

Batch Certificate

For Research Use Only

PRODUCT INFORMATION AND QUALITY CONTROL

NAME OF PRODUCT	PIK3CA-T1 mutations 12.5% AF cfDNA- PIK3CA E542K & Q546R
DESCRIPTION	PIK3CA-T1 mutations 12.5% AF cfDNA- PIK3CA E542K & Q546R is highly characterized human DNA from cell lines.
CATALOG NUMBER	SID-000099
BATCH NUMBER	00043
MANUFACTURING CONDITIONS	<ul style="list-style-type: none"> • Manufactured and sealed in class 2 safety cabinet • At room temperature
PACKAGE SIZE	<ul style="list-style-type: none"> • 2D barcoded tube with screw cap
PACKAGE TYPE	<ul style="list-style-type: none"> • Material: Polypropylen (PP)
DATE OF MANUFACTURE	13.03.2020
EXPIRY DATE	12.03.2022
CONCENTRATION	20 ng/μl (dsDNA)
QUANTITY	400 ng (dsDNA)
NOMINAL VOLUME	25 μl; (508 ng)
MUTATION	PIK3CA p.E542K (COSM760*, COSV55873227*, substitution, c.1624G>A, Exon 9) PIK3CA p.Q546R (COSM12459*, COSV55876869* substitution, c.1637A>G, Exon 9) <small>* GRCh38 COSMIC v90</small>
ALLELE FREQUENCY	12.5%
QUALITY	DNA quantity metrologically traceable to internationally certified reference material ¹ The copy number values are metrologically traceable to the natural units count 1 and ratio 1 and International System of Units (SI) derived units of volume.
STORAGE CONDITIONS	+2 - +8 °C
MANUFACTURING AND QUALITY CONTROL SITES	SensID GmbH Schillingallee 68, 18057 Rostock, Germany

¹ ERM_AD442K
Phone: +49 (0) 381 377 182 01

TEST METHOD AND ACCEPTANCE CRITERIA	Quality Control	Test Method	Acceptance criteria
	Fragmentation	Fragment Length Analysis Agilent High Sensitivity DNA Kit (Agilent Technologies)	peak size 167 bp ± 10% (151 bp – 181 bp)
	Quantification	Total DNA measurement: Spectrophotometry ssDNA [ng/μl] = (A260-A320)*38 ^{2,3} dsDNA measurement: Qubit dsDNA BR Assay Kit (Invitrogen)	Total DNA: n.a. ⁴ dsDNA: 17.5 – 22.5 ng/μl
	Allele Frequency	ddPCR Analysis using BioRad QX200™ System	AF 12.5% ±20% (10.0-15.0%)

RESULTS OF ANALYSIS	Result	PASS/FAIL									
Fragmentation	180 bp	PASS									
Quantity	28.7 ng/μl (total DNA) 20.3 ng/μl (dsDNA)	PASS									
Allele Frequency	<table border="1"> <thead> <tr> <th>Mutation</th><th>AF in %</th><th>PASS/FAIL</th></tr> </thead> <tbody> <tr> <td>PIK3CA E542K</td><td>10.2</td><td>PASS</td></tr> <tr> <td>PIK3CA Q546R</td><td>10.1</td><td>PASS</td></tr> </tbody> </table>	Mutation	AF in %	PASS/FAIL	PIK3CA E542K	10.2	PASS	PIK3CA Q546R	10.1	PASS	
Mutation	AF in %	PASS/FAIL									
PIK3CA E542K	10.2	PASS									
PIK3CA Q546R	10.1	PASS									

COMMENTS/REMARKS	Additional information:									
	Copy numbers (CN) of the respective measurements									
	<table><tr><th>Mutation</th><th>CN wt⁵/μl</th><th>CN mut⁶/μl</th></tr><tr><td>PIK3CA E542K</td><td>5830</td><td>660</td></tr><tr><td>PIK3CA Q546R</td><td>6637</td><td>744</td></tr></table>	Mutation	CN wt ⁵ /μl	CN mut ⁶ /μl	PIK3CA E542K	5830	660	PIK3CA Q546R	6637	744
	Mutation	CN wt ⁵ /μl	CN mut ⁶ /μl							
	PIK3CA E542K	5830	660							
PIK3CA Q546R	6637	744								
<i>Table 1 indicates the values of the QC assays performed by SensID GmbH with a DNA input of ~40 ng. The value for the respective mutation results from the mean value of five measured replicates (CN values are rounded). CN concentration values per microliter (μl), are based on droplet digital (ddPCR) assay counts dilution factors, and droplet volume measurements. The detection of the amount of CNs may vary depending on the assay used. Therefore, due to assay properties, there may be deviations in the observed number of copies and allele frequencies compared to the values given here.</i>										

² Protocol NK603 – Community Reference Laboratory for GM Food and Feed

³ Measured before filling in product tube

⁴ not applicable

⁵ Wild Type

⁶ Mutation

Batch Certificate

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PRODUCT INFORMATION AND QUALITY CONTROL

NAME OF PRODUCT	PIK3CA-11 mutations 12.5% AF cfDNA- PIK3CA E545D & H1047L
DESCRIPTION	PIK3CA-11 mutations 12.5% AF cfDNA- PIK3CA E545D & H1047L is highly characterized human DNA from cell lines.
CATALOG NUMBER	SID-000099
BATCH NUMBER	00044
MANUFACTURING CONDITIONS	<ul style="list-style-type: none"> • Manufactured and sealed in class 2 safety cabinet • At room temperature
PACKAGE SIZE	<ul style="list-style-type: none"> • 2D barcoded tube with screw cap
PACKAGE TYPE	<ul style="list-style-type: none"> • Material: Polypropylen (PP)
DATE OF MANUFACTURE	13.03.2020
EXPIRY DATE	12.03.2022
CONCENTRATION	20 ng/μl (dsDNA)
QUANTITY	400 ng (dsDNA)
NOMINAL VOLUME	25 μl; (473 ng)
MUTATION	PIK3CA p.E545D (COSM765*, COSV55874040*, substitution, c.1635G>T, Exon 9) PIK3CA p.H1047L (COSM776*, COSV55873401* substitution, c.3140A>T, Exon 20) <small>* GRCh38 COSMIC v90</small>
ALLELE FREQUENCY	12.5%
QUALITY	DNA quantity metrologically traceable to internationally certified reference material ⁷ The copy number values are metrologically traceable to the natural units count 1 and ratio 1 and International System of Units (SI) derived units of volume.
STORAGE CONDITIONS	+2 - +8 °C
MANUFACTURING AND QUALITY CONTROL SITES	SensID GmbH Schillingallee 68, 18057 Rostock, Germany

⁷ ERM_AD442K
Phone: +49 (0) 381 377 182 01

Net: www.sens-id.com
SensID GmbH, Schillingallee 68, 18057 Rostock, Germany

Mail: support@sens-id.com

VAT No: DE305142405, district court: Rostock HRB 14621
CEO: Björn Nowack

TEST METHOD AND ACCEPTANCE CRITERIA	Quality Control	Test Method	Acceptance criteria
	Fragmentation	Fragment Length Analysis Agilent High Sensitivity DNA Kit (Agilent Technologies)	peak size 167 bp ± 10% (151 bp – 181 bp)
	Quantification	Total DNA measurement: Spectrophotometry ssDNA [ng/μl] = (A260-A320)*38 ^{8,9} dsDNA measurement: Qubit dsDNA BR Assay Kit (Invitrogen)	Total DNA: n.a. ¹⁰ dsDNA: 17.5 – 22.5 ng/μl
	Allele Frequency	ddPCR Analysis using BioRad QX200™ System	AF 12.5% ±20% (10.0-15.0%)

RESULTS OF ANALYSIS	Result	PASS/FAIL									
Fragmentation	179 bp	PASS									
Quantity	27.0 ng/μl (total DNA) 18.9 ng/μl (dsDNA)	PASS									
Allele Frequency	<table border="1"> <thead> <tr> <th>Mutation</th><th>AF in %</th><th>PASS/FAIL</th></tr> </thead> <tbody> <tr> <td>PIK3CA E545D</td><td>13.2</td><td>PASS</td></tr> <tr> <td>PIK3CA H1047L</td><td>14.6</td><td>PASS</td></tr> </tbody> </table>	Mutation	AF in %	PASS/FAIL	PIK3CA E545D	13.2	PASS	PIK3CA H1047L	14.6	PASS	
Mutation	AF in %	PASS/FAIL									
PIK3CA E545D	13.2	PASS									
PIK3CA H1047L	14.6	PASS									

COMMENTS/REMARKS	Additional information:									
	Copy numbers (CN) of the respective measurements									
	<table><tr><th>Mutation</th><th>CN wt¹¹/μl</th><th>CN mut¹²/μl</th></tr><tr><td>PIK3CA E545D</td><td>3371</td><td>511</td></tr><tr><td>PIK3CA H1047L</td><td>4111</td><td>704</td></tr></table>	Mutation	CN wt ¹¹ /μl	CN mut ¹² /μl	PIK3CA E545D	3371	511	PIK3CA H1047L	4111	704
	Mutation	CN wt ¹¹ /μl	CN mut ¹² /μl							
	PIK3CA E545D	3371	511							
PIK3CA H1047L	4111	704								
<i>Table 2 indicates the values of the QC assays performed by SensID GmbH with a DNA input of ~40 ng. The value for the respective mutation results from the mean value of five measured replicates (CN values are rounded). CN concentration values per microliter (μl), are based on droplet digital (ddPCR) assay counts dilution factors, and droplet volume measurements. The detection of the amount of CNs may vary depending on the assay used. Therefore, due to assay properties, there may be deviations in the observed number of copies and allele frequencies compared to the values given here.</i>										

⁸ Protocol NK603 – Community Reference Laboratory for GM Food and Feed

⁹ Measured before filling in product tube

¹⁰ not applicable

¹¹ Wild Type

¹² Mutation

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PRODUCT INFORMATION AND QUALITY CONTROL

NAME OF PRODUCT	PIK3CA-11 mutations 12.5% AF cfDNA- PIK3CA E545G & H1047R
DESCRIPTION	PIK3CA-11 mutations 12.5% AF cfDNA- PIK3CA E545G & H1047R is highly characterized human DNA from cell lines.
CATALOG NUMBER	SID-000099
BATCH NUMBER	00045
MANUFACTURING CONDITIONS	<ul style="list-style-type: none"> • Manufactured and sealed in class 2 safety cabinet • At room temperature
PACKAGE SIZE	<ul style="list-style-type: none"> • 2D barcoded tube with screw cap
PACKAGE TYPE	<ul style="list-style-type: none"> • Material: Polypropylen (PP)
DATE OF MANUFACTURE	13.03.2020
EXPIRY DATE	12.03.2022
CONCENTRATION	20 ng/μl (dsDNA)
QUANTITY	400 ng (dsDNA)
NOMINAL VOLUME	25 μl; (450 ng)
MUTATION	PIK3CA p.E545G (COSM764*, COSV55873220*, substitution, c.1634A>G, Exon 9) PIK3CA p.H1047R (COSM775*, COSV55873195* substitution, c.3140A>G, Exon 20) <small>* GRCh38 COSMIC v90</small>
ALLELE FREQUENCY	12.5%
QUALITY	DNA quantity metrologically traceable to internationally certified reference material ¹³ The copy number values are metrologically traceable to the natural units count 1 and ratio 1 and International System of Units (SI) derived units of volume.
STORAGE CONDITIONS	+ 2 - 8 °C
MANUFACTURING AND QUALITY CONTROL SITES	SensID GmbH Schillingallee 68, 18057 Rostock, Germany

¹³ ERM_AD442K

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VAT No: DE305142405, district court: Rostock HRB 14621

CEO: Björn Nowack

TEST METHOD AND ACCEPTANCE CRITERIA	Quality Control	Test Method	Acceptance criteria
	Fragmentation	Fragment Length Analysis Agilent High Sensitivity DNA Kit (Agilent Technologies)	peak size 167 bp ± 10% (151 bp – 181 bp)
	Quantification	Total DNA measurement: Spectrophotometry ssDNA [ng/μl] = (A260-A320)*38 ^{14,15} dsDNA measurement: Qubit dsDNA BR Assay Kit (Invitrogen)	Total DNA: n.a. ¹⁶ dsDNA: 17.5 – 22.5 ng/μl
	Allele Frequency	ddPCR Analysis using BioRad QX200™ System	AF 12.5% ±20% (10.0-15.0%)

RESULTS OF ANALYSIS	Result	PASS/FAIL									
Fragmentation	179 bp	PASS									
Quantity	23.9 ng/μl (total DNA) 18.0 ng/μl (dsDNA)	PASS									
Allele Frequency	<table border="1"> <thead> <tr> <th>Mutation</th><th>AF in %</th><th>PASS/FAIL</th></tr> </thead> <tbody> <tr> <td>PIK3CA E545G</td><td>12.1</td><td>PASS</td></tr> <tr> <td>PIK3CA H1047R</td><td>12.9</td><td>PASS</td></tr> </tbody> </table>	Mutation	AF in %	PASS/FAIL	PIK3CA E545G	12.1	PASS	PIK3CA H1047R	12.9	PASS	
Mutation	AF in %	PASS/FAIL									
PIK3CA E545G	12.1	PASS									
PIK3CA H1047R	12.9	PASS									

COMMENTS/REMARKS	Additional information:									
	Copy numbers (CN) of the respective measurements									
	<table><tr><th>Mutation</th><th>CN wt¹⁷/μl</th><th>CN mut¹⁸/μl</th></tr><tr><td>PIK3CA E545G</td><td>4271</td><td>587</td></tr><tr><td>PIK3CA H1047R</td><td>3805</td><td>565</td></tr></table>	Mutation	CN wt ¹⁷ /μl	CN mut ¹⁸ /μl	PIK3CA E545G	4271	587	PIK3CA H1047R	3805	565
	Mutation	CN wt ¹⁷ /μl	CN mut ¹⁸ /μl							
	PIK3CA E545G	4271	587							
PIK3CA H1047R	3805	565								
<i>Table 3 indicates the values of the QC assays performed by SensID GmbH with a DNA input of ~40 ng. The value for the respective mutation results from the mean value of five measured replicates (CN values are rounded). CN concentration values per microliter (μl), are based on droplet digital (ddPCR) assay counts dilution factors, and droplet volume measurements. The detection of the amount of CNs may vary depending on the assay used. Therefore, due to assay properties, there may be deviations in the observed number of copies and allele frequencies compared to the values given here.</i>										

¹⁴ Protocol NK603 – Community Reference Laboratory for GM Food and Feed

¹⁵ Measured before filling in product tube

¹⁶ not applicable

¹⁷ Wild Type

¹⁸ Mutation

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PRODUCT INFORMATION AND QUALITY CONTROL

NAME OF PRODUCT	PIK3CA-T1 mutations 12.5% AF cfDNA- PIK3CA E545K & H1047Y
DESCRIPTION	PIK3CA-T1 mutations 12.5% AF cfDNA- PIK3CA E545K & H1047Y is highly characterized human DNA from cell lines.
CATALOG NUMBER	SID-000099
BATCH NUMBER	00046
MANUFACTURING CONDITIONS	<ul style="list-style-type: none"> • Manufactured and sealed in class 2 safety cabinet • At room temperature
PACKAGE SIZE	<ul style="list-style-type: none"> • 2D barcoded tube with screw cap
PACKAGE TYPE	<ul style="list-style-type: none"> • Material: Polypropylen (PP)
DATE OF MANUFACTURE	13.03.2020
EXPIRY DATE	12.03.2022
CONCENTRATION	20 ng/μl (dsDNA)
QUANTITY	400 ng (dsDNA)
NOMINAL VOLUME	25 μl; (485 ng)
MUTATION	PIK3CA p.E545K (COSM763*, COSV55873239*, substitution, c.1633G>A, Exon 9) PIK3CA p.H1047Y (COSM774*, COSV55876499* substitution, c.3139C>T, Exon 20) <small>* GRCh38 COSMIC v90</small>
ALLELE FREQUENCY	12.5%
QUALITY	DNA quantity metrologically traceable to internationally certified reference material ¹⁹ The copy number values are metrologically traceable to the natural units count 1 and ratio 1 and International System of Units (SI) derived units of volume.
STORAGE CONDITIONS	+2 - +8 °C
MANUFACTURING AND QUALITY CONTROL SITES	SensID GmbH Schillingallee 68, 18057 Rostock, Germany

¹⁹ ERM_AD442K

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CEO: Björn Nowack

TEST METHOD AND ACCEPTANCE CRITERIA	Quality Control	Test Method	Acceptance criteria
	Fragmentation	Fragment Length Analysis Agilent High Sensitivity DNA Kit (Agilent Technologies)	peak size 167 bp ± 10% (151 bp – 181 bp)
	Quantification	Total DNA measurement: Spectrophotometry ssDNA [ng/μl] = (A260-A320)*38 ^{20, 21} dsDNA measurement: Qubit dsDNA BR Assay Kit (Invitrogen)	Total DNA: n.a. ²² dsDNA: 17.5 – 22.5 ng/μl
	Allele Frequency	ddPCR Analysis using BioRad QX200™ System	AF 12.5% ±20% (10.0-15.0%)

RESULTS OF ANALYSIS	Result	PASS/FAIL									
Fragmentation	180 bp	PASS									
Quantity	29.5 ng/μl (total DNA) 19.4 ng/μl (dsDNA)	PASS									
Allele Frequency	<table border="1"> <thead> <tr> <th>Mutation</th><th>AF in %</th><th>PASS/FAIL</th></tr> </thead> <tbody> <tr> <td>PIK3CA E545K</td><td>12.2</td><td>PASS</td></tr> <tr> <td>PIK3CA H1047Y</td><td>11.0</td><td>PASS</td></tr> </tbody> </table>	Mutation	AF in %	PASS/FAIL	PIK3CA E545K	12.2	PASS	PIK3CA H1047Y	11.0	PASS	
Mutation	AF in %	PASS/FAIL									
PIK3CA E545K	12.2	PASS									
PIK3CA H1047Y	11.0	PASS									

COMMENTS/REMARKS	Additional information:									
	Copy numbers (CN) of the respective measurements									
	<table><tr><th>Mutation</th><th>CN wt²³/μl</th><th>CN mut²⁴/μl</th></tr><tr><td>PIK3CA E545K</td><td>2938</td><td>409</td></tr><tr><td>PIK3CA H1047Y</td><td>4187</td><td>520</td></tr></table>	Mutation	CN wt ²³ /μl	CN mut ²⁴ /μl	PIK3CA E545K	2938	409	PIK3CA H1047Y	4187	520
	Mutation	CN wt ²³ /μl	CN mut ²⁴ /μl							
	PIK3CA E545K	2938	409							
PIK3CA H1047Y	4187	520								
<i>Table 4 indicates the values of the QC assays performed by SensID GmbH with a DNA input of ~40 ng. The value for the respective mutation results from the mean value of five measured replicates (CN values are rounded). CN concentration values per microliter (μl), are based on droplet digital (ddPCR) assay counts dilution factors, and droplet volume measurements. The detection of the amount of CNs may vary depending on the assay used. Therefore, due to assay properties, there may be deviations in the observed number of copies and allele frequencies compared to the values given here.</i>										

²⁰ Protocol NK603 – Community Reference Laboratory for GM Food and Feed

²¹ Measured before filling in product tube

²² not applicable

²³ Wild Type

²⁴ Mutation

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PRODUCT INFORMATION AND QUALITY CONTROL

NAME OF PRODUCT	PIK3CA-T1 mutations 12.5% AF cfDNA- PIK3CA Q546E
DESCRIPTION	PIK3CA-T1 mutations 12.5% AF cfDNA- PIK3CA Q546E is highly characterized human DNA from cell lines.
CATALOG NUMBER	SID-000099
BATCH NUMBER	00047
MANUFACTURING CONDITIONS	<ul style="list-style-type: none"> • Manufactured and sealed in class 2 safety cabinet • At room temperature
PACKAGE SIZE	<ul style="list-style-type: none"> • 2D barcoded tube with screw cap
PACKAGE TYPE	<ul style="list-style-type: none"> • Material: Polypropylen (PP)
DATE OF MANUFACTURE	13.03.2020
EXPIRY DATE	12.03.2022
CONCENTRATION	20 ng/μl (dsDNA)
QUANTITY	400 ng (dsDNA)
NOMINAL VOLUME	25 μl; (523 ng)
MUTATION	PIK3CA p.Q546E (COSM6147*, COSV55882350* substitution, c.1636C>G, Exon 9) * GRCh38 COSMIC v90
ALLELE FREQUENCY	12.5%
QUALITY	<p>DNA quantity metrologically traceable to internationally certified reference material²⁵</p> <p>The copy number values are metrologically traceable to the natural units count 1 and ratio 1 and International System of Units (SI) derived units of volume.</p>
STORAGE CONDITIONS	+2 - +8 °C
MANUFACTURING AND QUALITY CONTROL SITES	<p>SensID GmbH</p> <p>Schillingallee 68, 18057 Rostock, Germany</p>

²⁵ ERM_AD442K

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CEO: Björn Nowack

TEST METHOD AND ACCEPTANCE CRITERIA	Quality Control	Test Method	Acceptance criteria
	Fragmentation	Fragment Length Analysis Agilent High Sensitivity DNA Kit (Agilent Technologies)	peak size 167 bp ± 10% (151 bp – 181 bp)
	Quantification	Total DNA measurement: Spectrophotometry ssDNA [ng/μl] = (A260-A320)*38 ^{26,27} dsDNA measurement: Qubit dsDNA BR Assay Kit (Invitrogen)	Total DNA: n.a. ²⁸ dsDNA: 17.5 – 22.5 ng/μl
	Allele Frequency	ddPCR Analysis using BioRad QX200™ System	AF 12.5% ±20% (10.0-15.0%)

RESULTS OF ANALYSIS	Result	PASS/FAIL						
Fragmentation	180 bp	PASS						
Quantity	29.9 ng/μl (total DNA) 20.9 ng/μl (dsDNA)	PASS						
Allele Frequency	<table border="1"> <tr> <th>Mutation</th><th>AF in %</th><th>PASS/FAIL</th></tr> <tr> <td>PIK3CA Q546E</td><td>12.8</td><td>PASS</td></tr> </table>	Mutation	AF in %	PASS/FAIL	PIK3CA Q546E	12.8	PASS	
Mutation	AF in %	PASS/FAIL						
PIK3CA Q546E	12.8	PASS						

COMMENTS/REMARKS	Additional information:						
	Copy numbers (CN) of the respective measurements						
	<table><tr><th>Mutation</th><th>CN wt²⁹/μl</th><th>CN mut³⁰/μl</th></tr><tr><td>PIK3CA Q546E</td><td>5585</td><td>821</td></tr></table>	Mutation	CN wt ²⁹ /μl	CN mut ³⁰ /μl	PIK3CA Q546E	5585	821
	Mutation	CN wt ²⁹ /μl	CN mut ³⁰ /μl				
PIK3CA Q546E	5585	821					
<p><i>Table 5 indicates the values of the QC assays performed by SensID GmbH with a DNA input of ~40 ng. The value for the respective mutation results from the mean value of five measured replicates (CN values are rounded). CN concentration values per microliter (μl), are based on droplet digital (ddPCR) assay counts dilution factors, and droplet volume measurements. The detection of the amount of CNs may vary depending on the assay used. Therefore, due to assay properties, there may be deviations in the observed number of copies and allele frequencies compared to the values given here.</i></p>							

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²⁷ Measured before filling in product tube

²⁸ not applicable

²⁹ Wild Type

³⁰ Mutation

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PRODUCT INFORMATION AND QUALITY CONTROL

NAME OF PRODUCT	PIK3CA-T1 mutations 12.5% AF cfDNA- PIK3CA E545A & C420R
DESCRIPTION	PIK3CA-T1 mutations 12.5% AF cfDNA- PIK3CA E545A & C420R is highly characterized human DNA from cell lines.
CATALOG NUMBER	SID-000099
BATCH NUMBER	00049
MANUFACTURING CONDITIONS	<ul style="list-style-type: none"> • Manufactured and sealed in class 2 safety cabinet • At room temperature
PACKAGE SIZE	<ul style="list-style-type: none"> • 2D barcoded tube with screw cap
PACKAGE TYPE	<ul style="list-style-type: none"> • Material: Polypropylen (PP)
DATE OF MANUFACTURE	23.03.2020
EXPIRY DATE	22.03.2022
CONCENTRATION	20 ng/μl (dsDNA)
QUANTITY	400 ng (dsDNA)
NOMINAL VOLUME	25 μl; (540 ng)
MUTATION	PIK3CA p.E545A (COSM12458*, COSV55873209*, substitution, c.1634A>C, Exon 9) PIK3CA p.C420R (COSM757*, COSV55874020* substitution, c.1258T>C, Exon 7) <small>* GRCh38 COSMIC v90</small>
ALLELE FREQUENCY	12.5%
QUALITY	DNA quantity metrologically traceable to internationally certified reference material ³¹ The copy number values are metrologically traceable to the natural units count 1 and ratio 1 and International System of Units (SI) derived units of volume.
STORAGE CONDITIONS	+2 - +8 °C
MANUFACTURING AND QUALITY CONTROL SITES	SensID GmbH Schillingallee 68, 18057 Rostock, Germany

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TEST METHOD AND ACCEPTANCE CRITERIA	Quality Control	Test Method	Acceptance criteria
	Fragmentation	Fragment Length Analysis Agilent High Sensitivity DNA Kit (Agilent Technologies)	peak size 167 bp ± 10% (151 bp – 181 bp)
	Quantification	Total DNA measurement: Spectrophotometry ssDNA [ng/μl] = (A260 – A320)*38 ^{32,33} dsDNA measurement: Qubit dsDNA BR Assay Kit (Invitrogen)	Total DNA: n.a. ³⁴ dsDNA: 17.5 – 22.5 ng/μl
	Allele Frequency	ddPCR Analysis using BioRad QX200™ System	AF 12.5% ±20% (10.0–15.0%)

RESULTS OF ANALYSIS		Result	PASS/FAIL							
	Fragmentation	178 bp	PASS							
	Quantity	33.6 ng/μl (total DNA) 21.6 ng/μl (dsDNA)	PASS							
	Allele									
	Frequency	<table><tr><th>Mutation</th><th>AF in %</th><th>PASS/FAIL</th></tr><tr><td>PIK3CA E545A³⁵</td><td>11.9</td><td>PASS</td></tr><tr><td>PIK3CA C420R</td><td>14.1</td><td>PASS</td></tr></table>	Mutation	AF in %	PASS/FAIL	PIK3CA E545A ³⁵	11.9	PASS	PIK3CA C420R	14.1
Mutation	AF in %	PASS/FAIL								
PIK3CA E545A ³⁵	11.9	PASS								
PIK3CA C420R	14.1	PASS								

COMMENTS/REMARKS	Additional information:									
	Copy numbers (CN) of the respective measurements									
	<table><tr><th>Mutation</th><th>CN wt³⁶/μl</th><th>CN mut³⁷/μl</th></tr><tr><td>PIK3CA E545A</td><td>5711</td><td>771</td></tr><tr><td>PIK3CA C420R</td><td>2316</td><td>381</td></tr></table>	Mutation	CN wt ³⁶ /μl	CN mut ³⁷ /μl	PIK3CA E545A	5711	771	PIK3CA C420R	2316	381
	Mutation	CN wt ³⁶ /μl	CN mut ³⁷ /μl							
	PIK3CA E545A	5711	771							
PIK3CA C420R	2316	381								
<i>Table 6 indicates the values of the QC assays performed by SensID GmbH with a DNA input of ~40 ng. The value for the respective mutation results from the mean value of five measured replicates (CN values are rounded). CN concentration values per microliter (μl), are based on droplet digital (ddPCR) assay counts dilution factors, and droplet volume measurements. The detection of the amount of CNs may vary depending on the assay used. Therefore, due to assay properties, there may be deviations in the observed number of copies and allele frequencies compared to the values given here.</i>										

³² Protocol NK603 – Community Reference Laboratory for GM Food and Feed

³³ Measured before filling in product tube

³⁴ not applicable

³⁵ A BLAST sequence analysis shows 98% homology of PIK3CA E545A mutation sequence to genome locus Homo sapiens chromosome 22, GRCh38.p13. Therefore, a higher false positive rate is expected and measured, most likely due to a cross reaction of gene probe to genome locus Homo sapiens chromosome 22, GRCh38.p13. Cross reaction AF / false positive rate is subtracted from AF for PIK3CA E545A mutation given in this CoA.

³⁶ Wild Type

³⁷ Mutation

Name and position/title of Person authorising the batch release:

Mr. Björn Nowack, Managing Director

Date of batch release: 24.03.2020

Signature batch release: Björn Nowack

This document was created electronically and is valid without a signature.