

## NGS Translocatie onderzoek: Archer FusionPlex CTL

Gene	Accession	Exon	AssayType	Description
<b>AKT1</b>	NM_005163	3	Mutation	E17K
<b>ALK</b>	NM_004304	2,4,6,10,16,17, 18,19,(intron19), 20,21,22,23	Fusion/ Expression Imbalance	5'
<b>ALK</b>	NM_004304	N/A	Expression	N/A
<b>ALK</b>	NM_004304	21,22,23,25	Imbalance	T1151ins,L1152R, C1156Y,F1174L, L1196M,G1202R, S1206Y,G1269A
<b>AXL</b>	NM_001699	18,19,20	Expression	3'
<b>AXL</b>	NM_001699	N/A	Mutation	N/A
<b>BRAF</b>	NM_004333	11,15	Mutation	G466V,G469, Y472,L597V, V600,D594G
<b>BRAF</b>	NM_004333	7,8,9,10,11	Fusion	5'
<b>BRAF</b>	NM_004333	7,8,10	Fusion	3'
<b>BRAF</b>	NM_004333	N/A	Expression	N/A
<b>CALCA</b>	NM_001741	N/A	Expression	N/A
<b>CCND1</b>	NM_053056	1,2,3,4	Fusion	5'
<b>CCND1</b>	NM_053056	1,2,4	Fusion	3'
<b>CCND1</b>	NM_053056	N/A	Expression	N/A
<b>CTNNB1</b>	NM_001904	3	Mutation	D32G,S37,G34
<b>DDR2</b>	NM_006182	17	Mutation	S768R,T765P, G774
<b>EGFR</b>	NM_005228	8	Splice Variant	5' (2-7exon skipping event) G719, A763insFQEA, T790M,L858R, L861Q,Y764,
<b>EGFR</b>	NM_005228	18,19,20,21	Mutation	V774,L777,L768, P753,L760, E709A,L747,varius deletions in exon 19
<b>EGFR</b>	NM_005228	N/A	Expression	N/A
<b>ERBB2</b>	NM_004448	23	Mutation	C775ins
<b>FGFR1</b>	NM_015850	13	Mutation	V561M
<b>FGFR1</b>	NM_015850	2,8,9,10,17	Fusion	5'
<b>FGFR1</b>	NM_015850	17	Fusion	3'
<b>FGFR1</b>	NM_015850	N/A	Expression	N/A
<b>FGFR2</b>	NM_000141	2,5,7,8,9,10	Fusion	5'

<b>FGFR2</b>	NM_000141	17	Fusion	3'
<b>FGFR2</b>	NM_000141	N/A	Expression	N/A
<b>FGFR3</b>	NM_000142	17,(intron17)	Fusion	3'
<b>FGFR3</b>	NM_000142	3,5,8,9,10	Fusion	5'
<b>FGFR3</b>	NM_000142	N/A	Expression	N/A
<b>GNAS</b>	NM_000516	8,9	Mutation	Various exon 8 and 9 mutations
<b>GNAS</b>	NM_000516	8,9	Mutation	N/A
<b>HRAS</b>	NM_005343	2,3	Mutation	G12,G13,Q61
<b>IDH1</b>	NM_005896	4	Mutation	R132
<b>IDH2</b>	NM_002168	4	Mutation	R172,R140
<b>KRAS</b>	NM_004985	2,3,4	Mutation	G12,G13, Q61,A146
<b>KRT20</b>	NM_019010	N/A	Expression	N/A
<b>KRT7</b>	NM_005556	N/A	Expression	N/A
<b>MAP2K1</b>	NM_002755	2,3	Mutation	Q56P,K57N,D67N
<b>MET</b>	NM_000245	2	Fusion	3'
<b>MET</b>	NM_000245	2,4,5,6,13,14, 16,17,21	Fusion	5'
<b>MET</b>	NM_000245	15	Splice	5'
<b>MET</b>	NM_000245	N/A	Expression Variant	N/A
<b>NRAS</b>	NM_002524	2,3	Mutation	G12,G13,Q61
<b>NRG1</b>	NM_004495	1,2,3,6	Fusion	5'
<b>NTRK1</b>	NM_002529	2,4,6,8,10, 11,12,13	Fusion/ Expression Imbalance	5'
<b>NTRK1</b>	NM_002529	N/A	Expression	N/A
<b>NTRK2</b>	NM_006180	5,7,9,11,12,13,14, 15,16,17	Imbalance	5'
<b>NTRK2</b>	NM_006180	N/A	Expression	N/A
<b>NTRK3</b>	NM_002530	4,7,10,13, 14,15,16	Fusion/ Expression Imbalance	5'
<b>NTRK3</b>	NM_001007156	15	Expression	5'
<b>NTRK3</b>	NM_002530	N/A	Imbalance	N/A
<b>PIK3CA</b>	NM_006218	10,21	Expression	E542K,E545, H1047
<b>PPARG</b>	NM_015869	1,2,3,5	Fusion/ Expression Imbalance	5'
<b>PTH</b>	NM_000315	N/A	Expression	N/A
<b>RAF1</b>	NM_002880	4,5,6,7,9, 10,11,12	Imbalance	5'
<b>RET</b>	NM_020630	11,13,14,15,16	Fusion	C634,M918T, V804,Y806D, E768D,A883F

<b>RET</b>	NM_020630	2,4,6,8,9,10, 11,12,13,14	Fusion/ Expression Imbalance	5'
<b>RET</b>	NM_020630	N/A	Expression	N/A
<b>ROS1</b>	NM_002944	2,4,7,31,32, 33,34,35,36,37	Imbalance	5'
<b>ROS1</b>	NM_002944	38	Expression	G2032R
<b>ROS1</b>	NM_002944	N/A	Fusion/ Expression Imbalance	N/A
<b>SLC5A5</b>	NM_000453	N/A	Expression	N/A
<b>THADA</b>	NM_022065	24,25,26,27,28, 29,30,36,37	Imbalance	3'
<b>THADA</b>	NM_022065	N/A	Mutation	N/A
<b>TTF1</b>	NM_007344	N/A	Expression	N/A
<b>CHMP2A</b>	NM_014453	3,4	Expression	N/A
<b>GPI</b>	NM_000175	15,16	Control	N/A
<b>RAB7A</b>	NM_004637	3,4	Control	N/A
<b>VCP</b>	NM_007126	14,15	Control	N/A

\*not all accession numbers covered are shown in table