 pmol of hairpin H2 by snap cooling 2  $\mu\text{L}$  of 3  $\mu\text{M}$  stock in hairpin storage buffer (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.*
4. Prepare hairpin solution by adding all snap-cooled hairpins to 100  $\mu\text{L}$  of amplification buffer at room temperature.
5. Remove the pre-amplification solution and drain excess buffer on slide by blotting edges on a Kimwipe.
6. Add 50–100  $\mu\text{L}$  of the hairpin solution on top of the tissue sample.  
*NOTE: Amount of hairpin solution depends on the size of the coverslip.*
7. Place a coverslip on the tissue sample and incubate overnight (12–16 h) in a dark humidified chamber at room temperature.
8. Immerse slide in 5  $\times$  SSCT at room temperature to float off coverslip.
9. Remove excess hairpins by incubating slide in 5  $\times$  SSCT at room temperature for:
  - (a) 2  $\times$  30 min
  - (b) 1  $\times$  5 min

### **Sample mounting for microscopy**

1. Dry slide by blotting edges on a Kimwipe.
2. Add 50  $\mu\text{L}$  of SlowFade Gold antifade mountant with DAPI on top of tissue section.
3. Place a coverslip on top for imaging.
4. A Zeiss Axio Observer Z1 fluorescent inverted microscope equipped with an Plan-Apochromat 63 $\times$ /1.4 Oil DIC M27 objective was used in our lab to acquire tissue section images. Samples were illuminated with X-Cite Series 120 (EXFO Photonics) and images were collected with an ImagEM-1K EM-CCD digital camera (Hamamatsu).

## Buffer recipes for in situ HCR v2.0

Probes, amplifiers, probe hybridization buffer, and probe wash buffer should be stored at -20 °C. Amplification buffer should be stored at 4 °C. Keep these reagents on ice at all times during probe and amplifier preparation. Make sure all solutions are well mixed before use.

### Probe hybridization buffer

50% formamide  
5× sodium chloride sodium citrate (SSC)  
9 mM citric acid (pH 6.0)  
0.1% Tween 20  
50 µg/mL heparin  
1× Denhardt's solution  
10% dextran sulfate

### For 40 mL of solution

20 mL formamide  
10 mL of 20× SSC  
360 µL 1 M citric acid, pH 6.0  
400 µL of 10% Tween 20  
200 µL of 10 mg/mL heparin  
800 µL of 50× Denhardt's solution  
8 mL of 50% dextran sulfate  
Fill up to 40 mL with ultrapure H<sub>2</sub>O

### Probe wash buffer

50% formamide  
5× sodium chloride sodium citrate (SSC)  
9 mM citric acid (pH 6.0)  
0.1% Tween 20  
50 µg/mL heparin

### For 40 mL of solution

20 mL formamide  
10 mL of 20× SSC  
360 µL 1 M citric acid, pH 6.0  
400 µL of 10% Tween 20  
200 µL of 10 mg/mL heparin  
Fill up to 40 mL with ultrapure H<sub>2</sub>O

### Amplification buffer

5× sodium chloride sodium citrate (SSC)  
0.1% Tween 20  
10% dextran sulfate

### For 40 mL of solution

10 mL of 20× SSC  
400 µL of 10% Tween 20  
8 mL of 50% dextran sulfate  
Fill up to 40 mL with ultrapure H<sub>2</sub>O

### 5× SSCT

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

### For 40 mL of solution

10 mL of 20× SSC  
400 µL of 10% Tween 20  
Fill up to 40 mL with ultrapure H<sub>2</sub>O

### 50% dextran sulfate

50% dextran sulfate

### For 40 mL of solution

20 g of dextran sulfate powder  
Fill up to 40 mL with ultrapure H<sub>2</sub>O

## **Reagents and supplies**

Formamide (Deionized) (Ambion Cat. # AM9342)

20× sodium chloride sodium citrate (SSC) (Life Technologies Cat. # 15557-044)

Heparin (Sigma Cat. # H3393)

50× Denhardt's solution (Life Technologies Cat. # 750018)

Dextran sulfate, mol. wt. > 500,000 (Sigma Cat. # D6001)

10% Tween 20 Solution (Bio-Rad Cat. # 161-0781)

SlowFade Gold antifade mountant with DAPI (Life Technologies Cat. # S36938)

## Citation Notes

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- **In situ HCR v3.0**  
Automatic background suppression for dramatically enhanced performance (signal-to-background and quantitative precision) and ease-of-use (no probe set optimization for new targets) (Choi *et al.*, 2018). v3.0 supports three quantitative modes: qHCR imaging – analog mRNA relative quantitation with subcellular resolution; qHCR flow cytometry – analog mRNA relative quantitation for high-throughput single-cell analysis; dHCR imaging – digital mRNA absolute quantitation. Protocols for v3.0 in diverse organisms are adapted from the zoo paper.
- **Quantitative HCR (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	whole-mount zebrafish embryos and larvae
whole-mount fruit fly embryos	whole-mount chicken embryos
whole-mount worm larvae	whole-mount mouse embryos
whole-mount sea urchin embryos	FFPE human tissue sections
- **Single-molecule HCR imaging**  
Single-molecule mRNA imaging in thick autofluorescent samples (e.g., 0.5 mm adult mouse brain sections) (Shah *et al.*, 2016).
- **Multiplexed quantitative HCR (qHCR) northern blots**  
Simultaneous quantification of RNA target size and abundance with signal amplification for up to 5 target RNAs (Schwarzkopf & Pierce, 2016).
- **In situ HCR v2.0**  
2nd generation in situ HCR technology (v2.0) using DNA probes and DNA HCR amplifiers: 10× increase in signal, 10× reduction in cost, dramatic increase in reagent durability (Choi *et al.*, 2014).
- **Shielded covalent (SC) probes**  
Highly selective covalent capture of DNA and RNA targets, overcoming the longstanding tradeoff between selectivity and durable target capture (Vieregg *et al.*, 2013).
- **In situ HCR v1.0**  
1st generation in situ HCR technology (v1.0) using RNA 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).
- **Hybridization chain reaction (HCR) mechanism**  
Isothermal enzyme-free molecular signal amplification (Dirks & Pierce, 2004).

We are happy to provide advice and technical support: [support@moleculartechnologies.org](mailto:support@moleculartechnologies.org).

– The Molecular Technologies Team

## 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. ([pdf](#), [supp info](#), [supp movie 1](#), [supp movie 2](#))
- 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. ([pdf](#), [supp info](#), [supp movie](#))
- 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. ([pdf](#), [supp info](#), [fruit fly movie](#), [sea urchin movie](#), [zebrafish movie](#), [chicken movie](#), [mouse movie](#))
- 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. *bioRxiv*, doi: <http://dx.doi.org/10.1101/285213>. ([pdf](#), [supp info](#))
- Dirks, R. M., & Pierce, N. A. (2004). Triggered amplification by hybridization chain reaction. *Proc Natl Acad Sci USA*, **101**(43), 15275–15278. ([pdf](#))
- Schwarzkopf, M., & Pierce, N. A. (2016). Multiplexed miRNA northern blots via hybridization chain reaction. *Nucleic Acids Res*, **44**(15), e129. ([pdf](#), [supp info](#))
- 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. ([pdf](#), [supp info](#), [supp movie 1](#), [supp movie 2](#), [supp movie 3](#))
- 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**, doi:10.1242/dev.156869. ([pdf](#), [supp info](#), [Read-out/Read-in 1.0 package](#))
- Vieregg, J. R., Nelson, H. M., Stoltz, B. M., & Pierce, N. A. (2013). Selective nucleic acid capture with shielded covalent probes. *J Am Chem Soc*, **135**(26), 9691–9699. ([pdf](#), [supp info](#))