## RUO

### **Technical sheet**

# Idylla™ BRAF Mutation Assay

The Idylla<sup>TM</sup> BRAF Mutation Assay, performed on the Biocartis Idylla<sup>TM</sup> System, is a molecular assay for the qualitative detection of V600E/E2/D and V600K/R/M mutations in codon 600 of the BRAF gene.

The Idylla<sup>TM</sup> BRAF Mutation Assay, from sample-to-result, starts with formalin-fixed paraffin-embedded (FFPE) tissue to liberate DNA for subsequent real-time PCR amplification and detection.



### **Features**

| BRAF mutation detection                      |   |                       |  |
|--|---|-----------------------|--|
| Codon 600                                    | BRAF V600E                                | (c.1799T>A)           |  |
|  | BRAF V600E2                               | (c.1799_1800delinsAA) |  |
|  | BRAF V600D                                | (c.1799_1800delinsAT; |  |
|  |   | c.1799_1800delinsAC)  |  |
|  | BRAF V600K                                | (c.1798_1799delinsAA) |  |
|  | BRAF V600R                                | (c.1798_1799delinsAG) |  |
|  | BRAF V600M                                | (c.1798G>A)           |  |
|  | BRAF Wild Type                            | (c.1799T)             |  |
| RNaseP (acting as Sample Processing Control) |   |                       |  |
| Specimen requirements                        |   |                       |  |
| Sample Type                                  | FFPE tissue sections (5 to 10µm)          |                       |  |
| Neoplastic cells                             | ≥50%, if less macrodissection is required |                       |  |
| Τ.   | 50-600mm² (5µm                            | 50-600mm² (5µm)       |  |
| Tissue area                                  | 25-300mm² (10µr                           | 25-300mm² (10μm)      |  |
| Total turnaround time                        |   |                       |  |
| Time   | 90 minutes                                | 90 minutes            |  |
| Result Reporting                             |   |                       |  |
| Report                                       | Qualitative genotype call                 |                       |  |

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