

## HCR v3.0 protocol for bacteria in suspension

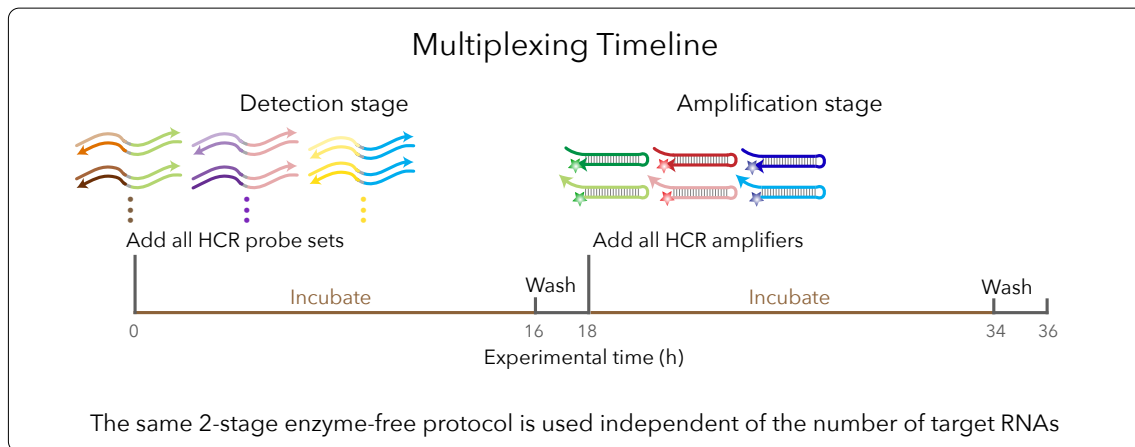
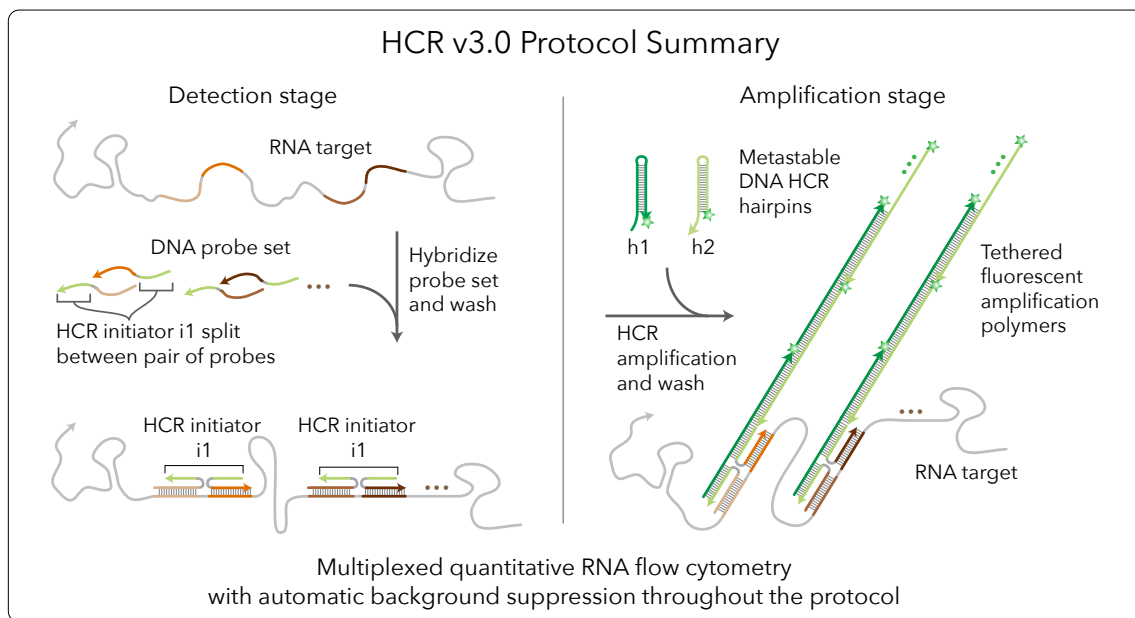
This protocol has been optimized for *Escherichia coli* and should only be used as a template for other types of bacteria.  
Technical support: [support@molecularinstruments.com](mailto:support@molecularinstruments.com)

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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.



## Sample preparation

1. Grow *E. coli* from streaked plate or frozen glycerol stocks in 2–3 mL of LB media overnight in a 37 °C shaker.
2. Dilute to make a 5 mL liquid culture with  $OD_{600} = 0.05$ .
3. Incubate in a 37 °C shaker until  $OD_{600} \approx 0.5$  (exponential phase).
4. Aliquot 1 mL of cells and centrifuge for 10 min.  
*NOTE: Centrifugation should be as gentle as possible to pellet cells. For E. coli, all centrifugation steps are done at  $4000 \times g$ .*
5. Remove supernatant and re-suspend cells in 750  $\mu\text{L}$  of  $1 \times$  phosphate-buffered saline (PBS).  
*NOTE: remove all solutions via pipetting throughout the protocol.*
6. Add 250  $\mu\text{L}$  of 4% formaldehyde and incubate overnight at 4 °C.  
*CAUTION: use formaldehyde with extreme care as it is a hazardous material.*
7. Centrifuge for 10 min and remove supernatant.
8. Re-suspend cells in 150  $\mu\text{L}$  of  $1 \times$  PBS.
9. Add 850  $\mu\text{L}$  of 100% MeOH and store cells at -20 °C before use.

## Buffer recipes for sample preparation

### LB media

5 g of Novagen LB Broth Miller powder  
Fill up to 200 mL with ultrapure H<sub>2</sub>O  
Autoclave at 121 °C for 20 min

### 4% formaldehyde

4% formaldehyde  
 $1 \times$  PBS

### For 4 mL of solution

1 mL of 16% formaldehyde  
0.4 mL of  $10 \times$  PBS  
Fill up to 4 mL with ultrapure H<sub>2</sub>O

*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. Transfer 150  $\mu\text{L}$  of cells into a 1.5 mL eppendorf tube.
2. Centrifuge for 5 min and remove supernatant.
3. Wash cells with 500  $\mu\text{L}$  of 1 $\times$  PBST. Centrifuge for 5 min and remove the supernatant.
4. Re-suspend the pellet with 400  $\mu\text{L}$  of probe hybridization buffer (with low M.W. dextran sulfate) and pre-hybridize for 1 h at 37  $^{\circ}\text{C}$ .  
*CAUTION: probe hybridization buffer contains formamide, a hazardous material.*  
*NOTE: Probe hybridization buffer with low M.W. dextran sulfate will be used in this protocol.*
5. In the meantime, prepare probe solution by adding 2 pmol of each probe set (e.g. 2  $\mu\text{L}$  of 1  $\mu\text{M}$  stock) to 100  $\mu\text{L}$  of probe hybridization buffer pre-heated to 37  $^{\circ}\text{C}$ .
6. Add the probe solution directly to the sample to reach a final probe concentration of 4 nM.
7. Incubate the sample overnight at 37  $^{\circ}\text{C}$ .
8. Add 1 mL of probe wash buffer to the sample to dilute the probes.  
*CAUTION: probe wash buffer contains formamide, a hazardous material.*  
*NOTE: pre-heat probe wash buffer to 37  $^{\circ}\text{C}$  before use.*
9. Centrifuge for 5 min and remove the wash solution.
10. Re-suspend the cell pellet with 500  $\mu\text{L}$  of probe wash buffer (pre-heated to 37  $^{\circ}\text{C}$ ).
11. Incubate for 5 min at 37  $^{\circ}\text{C}$  and remove the wash solution by centrifugation for 5 min.
12. Repeat steps 10 and 11 for two additional times but with 10 min incubation.
13. Proceed to hairpin amplification.

### Amplification stage

1. Re-suspend the cell pellet with 150  $\mu\text{L}$  of amplification buffer (with low M.W. dextran sulfate) and pre-amplify for 30 min at room temperature.  
*NOTE: Amplification buffer with low M.W. dextran sulfate will be used in this protocol. Equilibrate amplification buffer to room temperature before use.*
2. Separately prepare 15 pmol of hairpin h1 and 15 pmol of hairpin h2 by snap cooling 5  $\mu\text{L}$  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 100  $\mu\text{L}$  of amplification buffer at room temperature.
4. Add the hairpin solution directly to the sample to reach a final hairpin concentration of 60 nM.

5. Incubate the sample overnight (>12 h) in the dark at room temperature.
6. Add 1 mL of 5× SSCT at room temperature to the sample to dilute the hairpins.
7. Centrifuge for 5 min and remove the wash solution.
8. Re-suspend the cell pellet with 500  $\mu$ L of 5× SSCT.
9. Incubate for 5 min at room temperature and remove the wash solution by centrifugation for 5 min.
10. Repeat steps 8 and 9 for two additional times but with 10 min incubation.
11. Re-suspend the cell pellet in desired buffer (e.g. 5× SSCT) and volume.
12. Samples can be stored at 4 °C protected from light before microscopy or flow cytometry.

*NOTE: filter cells before flow cytometry.*

## 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

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