

Report panelDF

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| Full name: | Hereditary colorectal cancer (Adenomatous polyposis, Lynch, Peutz- Jeghers, juvenile polyposis, PPAP, NAP) - UGent |
| Laboratory: | <u>Centrum Medische Genetica - UZ Gent</u> |
| Created: | 04 Jun 2019 - 10:59 |
| Changed: | 13 Dec 2022 - 12:11 |

Related Diseases

- Familial adenomatous polyposis
- Lynch syndrome
- MSH3-related attenuated familial adenomatous polyposis
- MUTYH-related attenuated familial adenomatous polyposis
- NTHL1-related attenuated familial adenomatous polyposis
- Peutz-Jeghers syndrome
- Polymerase proofreading-related adenomatous polyposis

Related Analytes

| GENE | % OF CODING SEQUENCE SUFFICIENTLY COVERED TO DETECT HETEROZYGOUS MUTATIONS | COPY NUMBER VARIATION | COMMENTS |
|---------------|--|-----------------------|--|
| <u>APC</u> | 100.00 | 1 | |
| <u>AXIN2</u> | 100.00 | 0 | |
| <u>BMPR1A</u> | 100.00 | 1 | |
| <u>EPCAM</u> | 100.00 | 1 | CNV for EPCAM exon 7-9 and region between EPCAM and MSH2 |
| <u>GREM1</u> | 100.00 | 1 | CNV for recurrent 40kb duplication |
| <u>MLH1</u> | 100.00 | 1 | |
| <u>MSH2</u> | 100.00 | 1 | |

| GENE | % OF CODING SEQUENCE SUFFICIENTLY COVERED TO DETECT HETEROZYGOUS MUTATIONS | COPY NUMBER VARIATION | COMMENTS |
|--------------|--|-----------------------|----------|
| <u>MSH3</u> | 100.00 | 0 | |
| <u>MSH6</u> | 100.00 | 1 | |
| <u>MUTYH</u> | 100.00 | 1 | |
| <u>NTHL1</u> | 100.00 | 0 | |
| <u>PMS2</u> | 100.00 | 1 | |
| <u>POLD1</u> | 100.00 | 0 | |
| <u>POLE</u> | 100.00 | 1 | |
| <u>PTEN</u> | 100.00 | 1 | |
| <u>RNF43</u> | 100.00 | 0 | |
| <u>SMAD4</u> | 100.00 | 1 | |
| <u>STK11</u> | 100.00 | 1 | |