

## ForeSENTIA GENES AND GENETIC ALTERATIONS TESTED PER PANEL

### GENES TESTED

The following table indicates the genes tested per panel.

<b>Pan-Cancer Core</b> 80 genes	AKT1, ALK, APC, AR, ARAF, ATM, ATRX, BARD1, BRAF, BRCA1, BRCA2, BRIP1, CDH1, CDKN2A, CHEK2, CIC, CTNNB1, DDR2, DICER1, EGFR, ERBB2, ERBB3, ESR1, FBXW7, FGFR1, FGFR2, FGFR3, FLT3, FOXA1, FOXL2, FUBP1, GATA3, GNA11, GNAQ, GNAS, H3F3A, IDH1, IDH2, JAK2, KEAP1, KIT, KRAS, MAP2K1, MAP3K1, MET, MLH1, MRE11, MSH2, MSH6, MTOR, MYC, MYCN, NBN, NF1, NPM1, NRAS, NTRK1, NTRK2, NTRK3, PALB2, PDGFRA, PIK3CA, PIK3CB, PMS2, POLE, PTEN, RAD51C, RAD51D, RAF1, RB1, RET, ROS1, RUNX1, SMAD4, SPOP, STK11, TERT, TMPRSS2, TP53, 1p/19q codeletion
<b>Breast/Gynecological</b> 48 genes	AKT1, ATM, BARD1, BRAF, BRCA1, BRCA2, BRIP1, CHEK2, CTNNB1, DICER1, EGFR, ERBB2, ERBB3, ESR1, FBXW7, FGFR1, FGFR2, FGFR3, FOXA1, FOXL2, GATA3, KIT, KRAS, MAP3K1, MET, MLH1, MRE11, MSH2, MSH6, MTOR, NBN, NRAS, NTRK1, NTRK2, NTRK3, PALB2, PIK3CA, PIK3CB, PMS2, POLE, PTEN, RAD51D, RAF1, RET, RUNX1, SMAD4, TP53
<b>Colorectal</b> 34 genes	AKT1, APC, ATM, BRAF, BRCA1, BRCA2, CTNNB1, EGFR, ERBB2, FBXW7, FGFR1, FGFR2, FGFR3, GNAS, KRAS, MET, MLH1, MSH2, MSH6, MTOR, NRAS, NTRK1, NTRK2, NTRK3, PALB2, PDGFRA, PIK3CA, PIK3CB, PMS2, POLE, PTEN, RAF1, SMAD4, TP53
<b>Glioma</b> 22 genes	ATRX, BRAF, CDKN2A, CIC, CTNNB1, EGFR, FGFR3, FUBP1, H3F3A, IDH1, IDH2, MET, MYC, MYCN, NF1, NTRK1, NTRK2, NTRK3, POLE, PTEN, TERT, TP53, 1p/19q codeletion
<b>Lung (NSCLC)</b> 36 genes	AKT1, ALK, APC, ARAF, ATM, BRAF, BRCA2, CTNNB1, DDR2, EGFR, ERBB2, ERBB3, ERBB4, FBXW7, FGFR1, FGFR2, FGFR3, JAK2, KEAP1, KRAS, MAP2K1, MET, NRAS, NTRK1, NTRK2, NTRK3, PDGFRA, PIK3CA, POLE, PTEN, RAF1, RET, ROS1, SMAD4, STK11, TP53
<b>Melanoma</b> 22 genes	AKT1, ALK, BRAF, CTNNB1, ERBB2, GNA11, GNAQ, KIT, KRAS, MAP2K1, MYC, NF1, NRAS, NTRK1, NTRK2, NTRK3, PIK3CA, POLE, PTEN, RET, ROS1, TP53
<b>Prostate</b> 36 genes	AKT1, APC, AR, ATM, BARD1, BRAF, BRCA1, BRCA2, CHEK2, CTNNB1, ERBB2, FGFR1, FGFR2, FGFR3, FOXA1, MLH1, MSH2, MSH6, MYC, MYCN, NRAS, NTRK1, NTRK2, NTRK3, PALB2, PIK3CA, PIK3CB, PMS2, POLE, PTEN, RAD51C, RAD51D, RB1, SPOP, TMPRSS2, TP53

### GENETIC ALTERATIONS TESTED

The following table shows the types of genetic alterations and coding exon coverage for each gene.

<b>AKT1</b> Exon 4 (NM_001382430.1)	<b>ALK</b> Exons 18-20, 22-23 (NM_004304.5)	<b>APC</b> Full exonic coverage (NM_000038.6)	<b>AR</b> Exons 1-2, 4-8 (NM_000044.6)	<b>ARAF</b> Exon 7 (NM_001654.5)	<b>ATM</b> Full exonic coverage (NM_000051.4)
▲	▲ ■	▲	▲ ●	▲	▲
<b>ATRX</b> Exons 5, 7, 9, 11, 13-14, 17, 20-22, 29-31, 35 (NM_000489.6)	<b>BARD1</b> Full exonic coverage (NM_000465.4)	<b>BRAF</b> Exons 9-12, 15 (NM_004333.6)	<b>BRCA1</b> Full exonic coverage (NM_007294.4)	<b>BRCA2</b> Full exonic coverage (NM_000059.4)	<b>BRIP1</b> Full exonic coverage (NM_032043.3)
▲	▲	▲ ■	▲	▲	▲
<b>CDH1</b> Full exonic coverage (NM_004360.5)	<b>CDKN2A</b> Full exonic coverage (NM_000077.5) (NM_058195.4)	<b>CHEK2</b> Exons 1-10 (NM_007194.4)	<b>CIC</b> Exon 19, 21 (NM_0015125.4)	<b>CTNNB1</b> Exon 3 (NM_001904.4)	<b>DDR2</b> Exon 17 (NM_006182.4)
▲	●	▲	▲	▲	▲
<b>DICER1</b> Exons 2-26 (NM_177438.3)	<b>EGFR</b> Exons 2-5, 8-10, 12, 14-15, 17-24, 27-28 (NM_005228.5)	<b>ERBB2</b> Exons 3-4, 7-8, 10, 19-24 (NM_004448.4)	<b>ERBB3</b> Exons 3, 6-11, 20-21 (NM_001982.4)	<b>ERBB4</b> Exons 7, 15, 19, 23 (NM_005235.3)	<b>ESR1</b> Exons 2-5, 7-8 (NM_000125.4)
▲	▲ ●	▲ ●	▲	▲	▲ ●

Selected non-coding regions which are covered by the test are indicated above.

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▲ Single Nucleotide Variant / Indels

● Copy Number Alterations

■ Rearrangements

<b>FBXW7</b> Exons 6-7, 9-14 (NM_001349798.2)	<b>FGFR1</b> Exon 5-9, 12-13, 16, 18 (NM_023110.3)	<b>FGFR2</b> Exons 2-9, 11, 13-18 (NM_000141.5)	<b>FGFR3</b> Exons 4, 7-8, 11-12, 16-18 (NM_000142.5)	<b>FLT3</b> Exon 20 (NM_004119.3)	<b>FOXA1</b> Exon 2 (NM_004496.5)
▲	Selected non-coding regions covered ●	Selected non-coding regions covered ●	Selected non-coding regions covered ● ■	▲	▲
<b>FOXL2</b> Exon 1 (NM_023067.4)	<b>FUBP1</b> Exon 14 (NM_003902.5)	<b>GATA3</b> Exons 5-6 (NM_001002295.2)	<b>GNA11</b> Exons 4-5 (NM_002067.5)	<b>GNAQ</b> Exons 5, 7 (NM_002072.5)	<b>GNAS</b> Exon 8 (NM_000516.7)
▲	▲	▲	▲	▲	▲
<b>H3F3A</b> Exon 2 (NM_002107.7)	<b>IDH1</b> Exon 4 (NM_005896.4)	<b>IDH2</b> Exon 4 (NM_002168.4)	<b>JAK2</b> Exon 14 (NM_004972.4)	<b>KEAP1</b> Exons 2-4 (NM_203500.2)	<b>KIT</b> Exons 2-21 (NM_000222.3)
▲	▲	▲	▲	▲	▲ ●
<b>KRAS</b> Exons 2-4 (NM_004985.5)	<b>MAP2K1</b> Exons 2-3, 6-7 (NM_002755.4)	<b>MAP3K1</b> Exons 5-6, 16-18 (NM_005921.2)	<b>MET</b> Exons 2-21 (NM_000245.4)	<b>MLH1</b> Full exonic coverage (NM_000249.4)	<b>MRE11</b> Exons 1-18, 20 (NM_005591.4)
▲ ●	▲	▲	▲ ●	Selected non-coding regions covered ▲	Selected non-coding regions covered ▲
<b>MSH2</b> Full exonic coverage (NM_000251.3)	<b>MSH6</b> Full exonic coverage (NM_000179.3)	<b>MTOR</b> Exons 43, 47, 53, 56 (NM_004958.4)	<b>MYC</b> Exons 1, 3 (NM_002467.6)	<b>MYCN</b> Exon 3 (NM_005378.6)	<b>NBN</b> Full exonic coverage (NM_002485.5)
Selected non-coding regions covered ▲	▲	▲	●	●	▲
<b>NF1</b> Exons 6, 12, 17-18, 21-22, 25, 27-28, 34, 37, 40, 44-47, 49, 53 (NM_001042492.3)	<b>NPM1</b> Exon 11 (NM_002520.7)	<b>NRAS</b> Exons 2-4 (NM_002524.5)	<b>NTRK1</b> Exons 7-15 (NM_002529.4)	<b>NTRK2</b> Exons 10-11, 13-14 (NM_006180.6)	<b>NTRK3</b> Exons 14-15 (NM_001012338.3)
▲	▲	▲	▲ ■	Selected non-coding regions covered ■	Selected non-coding regions covered ■
<b>PALB2</b> Full exonic coverage (NM_024675.4)	<b>PDGFRA</b> Exon 18 (NM_006206.6)	<b>PIK3CA</b> Exons 2-6, 8, 10, 15-17, 19-21 (NM_006218.4)	<b>PIK3CB</b> Exons 13, 15, 24 (NM_006219.3)	<b>PMS2</b> Exons 6-8, 10 (NM_000535.7)	<b>POLE</b> Exons 2-49 (NM_006231.4)
Selected non-coding regions covered ▲	▲	▲ ●	▲	▲	Selected non-coding regions covered ▲
<b>PTEN</b> Full exonic coverage (NM_000314.8)	<b>RAD51C</b> Full exonic coverage (NM_058216.3)	<b>RAD51D</b> Full exonic coverage (NM_002878.4)	<b>RAF1</b> Exon 7 (NM_002880.4)	<b>RB1</b> Exons 1-13, 16-27 (NM_000321.3)	<b>RET</b> Full exonic coverage (NM_020975.6)
Selected non-coding regions covered ▲ ●	Selected non-coding regions covered ▲	Selected non-coding regions covered ▲	▲	Selected non-coding regions covered ●	Selected non-coding regions covered ▲ ■
<b>ROS1</b> Exons 32-37 (NM_001378902.1)	<b>RUNX1</b> Exon 5 (NM_001754.4)	<b>SMAD4</b> Full exonic coverage (NM_005359.6)	<b>SPOP</b> Exons 4-5 (NM_001007228.2)	<b>STK11</b> Full exonic coverage (NM_000455.5)	<b>TERT</b> Exon 1 (NM_198253.3)
Selected non-coding regions covered ■	▲	Selected non-coding regions covered ▲	▲	Selected non-coding regions covered ▲	Selected non-coding regions covered ▲
<b>TMPPRSS2</b> Exons 2-4 (NM_005656.4)	<b>TP53</b> Full exonic coverage (NM_000546.6)				
Selected non-coding regions covered ■	Selected non-coding regions covered ▲ ●				

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