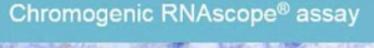
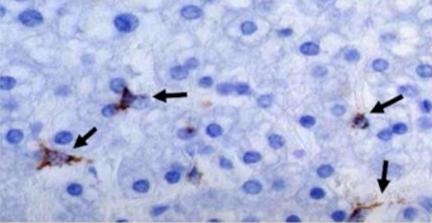
# TOPICS

- RNAscope Introduction and Overview: Embryonic and postnatal skeletal development
  - RNAscope technology: A novel RNA ISH technology, Wang et al. J. Mol.Diagn. 2011.
  - Chromogenic Red and the Multiplex Fluorescent Assays (Advanced Cell Diagnostics: ACD)
  - RNAscope assay workflow
- Tips for an RNAscope Assay
  - Sample fixation
  - Pretreatment

# Current State of the Art Tissue Based RNA Analysis with RNAscope Technology and Quantitative Analysi S. ISOTOPS ANA Share RNA ISH

#### IGFBP-3 expression in human liver tissue





- Standard oligo probes
- 10 min chromogenic reaction

ISOTOPIC RNA ISH

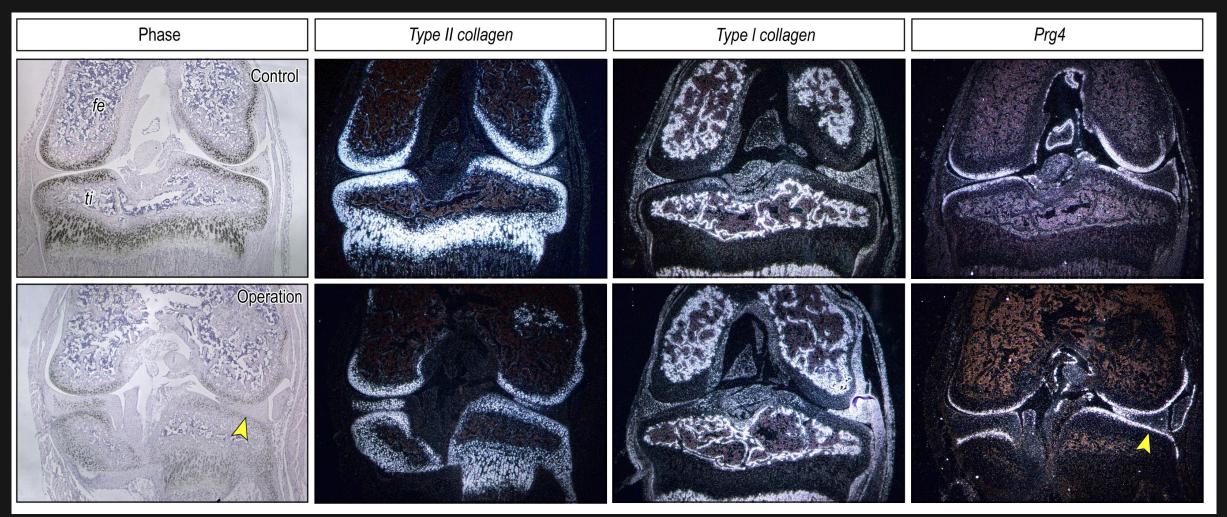
- S<sup>35</sup> labeled probes
- 48 hrs radiograph exposure

RNAscope® Assay: More signal, less background, faster detection

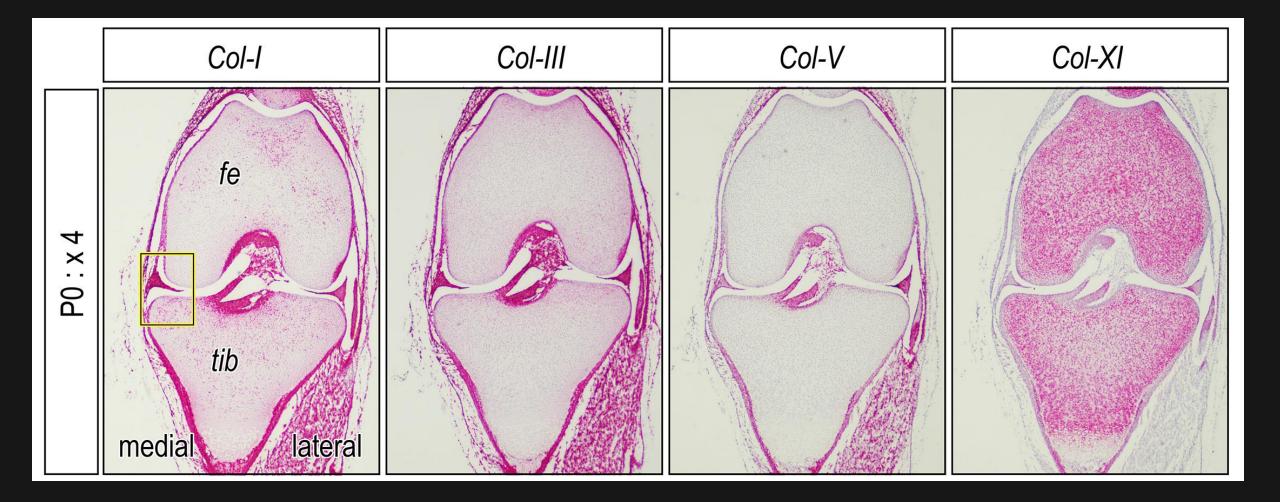
Monitoring, Inspection, Isotope, Dry & Liquid Waste, Emulsion, Dark room.....

#### Obtained from ACD

# ISH with 35S-labeled RNA probes



# RNAscope 2.5 HD Assay: RED

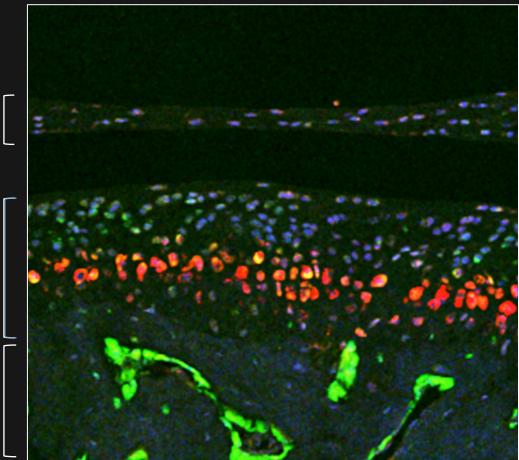


# **RNAscope Multiplex Fluorescent Assay**

# Articular Disc

Condylar (Fibro)Cartilage

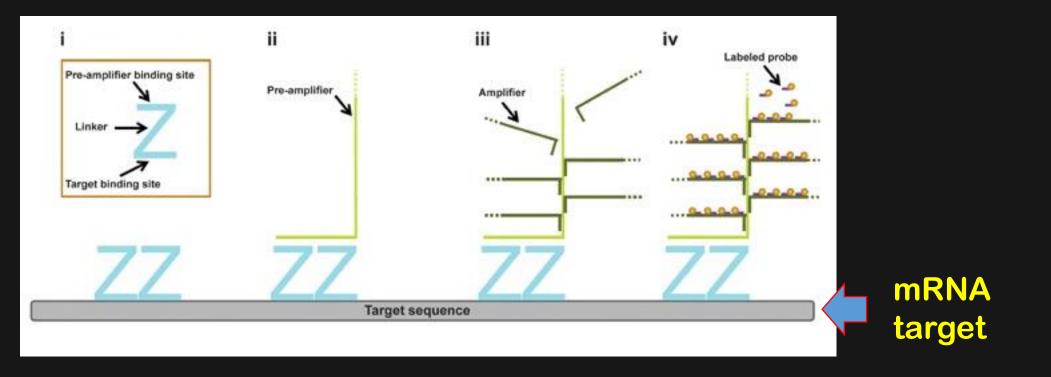
Subchondral bone



Fibrochondrocytes

Immature chondrocytes Osteoblasts and osteocytes

Type I collagen-<mark>Green</mark>, Type II collagen-<mark>Red</mark>, Dapi-<mark>Blue</mark>

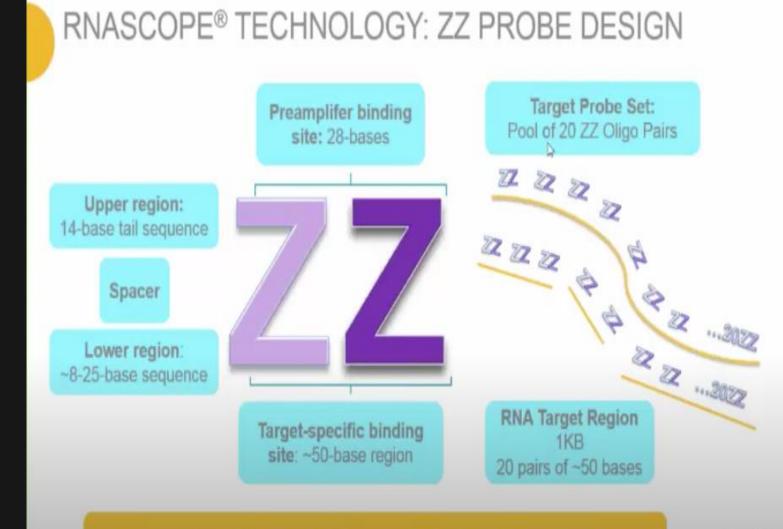


• **Step1**: Double Z target probes hybridize to the RNA target.

Obtained from ACD

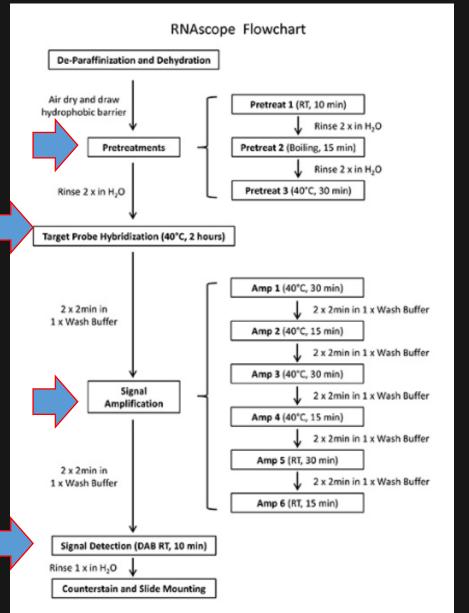
- Step2: Pre-amplifiers hybridize to the 28-base binding site formed by each double Z probe.
- Step3: Amplifiers are then bound to the multiple biding sites on each preamplifier.
- Step4: Labeled probes, containing a fluorescent molecule or chromogenic enzyme, bind to the binding sites on each amplifier.

# Each Target Z Probe Contains Three Elements



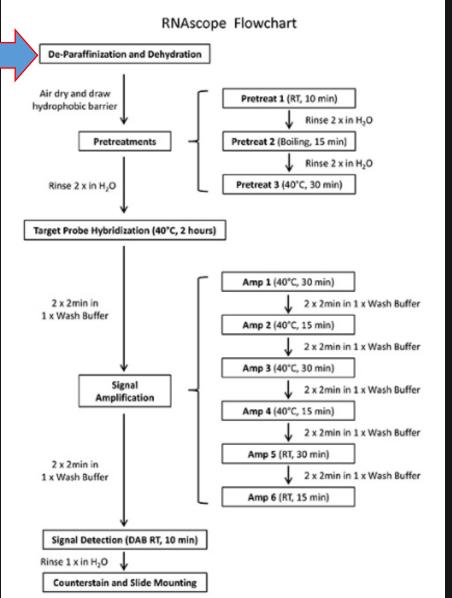
- The lower region of a double Z probe pair is about 50 bases long and is complementary to the target RNA.
- A spacer sequence links the upper and lower regions.
- The upper region of a double Z probe pair forms a 28 base biding site for the preamplifier.
- The Z probe pair needs to hybridize to the target RNA in tandem.
  Obtained from ACD

### Flowchart of RNAscope assay with FFPE sections, Chromogenic Red or Brown



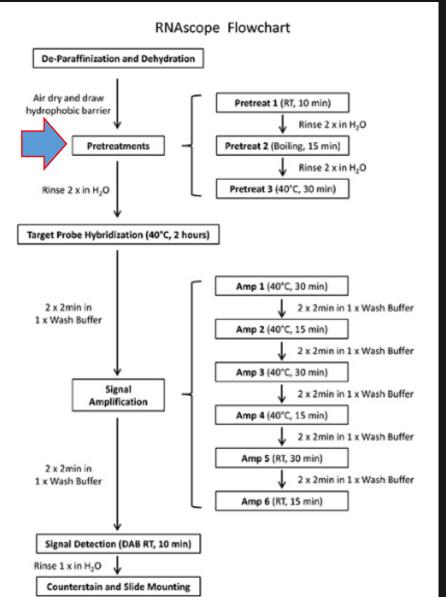
 RNA scope assay has a highly streamlined workflow that is similar to IHC. It consists of four major steps:

- Pretreatments
- Hybridization
- Signal amplifications
- Detection



# • Deparaffinization and Dehydration

- After baking (for 10 min at 60°C), deparaffinize tissue sections in Xylene for 2x5 min with agitation, and dehydrate in 100% Et-OH for 2x5 min with agitation.
- Air dry for 10 min at 40°C and draw a hydrophobic barrier around the tissue section with a Hydrophobic Barrier Pen. Dry the hydrophobic barrier completely at 40°C for 30 min.

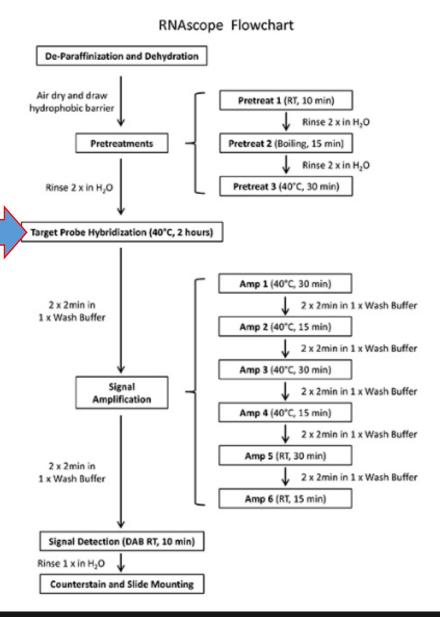


#### Pretreatments: ACD

- H2O2 block: RT, 10 min
- Epitope retrieval (pH6, citrate buffer): boiling, 10min
- Protease digestion: 40°C, 30min

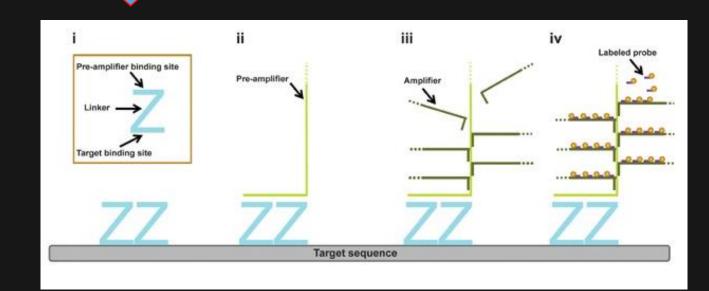
#### Embryonic/Postnatal Bone/(fibro)cartilage

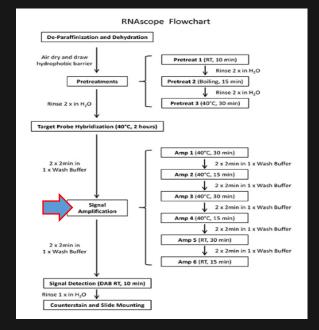
- H2O2 block: RT, 10 min
- Epitope retrieval
- Protease digestion
- Custom pretreatment reagent (Cat. 300040, \$90) @ 40°C for 30 min.

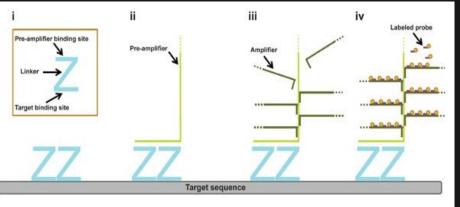


# Target Probe Hybridization:

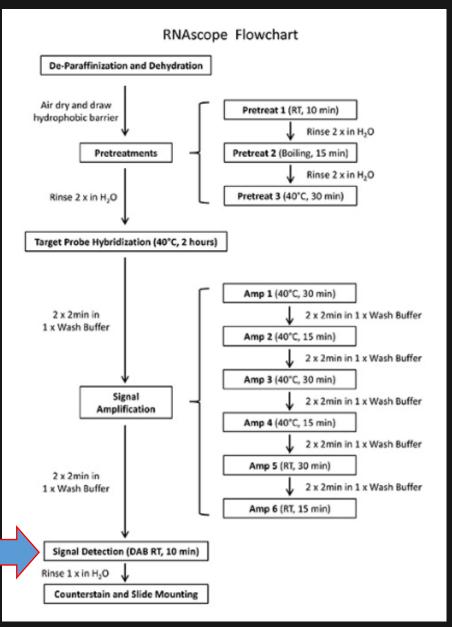
- Pre-warm probes @ 40°C, and then 40°C, 2hr
- 3-4 drops of probe solution: the solution still contains large amounts of un-hybridized Z probes.
- Do not use PARAFILM to cover the sections





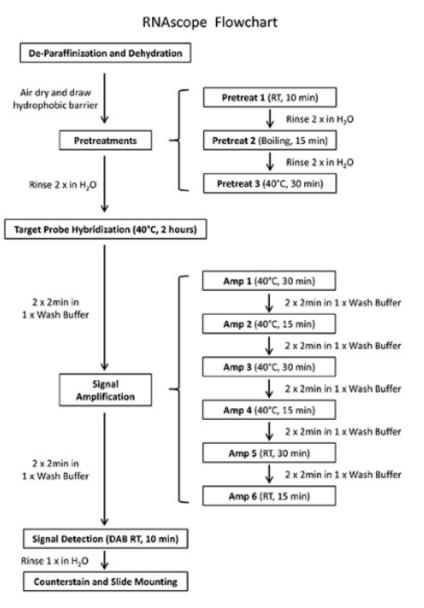


- Signal Amplification is achieved through Amp1-6.
  - Bring Amp1-6 to RT
  - Step 1: Pre-amplifier binds to the upper region of each double Z probe pair, 40°C, 30min
  - Step 2: Background reducer, 40°C, 15min
  - Step 3: Amplifiers bind to the binding site on each Pre-amplifier, 40°C, 30min
  - Step 4: Labeled probes, containing chromogenic enzymes or fluorescent molecules, adhere to the binding site on the amplifiers, 40°C, 15min
  - Amp5: **RT**, 30min or 2hr
  - Amp6: **RT**, 15min



#### Signal Detection

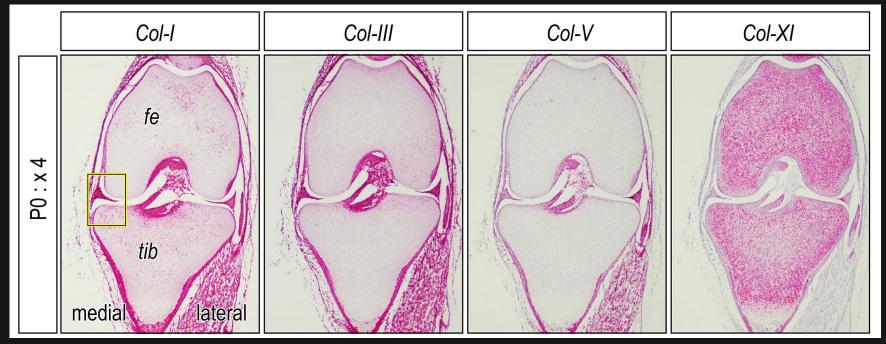
- Enzyme reaction is very quick in some cases (Prg4 2-3 min, Collagens 2-4 min), you need to observe the slide under the microscope.
- Dip the slide in H2O, postfix with 1-2%PFA for 10 min, and then counterstain with hematoxylin.



# Counterstaining

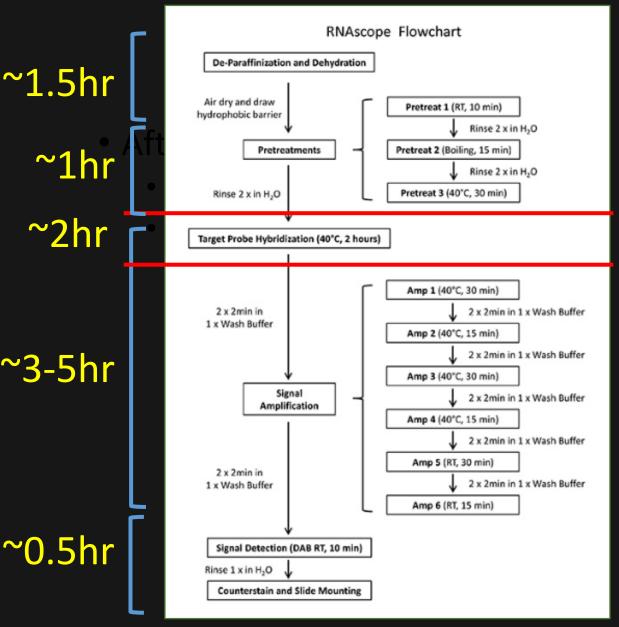
 Stain tissue section with hematoxylin solution (Hematoxylin QS, Vector: 1:125 with H2O ) for 1 min at RT, rinse with H2O until slides are clear. Dry slide O/N, dip slides in xylene and mount with xylene-based mounting media.

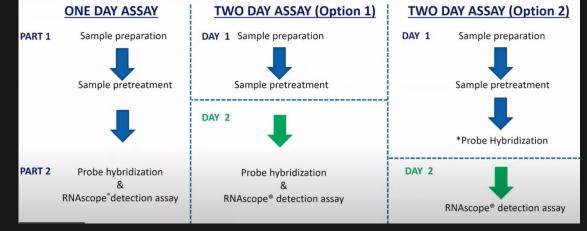
# RNAscope 2.5 HD Assay: RED



- ACD developed universal negative control probes targeting the DapB gene from the Bacillus. You can also order the sense probe with the sense direction.
- Cyclophilin B (PPIB) is used as a positive control probe. It is expressed at a low level , and so provides a rigorous control for sample quality. If PPIB is positive, then any target probe will detect your target RNAs.

# ONE DAY / TWO DAY ASSAY





**Obtained from ACD** 

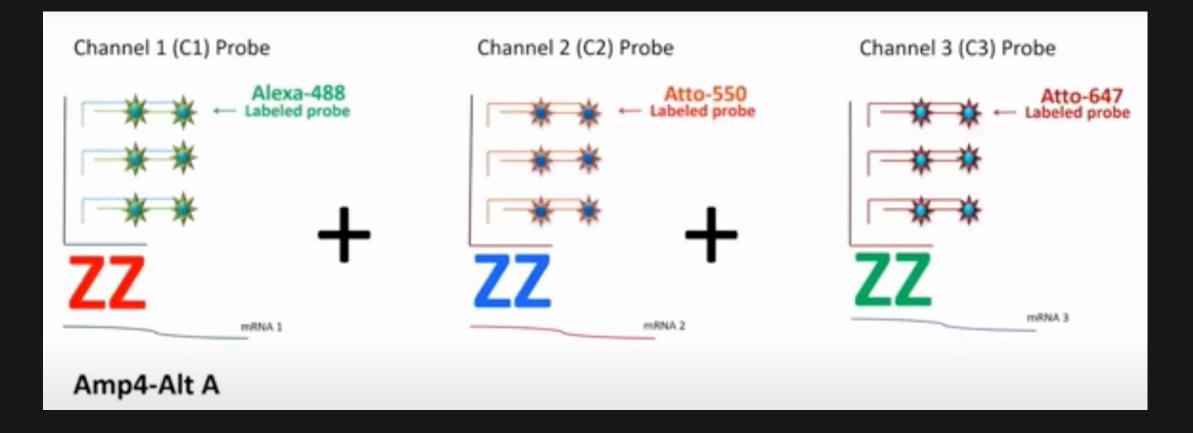
After hybridization RT, 2 min in 1xWash Buffer (0.1xSSC) Store @RT, O/N in 5xSSC (low stringency, keep slide from drying)

Next day, wash with 1xWash Buffer, then start amplification.

# **Critical Factors for Success of the RNAscope assay**

- Tissue should be fixed with fresh 4%PFA by injection. ACD recommends fresh 10% neutral buffered formalin at RT for 16-32 hr.
- Tissue decalcification: 10%EDTA with 2%PFA for 2-14 days, Formic acid (96%, Spectrum, F1089) @RT, for 2 days. But, reporter activity, IHC.....
- HybEZ oven is highly recommended since it enables optimal control of temperature and humidity for probe hybridization and signal amplification.
- Keep the tissue section from drying after hybridization.

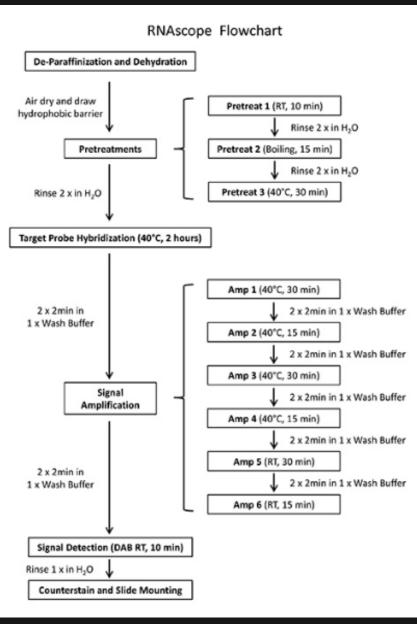
# **Multiplex Fluorescent Assays**

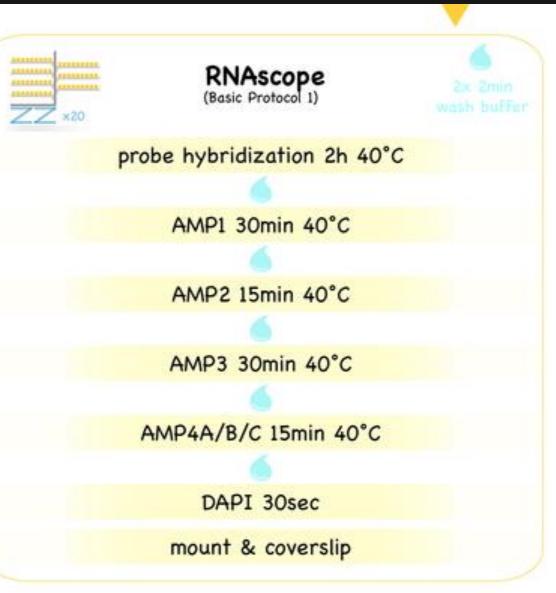


**Obtained from ACD** 

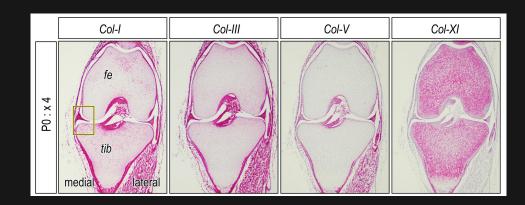
# **Chromogenic Red/Brown**

# **Multiplex Fluorescent**





#### RNAscope 2.5 HD Assay: RED



#### Channel 3 (C30 Channel 1 (C1) Channel 2 (C2) probe channel channel sensitivity highest weakest high cell type analysis of cell type marker 1 cell type marker 2 gene of interest (e.g. vGLUT1/2) (e.g. GAD1/2) target gene expression ..... ...... AAAAAAAAA ..... ...... \*\*\*\*\*\*\*\* \*\*\*\*\*\*\*\* ..... ........ ...... ...... \*\*\*\*\*\*\*\* \*\*\*\*\*\*\*\* \*\*\*\*\*\*\*\* AMP4A ........ ...... \*\*\*\*\*\*\*\* ...... ....... ...... ...... Atto550 7 Alexa488 77 Atto647 ...... ...... \*\*\*\*\*\*\*\* ......... ...... \*\*\*\*\*\*\*\* ........ ...... \*\*\*\*\*\*\*\* ...... ....... \*\*\*\*\*\*\*\*\* ..... ...... ...... AMP4B ...... ........ \*\*\*\*\*\*\*\* \*\*\*\*\*\*\*\* ...... ....... (recommended) Atto550 Alexa488 7 Atto647 -\*\*\*\*\*\*\*\* \*\*\*\*\*\*\*\* ....... \*\*\*\*\*\*\*\* \*\*\*\*\*\*\*\* \*\*\*\*\*\*\*\* ....... \*\*\*\*\*\*\*\* ...... ...... \*\*\*\*\*\*\*\* \*\*\*\*\*\*\*\* AMP4C ....... \*\*\*\*\*\*\*\* ...... ...... \*\*\*\*\*\*\*\* ........ ...... ....... \*\*\*\*\*\*\*\* Atto647 Atto550 Alexa488

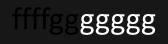
#### Figure 2.

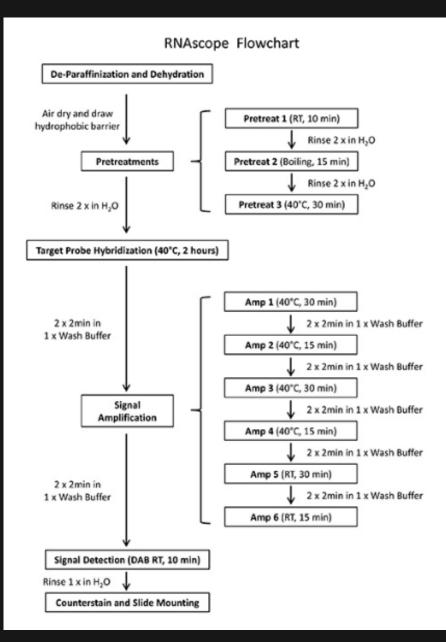
Multiplexing of the three channels in RNAscope. Detected fluorophores of the three channels (C1, C2, C3) can be adjusted by distinct amplification solution 4 (AMP4A, AMP4B, AMP4C). The sensitivity of the three channels is C1 > C2 >> C3. Therefore, we recommend examining the expression of a target gene (lowest expected expression) in different cell types using a channel 1 (C1) probe against this gene of interest and cell type-specific marker genes in Channel 2 & 3.

#### **Obtained from ACD**

# •The manual RNAscope procedure discussed here has been fully automated.

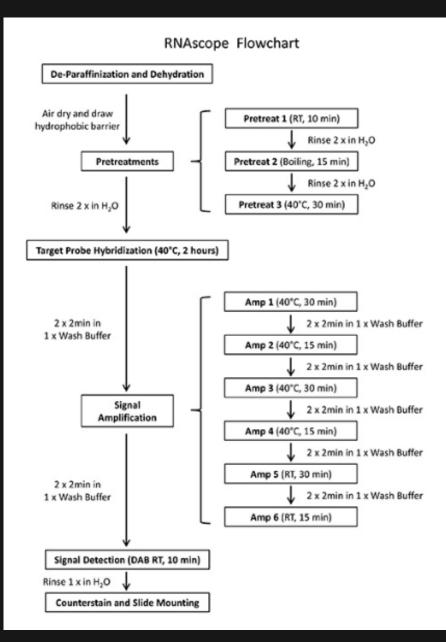
This should greatly facilitate standardization of assay conditions and save precious manual labor.





# Signal Detection

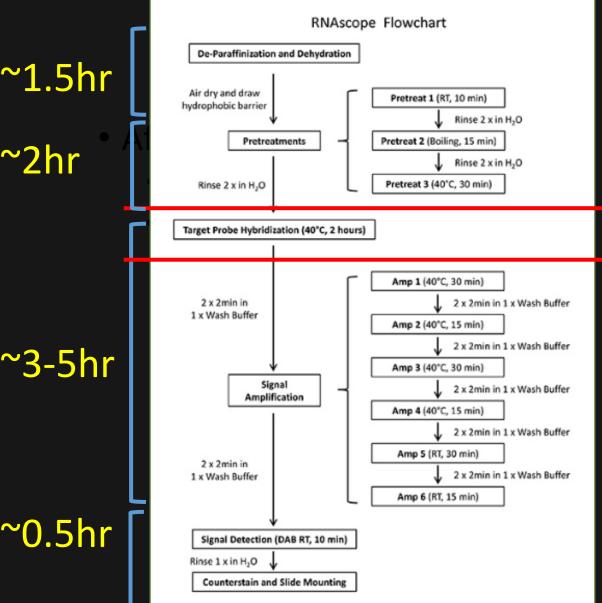
- Amp1: preamplifier
- Amp2 background reducer
- Amp3 amplifier
- Pam4 labeled probe
- Amp5
- Amp6

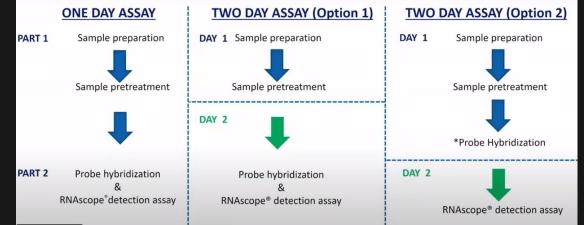


# Signal Detection

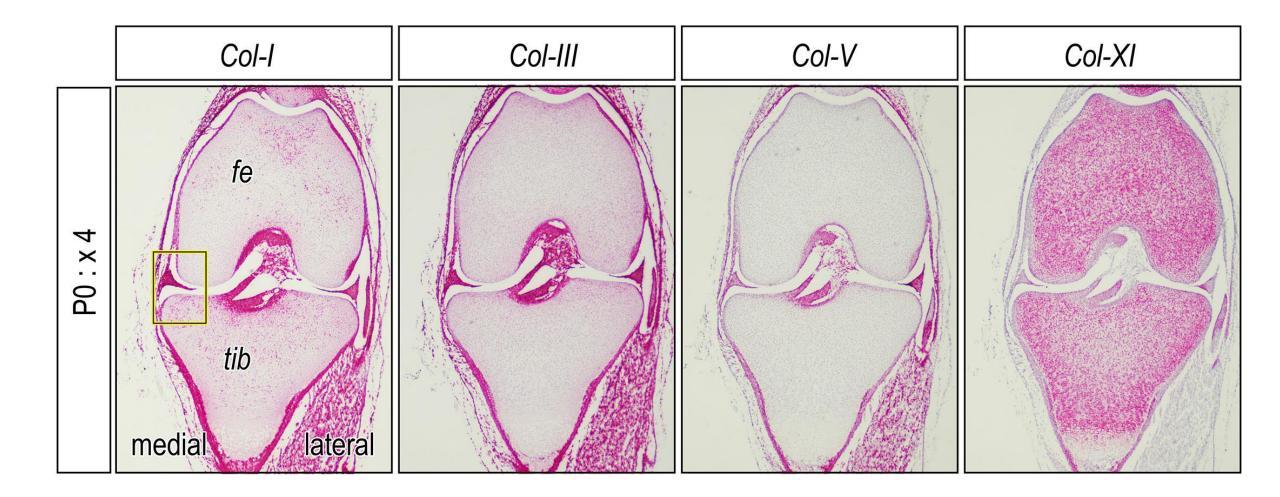
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- Pam4 labeled probe
- Amp5
- Amp6

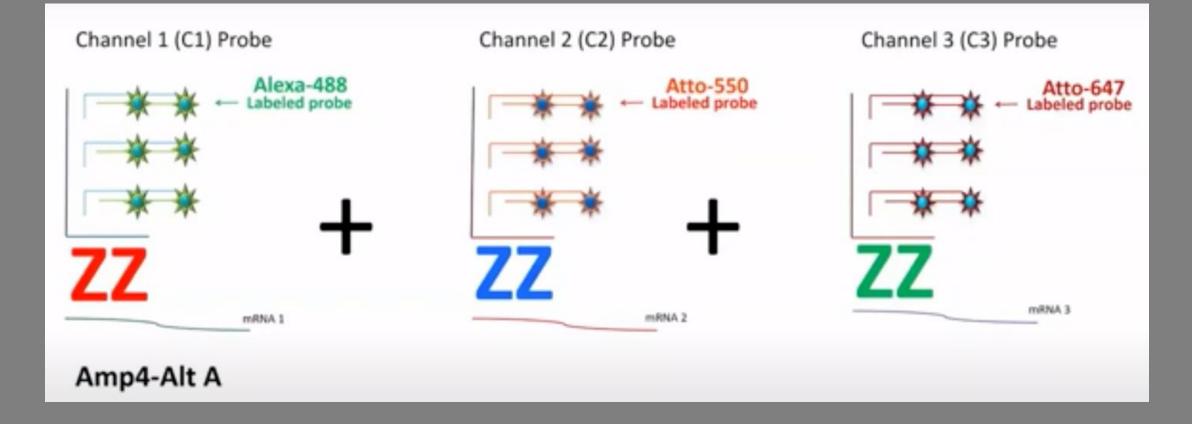
# Flowchart of RNAscope assay

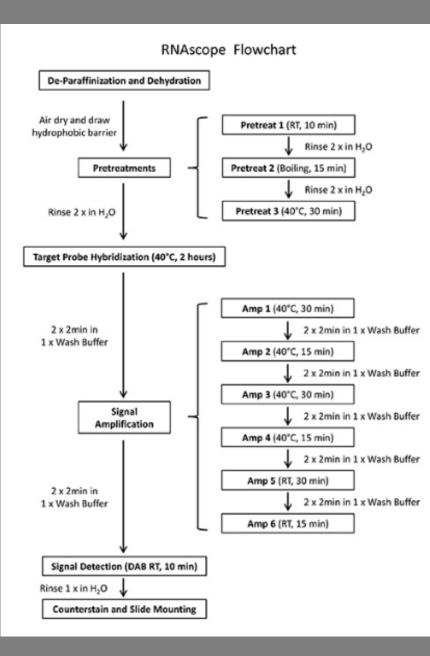


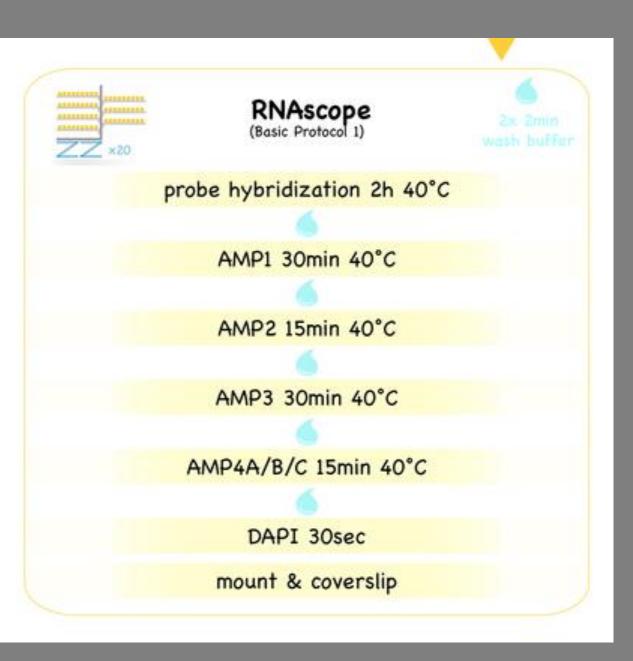


# After hybridization RT, 2 min in 1xWash Buffer Store @RT, O/N in 5xSSC







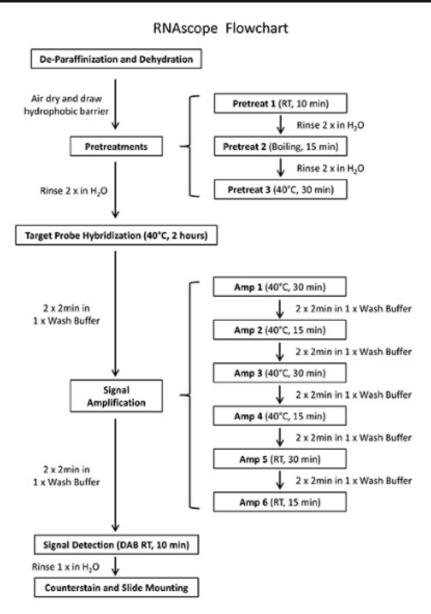


probe channel	Channel 1 (C1)	Channel 2 (C2)	Channel 3 (C3
channel sensitivity	highest	weakest	high
cell type analysis of arget gene expression	gene of interest	cell type marker 1 (e.g. vGLUT1/2)	cell type marker 2 (e.g. GAD1/2)
AMP4A	Alexa488	Atto550	Atto647
AMP4B (recommended)	Atto550	Alexa488	Atto647
AMP4C	Atto550	Atto647	Alexa488

#### Figure 2.

Multiplexing of the three channels in RNAscope. Detected fluorophores of the three channels (C1, C2, C3) can be adjusted by distinct amplification solution 4 (AMP4A, AMP4B, AMP4C). The sensitivity of the three channels is C1 > C2 >> C3. Therefore, we recommend examining the expression of a target gene (lowest expected expression) in different cell types using a channel 1 (C1) probe against this gene of interest and cell type-specific marker genes in Channel 2 & 3.

# Flowchart of RNAscope assay

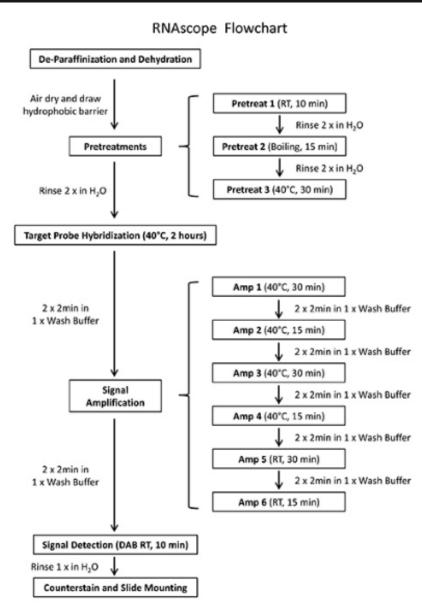


#### Signal Detection:

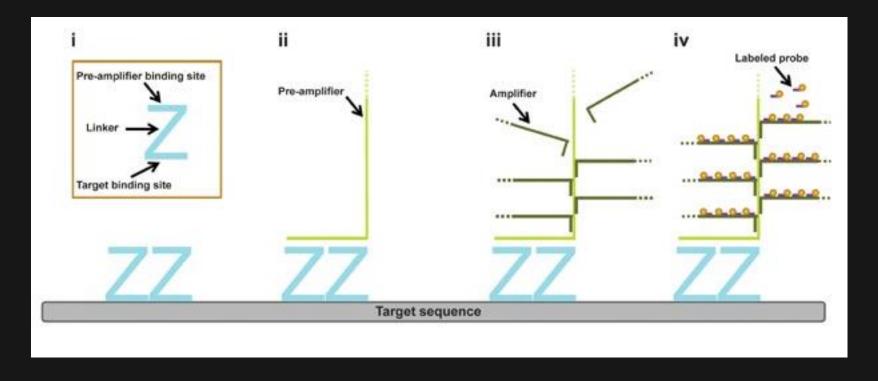
 Incubate tissue sections with 1:1
DAB Mixture by mixing equal volume of Brown-A and Brown-B for 10 min at RT, rinse twice in dH<sub>2</sub>O.

 Stop the substrate reaction to prevent overproduction of a colored precipitate. When the enzyme substrate is added, you need to observe the reaction very often.

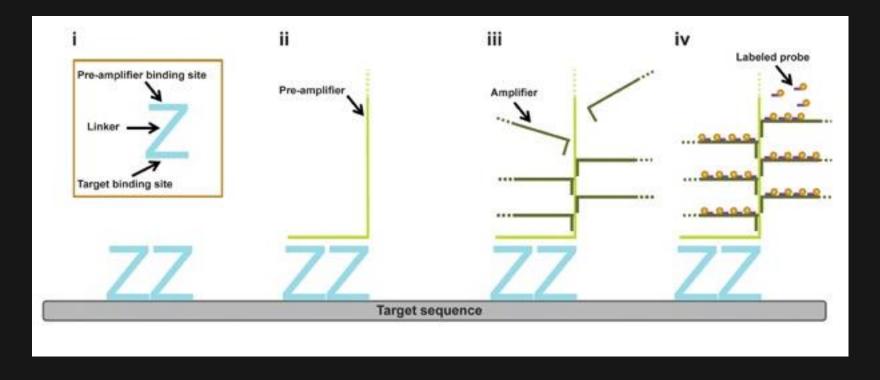
# Flowchart of RNAscope assay



- Fix tissue specimen with 4% PFA for 24 hr.
- 5 um thickness
- Slides are stored at 4 C
- The mounted tissue slide should be baked in a at 60 C prior to the assay.
- Pretreatment: for bone and cartilage: Hybridization of multiple amplifiers to preamplifier.
- Pre-warm target probes at 40 C for 10 min.



- (i) Hybridization of each Z-shaped target probe hybridizes to the RNA target.
- (ii) Hybridization of the pre-amplifier to the upper portion of the Z-probe pairs.
- (iii) Hybridization of multiple amplifiers to the preamplifier.
- (iv) Hybridization of multiple labeled probes to amplifier.



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A@D

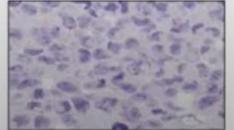
#### Current State of the Art Tissue Based RNA Analysis with RNAscope Technology and Quantitative Analysi DATA ANALYSIS: SEMI-QUANTITATIVE METHOD

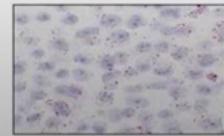
Semi-quantitative image analysis

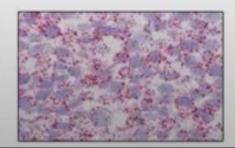
Score assigned to a sample based on average number of target dots/cell performed manually

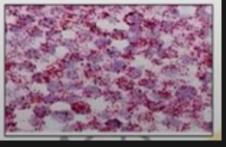
Score	Criteria		
0	No staining or <1 dot/ 10 cells		
1	1-3 dots/cell		
2	4-9 dots/cell. None or very few dot clusters		
3	10-15 dots/cell and <10% dots are in clusters		
4	>15 dots/cell and >10% dots are in clusters		

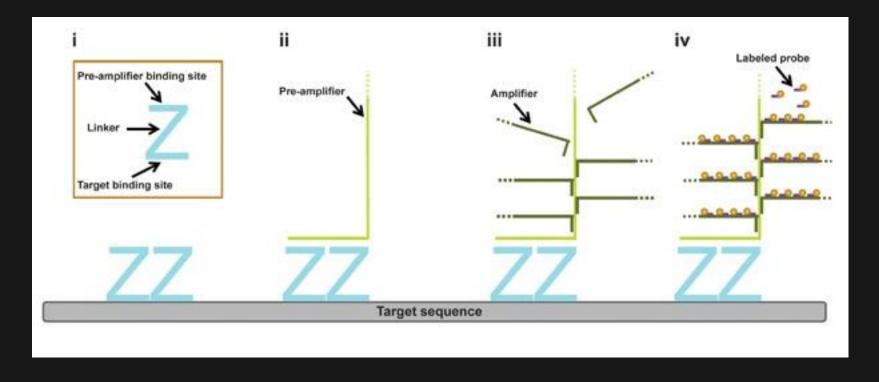




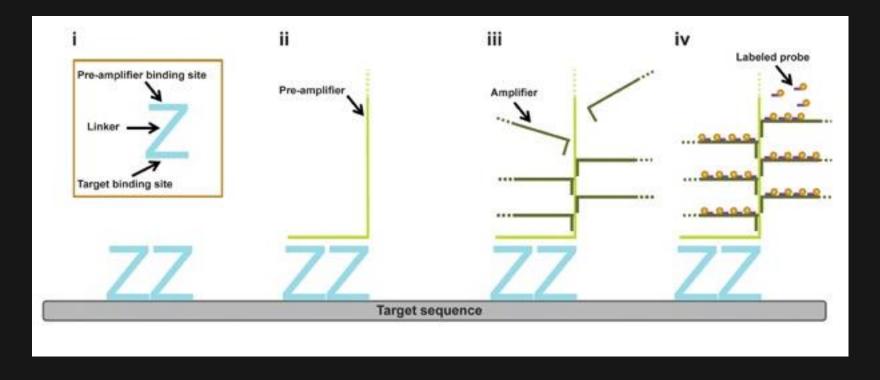




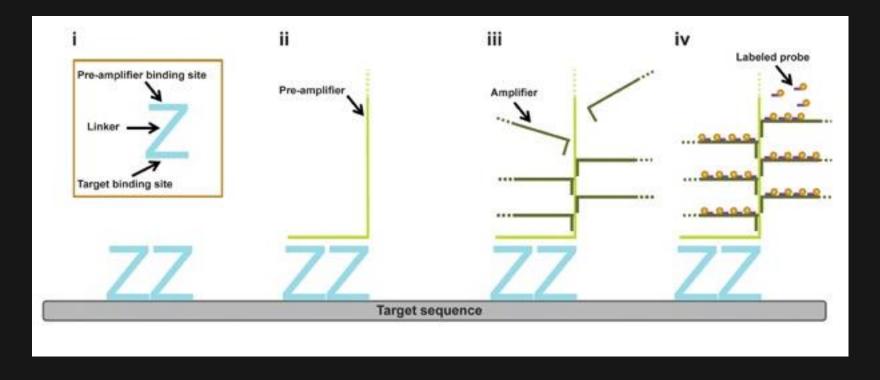




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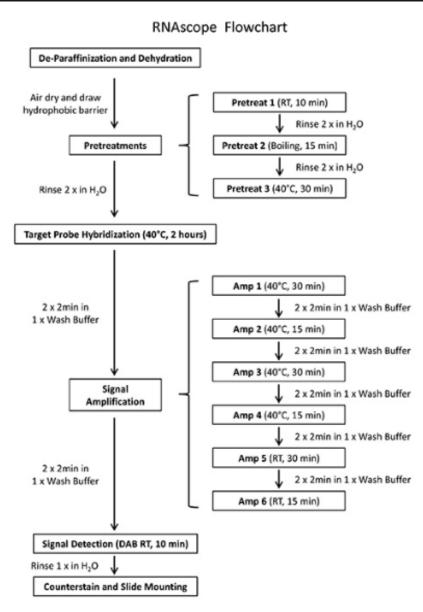


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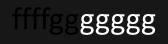


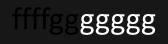
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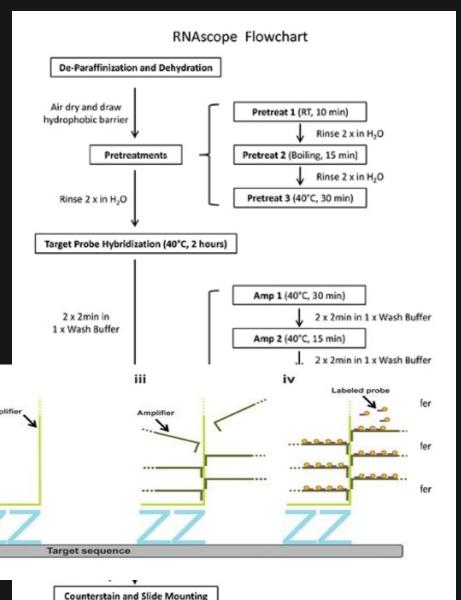
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- Pre-warm target probes at 40 C for 10 min.







## • Signal Amplification is achieved through Amp1-

- Amp1: **Pre-amplifier** binds to the upper region of each double Z probe pair, 40°C, 30min
- Amp2: Background reducer, 40°C, 15min
- Amp3: **Amplifiers** bind to the binding site of Preamplifier, 40°C, 30min
- Amp4: Background reducer, 40°C, 15min
- Amp5: **Amplifiers** bind to the binding site of Preamplifier, RT, 30min or 2hr
- Amp5: **RT**, 30min or 2hr
- Amp6: **RT**, 15min

**6**.