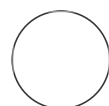




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Spatial Transcriptomics for FFPE utilizing 10x Genomics Visium

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ABSTRACT

To detect gene expression spatially mapped onto formalin-fixed paraffin-embedded tissue samples adapted from the 10x Genomics Visium FFPE v1 protocol.

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Protocol status: Working
We use this protocol and it's working

Created: Jan 18, 2023**Last Modified:** Jan 26, 2023**PROTOCOL integer ID:**
75508

Quality Assessment

- 1 Follow the tissue preparation guide as outlined below to determine if your starting FFPE tissue block is acceptable for the 10x Visium platform
Start [here](#)
Utilize Visium Tissue Section Test Slide (PN-2000460) to do a quality assessment of each sample
Be sure to trim a few sections and place in a 1.5ml tube for RNA quality check (no more than 40µm)
Follow the steps starting on pg. 10 of the above user guide
- 2 Perform all deparaffinization/staining/imaging steps utilizing the test slide
Follow the protocol as described by 10x [here](#)
- 3 Perform RNA extraction using the Qiagen RNeasy FFPE kit (Cat no. 73504)
Follow the steps outlined in the RNeasy Handbook which can be found below
 HB-0375-006_HB_RNeasy_FFPE_0721_WW (1).pdf
Start with your pre-cut sections in a 1.5ml tube on pg. 15
- 4 After RNA extraction test the concentration of your samples using a nanodrop
Determine the DV200 value of your samples utilizing the Agilent 4200 TapeStation system or similar quantitative assessment
If utilizing Agilent TapeStation to prepare your RNA samples utilize this [guide](#)
For first time analysis of DV200 you can set up your parameters as described [here](#)

Visium FFPE v1

- 5 If the tissue adheres to the slide and has a DV200 value >50% that sample can be used for the Visium Spatial Transcriptomics platform
- 6 Utilize the CG000408 linked above to section onto the Visium Spatial Gene Expression Slide (PN-2000233)
- 7 After sectioning start the protocol utilizing CG000409 linked above to perform deparaffinization/staining/imaging steps
Be sure to either store your slide or perform decrosslinking to proceed with the rest of the protocol outlined in the next step
- 8 If decrosslinking is complete, utilize the following user guide outlined [here](#)

Start probe hybridization on pg. 39

- 9 Follow the sequencing requirements on pg. 60 to generate your Visium libraries