

Ultimate ORF ID	Neg Signal.Use d	Signal.Use d	Ratio to Neg	Assay	Z-Factor	Z-Score	CI P-Value	Replicate Spot CV	Description
IOH14808	30	65492	2183	5 µg/ml	1.00	14.95	1.1E-05	0%	GABA(A) receptor-associated protein like 1 (GABARAPL1)
IOH4300	37	65492	1770	5 µg/ml	0.96	14.95	1.1E-05	0%	actin binding LIM protein 1 (ABLM1)
IOH43439	38	65492	1746	5 µg/ml	1.00	14.95	1.1E-05	0%	chromosome 11 open reading frame 63 (C11orf63), transcript variant 2
IOH21794	39	65492	1701	5 µg/ml	1.00	14.95	1.1E-05	0%	tumor protein D52-like 3 (TPD52L3)
IOH56887	72	65492	910	5 µg/ml	1.00	14.95	1.1E-05	0%	Splicing factor 1
IOH5807	26	65490	2568	5 µg/ml	1.00	14.95	1.1E-05	0%	PDZ domain containing 1 (PDZK1)
IOH5166	27	65490	2426	5 µg/ml	1.00	14.95	1.1E-05	0%	chromosome 16 open reading frame 80 (C16orf80)
IOH4025	23	65489	2847	5 µg/ml	1.00	14.95	1.1E-05	0%	solute carrier family 9 (sodium/hydrogen exchanger), member 3 regulator 1 (SLC9A3R1)
IOH61586	27	65489	2426	5 µg/ml	0.99	14.95	1.1E-05	0%	Myelin basic protein
IOH21704	29	65489	2258	5 µg/ml	1.00	14.95	1.1E-05	0%	chromosome 8 open reading frame 59 (C8orf59)
IOH7230	32	65488	2047	5 µg/ml	1.00	14.95	1.1E-05	0%	zinc finger, MYM-type 5 (ZMYM5)
IOH4088	53	65487	1247	5 µg/ml	0.99	14.95	1.1E-05	0%	proline/serine-rich coiled-coil 1 (PSRC1), transcript variant 1
IOH3994	45	65487	1472	5 µg/ml	0.99	14.95	1.1E-05	0%	small trans-membrane and glycosylated protein (LOC57228), transcript variant 2
IOH10928	34	65486	1926	5 µg/ml	1.00	14.95	1.1E-05	0%	melanoma antigen family B, 1 (MAGEB1), transcript variant 1
IOH12150	57	65483	1159	5 µg/ml	0.96	14.95	1.1E-05	0%	leukocyte receptor cluster (LRC) member 1 (LENG1)
IOH40668	29	65478	2258	5 µg/ml	1.00	14.95	1.1E-05	0%	ubiquitin protein ligase E3 component n-recognin 2 (UBR2)
IOH3488	39	65443	1700	5 µg/ml	1.00	14.94	1.1E-05	0%	LSM4 homolog, U6 small nuclear RNA associated (S. cerevisiae) (LSM4)
	5	41898	8380	5 µg/ml	0.81	9.47	2.8E-05	6%	NRP1 / Neuropilin-1 Protein
IOH28970	27	65483	2471	50 µg/ml	0.98	7.60	4.7E-05	0%	chromosome 22 open reading frame 9 (C22orf9)
	29	65480	2298	50 µg/ml	1.00	7.60	4.7E-05	0%	NIMA (never in mitosis gene a)-related kinase 1 (NEK1)
IOH56964	16	65479	4092	50 µg/ml	1.00	7.60	4.7E-05	0%	Apoptosis-inducing factor 2

IOH4887	284	65479	231	50 µg/ml	0.99	7.60	4.7E-05	0%	APEX nuclease (apurinic/aprimidinic endonuclease) 2 (APEX2), nuclear gene encoding mitochondrial protein
IOH27737	42	65478	1559	50 µg/ml	0.99	7.60	4.7E-05	0%	sciellin (SCEL)
IOH21824	38	65478	1746	50 µg/ml	0.99	7.60	4.7E-05	0%	dermatopontin (DPT)
IOH40615	44	65478	1505	50 µg/ml	0.99	7.60	4.7E-05	0%	serine/threonine kinase 40 (STK40)
IOH4329	21	65477	3118	50 µg/ml	1.00	7.60	4.7E-05	0%	peptidylprolyl isomerase A (cyclophilin A) (PPIA)
IOH42260	38	65477	1746	50 µg/ml	0.99	7.60	4.7E-05	0%	F-box protein 21 (FBXO21), transcript variant 2
IOH6102	45	65477	1471	50 µg/ml	0.99	7.60	4.7E-05	0%	eukaryotic translation initiation factor 4E family member 2 (EIF4E2)
IOH41337	5	65476	14550	50 µg/ml	0.99	7.60	4.7E-05	0%	syndecan binding protein (syntenin) (SDCBP), transcript variant 2
IOH63097	31	65476	2147	50 µg/ml	0.99	7.60	4.7E-05	0%	Myelin basic protein (MBP), transcript variant 4, mRNA
IOH57049	17	65475	3851	50 µg/ml	1.00	7.60	4.7E-05	0%	Cytochrome b-c1 complex subunit 7
IOH44469	26	65475	2518	50 µg/ml	0.99	7.60	4.7E-05	0%	PREDICTED: Homo sapiens hypothetical LOC401363 (LOC401363)
IOH10469	33	65475	2015	50 µg/ml	0.99	7.60	4.7E-05	0%	programmed cell death 6 (PDCD6)
IOH7249	35	65475	1871	50 µg/ml	0.99	7.60	4.7E-05	0%	myotilin (MYOT)
IOH4744	56	65475	1169	50 µg/ml	0.99	7.60	4.7E-05	0%	ubiquitin fusion degradation 1 like (yeast) (UFD1L), transcript variant 1, mRNA.
IOH42993	23	65475	2847	50 µg/ml	1.00	7.60	4.7E-05	0%	PREDICTED: Homo sapiens hypothetical protein FLJ36032 (FLJ36032)
IOH22975	28	65475	2381	50 µg/ml	0.99	7.60	4.7E-05	0%	Protein FAM54A
	3	65474	21825	50 µg/ml	0.99	7.60	4.7E-05	0%	Mitogen-activated protein kinase 1; see catalog number for detailed information on wild-type or point mutant status
IOH21984	66	65473	1000	50 µg/ml	0.98	7.60	4.7E-05	0%	baculoviral IAP repeat-containing 4 (BIRC4)
IOH28010	35	65473	1898	50 µg/ml	1.00	7.60	4.7E-05	0%	mitogen-activated protein kinase-activated protein kinase 5 (MAPKAPK5), transcript variant 1
IOH61973	26	65473	2518	50 µg/ml	0.98	7.60	4.7E-05	0%	Peptidyl-prolyl cis-trans isomerase NIMA-interacting 4
IOH14566	32	65472	2046	50 µg/ml	0.99	7.60	4.7E-05	0%	DnaJ (Hsp40) homolog, subfamily C, member 7 (DNAJC7)

IOH4688	43	65472	1541	50 µg/ml	0.99	7.60	4.7E-05	0%	death inducer-obliterator 1 (DIDO1)
IOH62268	27	65471	2471	50 µg/ml	0.99	7.60	4.7E-05	0%	Homeobox protein Meis1
IOH43381	33	65470	2014	50 µg/ml	1.00	7.60	4.7E-05	0%	PREDICTED: Homo sapiens hypothetical LOC150371 (LOC150371)
IOH40618	33	65470	1984	50 µg/ml	0.99	7.60	4.7E-05	0%	hypothetical protein FLJ22795 (FLJ22795)
IOH23003	107	65470	615	50 µg/ml	0.99	7.60	4.7E-05	0%	mitochondrial ribosomal protein L19 (MRPL19), nuclear gene encoding mitochondrial protein
	27	65470	2425	50 µg/ml	1.00	7.60	4.7E-05	0%	aurora kinase A (AURKA), transcript variant 2
IOH40642	30	65470	2219	50 µg/ml	0.98	7.60	4.7E-05	0%	LATS, large tumor suppressor, homolog 1 (Drosophila) (LATS1)
IOH27153	44	65469	1505	50 µg/ml	0.97	7.60	4.7E-05	0%	melanoma antigen family B, 4 (MAGEB4)
IOH42688	50	65469	1323	50 µg/ml	1.00	7.60	4.7E-05	0%	PREDICTED: Homo sapiens hypothetical protein LOC283663 (LOC283663), mRNA
IOH46139	27	65468	2470	50 µg/ml	0.99	7.60	4.7E-05	0%	olfactomedin 1 (OLFM1), transcript variant 1
IOH46132	32	65468	2046	50 µg/ml	0.99	7.60	4.7E-05	0%	protein tyrosine phosphatase, non-receptor type 7 (PTPN7), transcript variant 2
IOH42441	43	65468	1523	50 µg/ml	0.99	7.60	4.7E-05	0%	PREDICTED: Homo sapiens hypothetical LOC401394 (LOC401394)
IOH5992	58	65468	1139	50 µg/ml	0.99	7.60	4.7E-05	0%	chromosome 6 open reading frame 134 (C6orf134)
IOH27781	29	65468	2258	50 µg/ml	0.99	7.60	4.7E-05	0%	chromosome X open reading frame 56 (CXorf56)
IOH21165	32	65467	2046	50 µg/ml	0.99	7.60	4.7E-05	0%	Serine/threonine-protein kinase 6
IOH59050	17	65465	3968	50 µg/ml	0.98	7.60	4.7E-05	0%	Kallikrein 8 (neuropsin/ovasin) (KLK8), transcript variant 1, mRNA
IOH41289	34	65465	1954	50 µg/ml	0.99	7.60	4.7E-05	0%	GABA(A) receptor-associated protein (GABARAP)
IOH42885	44	65465	1488	50 µg/ml	0.99	7.60	4.7E-05	0%	hypothetical protein LOC150577 (LOC150577)
IOH27934	36	65464	1844	50 µg/ml	0.99	7.60	4.7E-05	0%	Delta(3,5)-Delta(2,4)-dienoyl-CoA isomerase, mitochondrial
IOH45655	35	65463	1870	50 µg/ml	0.99	7.60	4.7E-05	0%	Probable ATP-dependent RNA helicase DDX6
IOH9802	26	65462	2518	50 µg/ml	0.99	7.60	4.7E-05	0%	outer dense fiber of sperm tails 2 (ODF2)
IOH5926	43	65462	1522	50 µg/ml	0.99	7.60	4.7E-05	0%	chromosome 19 open reading frame 43 (C19orf43)
IOH40597	61	65461	1073	50 µg/ml	0.99	7.60	4.7E-05	0%	Protein midA homolog, mitochondrial
IOH60410	26	65458	2567	50 µg/ml	0.99	7.60	4.7E-05	0%	Nephrocystin-1
IOH10203	37	65453	1769	50 µg/ml	0.99	7.60	4.7E-05	0%	Protein FAM107A
IOH12969	43	65453	1540	50 µg/ml	0.99	7.60	4.7E-05	0%	sciellin (SCEL)

IOH13203	40	65452	1657	50 µg/ml	0.99	7.60	4.7E-05	0%	SH2 domain protein 2A (SH2D2A)
IOH7023	50	65452	1309	50 µg/ml	0.99	7.60	4.7E-05	0%	Uncharacterized protein C16orf14
IOH44018	41	65451	1616	50 µg/ml	0.99	7.60	4.7E-05	0%	PREDICTED: Homo sapiens hypothetical protein LOC339685 (LOC339685)
IOH4903	27	65450	2470	50 µg/ml	0.99	7.60	4.7E-05	0%	mitochondrial ribosomal protein L11 (MRPL11), nuclear gene encoding mitochondrial protein, transcript variant 1
IOH21753	112	65450	587	50 µg/ml	0.99	7.60	4.7E-05	0%	Ral GEF with PH domain and SH3 binding motif 1 (RALGPS1)
IOH40097	38	65448	1722	50 µg/ml	0.99	7.60	4.7E-05	0%	abhydrolase domain containing 11 (ABHD11)
IOH22742	100	65447	658	50 µg/ml	0.99	7.60	4.7E-05	0%	Putative E3 ubiquitin-protein ligase SH3RF2
IOH45485	29	65446	2257	50 µg/ml	0.99	7.60	4.7E-05	0%	mitochondrial ribosomal protein L55 (MRPL55), nuclear gene encoding mitochondrial protein, transcript variant 4
IOH42078	34	65444	1954	50 µg/ml	0.95	7.60	4.7E-05	0%	chromosome 22 open reading frame 9 (C22orf9), transcript variant 2
IOH40656	38	65444	1745	50 µg/ml	0.99	7.60	4.7E-05	0%	mitochondrial GTPase 1 homolog ( <i>S. cerevisiae</i> ) (MTG1)
IOH21756	33	65443	2014	50 µg/ml	0.99	7.60	4.7E-05	0%	DNA repair and recombination protein RAD54B
IOH13449	38	65439	1745	50 µg/ml	0.99	7.60	4.7E-05	0%	PRKR interacting protein 1 (IL11 inducible) (PRKRIP1)
IOH50108	34	65438	1953	50 µg/ml	0.99	7.60	4.7E-05	0%	G antigen, family D, 2 (GAGED2), transcript variant 3, mRNA
IOH25922	40	65437	1636	50 µg/ml	0.99	7.60	4.7E-05	0%	TBC1 domain family, member 10C (TBC1D10C)
IOH41478	31	65434	2145	50 µg/ml	0.98	7.60	4.7E-05	0%	quaking homolog, KH domain RNA binding (mouse) (QKI), transcript variant 4
IOH26570	30	65427	2181	50 µg/ml	0.97	7.59	4.7E-05	0%	leucine rich repeat containing 6 (LRRC6)
IOH61531	32	65423	2077	50 µg/ml	0.98	7.59	4.7E-05	0%	hypothetical protein LOC51233, mRNA (cDNA clone MGC:75009 IMAGE:5170001), complete cds.
IOH41283	74	65417	884	50 µg/ml	0.99	7.59	4.7E-05	0%	2',5'-oligoadenylate synthetase 1, 40/46kDa (OAS1), transcript variant 1
IOH12221	51	65416	1283	50 µg/ml	0.99	7.59	4.7E-05	0%	Protein mab-21-like 2
IOH21099	77	65410	849	50 µg/ml	0.97	7.59	4.7E-05	0%	dual specificity phosphatase 22 (DUSP22)
IOH14530	300	65410	218	50 µg/ml	0.99	7.59	4.7E-05	0%	tetra-peptide repeat homeobox-like (TPRXL)

IOH39895	40	65408	1656	50 µg/ml	0.99	7.59	4.7E-05	0%	chondrosarcoma associated gene 1 (CSAG1)
IOH10871	63	65401	1046	50 µg/ml	0.97	7.59	4.7E-05	0%	chromosome 10 open reading frame 63 (C10orf63)
IOH25846	35	65388	1868	50 µg/ml	0.99	7.59	4.7E-05	0%	NDUFA12-like (NDUFA12L)
IOH11297	32	65367	2075	50 µg/ml	0.99	7.59	4.7E-05	0%	deoxynucleotidyltransferase, terminal (DNTT), transcript variant 1
IOH6250	44	65365	1503	50 µg/ml	0.99	7.59	4.7E-05	0%	chromosome 16 open reading frame 48 (C16orf48)
IOH26289	34	65361	1951	50 µg/ml	0.99	7.59	4.7E-05	0%	CWC15 homolog ( <i>S. cerevisiae</i> ) (HSPC148)
IOH21738	53	65360	1233	50 µg/ml	0.99	7.59	4.8E-05	0%	pescadillo homolog 1, containing BRCT domain (zebrafish) (PES1)
IOH14526	32	65340	2074	50 µg/ml	0.98	7.58	4.8E-05	0%	Rho guanine nucleotide exchange factor (GEF) 5 (ARHGEF5)
IOH5640	33	65319	2010	50 µg/ml	0.98	7.58	4.8E-05	0%	tRNA phosphotransferase 1 (TRPT1), transcript variant 2
IOH45842	31	65270	2105	50 µg/ml	0.99	7.58	4.8E-05	0%	peptidylprolyl isomerase E (cyclophilin E) (PPIE), transcript variant 3
IOH12275	76	64668	851	50 µg/ml	0.94	7.50	5.0E-05	2%	coiled-coil domain containing 131 (CCDC131)
IOH12012	84	64313	770	50 µg/ml	0.92	7.46	5.1E-05	3%	UPF0498 protein KIAA1191
IOH55892	31	63798	2092	50 µg/ml	0.80	7.39	5.3E-05	4%	Transcription factor 25
IOH27186	37	63521	1717	50 µg/ml	0.86	7.36	5.3E-05	4%	egl nine homolog 2 ( <i>C. elegans</i> ) (EGLN2), transcript variant 1
IOH6405	23	63382	2756	50 µg/ml	0.97	7.34	5.1E-05	1%	Tripartite motif-containing protein 55
IOH12990	54	62665	1160	50 µg/ml	0.80	7.26	5.7E-05	6%	Adrenodoxin-like protein, mitochondrial
IOH46168	42	62601	1508	50 µg/ml	0.98	7.25	5.2E-05	0%	cell cycle associated protein 1 (CAPRIN1), transcript variant 1
IOH61486	30	62588	2122	50 µg/ml	0.79	7.25	5.7E-05	7%	splicing factor, arginine/serine-rich 2, mRNA (cDNA clone MGC:87440 IMAGE:4837392), complete cds
	24	62142	2644	50 µg/ml	0.89	7.19	5.5E-05	3%	casein kinase 1, alpha 1 (CSNK1A1), transcript variant 1
IOH28084	25	61306	2452	50 µg/ml	0.98	7.09	5.4E-05	0%	T-cell leukemia/lymphoma 1B (TCL1B)
IOH40016	12	60458	5257	50 µg/ml	0.85	6.99	5.9E-05	4%	Josephin domain containing 2 (JOSD2)
IOH5138	156	59775	384	50 µg/ml	0.86	6.90	6.1E-05	4%	protein regulator of cytokinesis 1 (PRC1), transcript variant 1

IOH27824	27	59588	2207	50 µg/ml	0.57	6.88	7.0E-05	14%	WW domain binding protein 5 (WBP5), transcript variant 1
IOH12531	38	59584	1589	50 µg/ml	0.87	6.88	6.1E-05	4%	iron-sulfur cluster scaffold homolog (E. coli) (ISCU)
IOH3444	30	59216	2007	50 µg/ml	0.74	6.83	6.6E-05	9%	chromogranin B (secretogranin 1) (CHGB)
IOH5568	32	58930	1842	50 µg/ml	0.95	6.80	6.0E-05	1%	alkB, alkylation repair homolog 7 (E. coli) (ALKBH7)
IOH45459	26	58358	2289	50 µg/ml	0.82	6.73	6.5E-05	6%	WAS protein family homolog 1
IOH40048	21	56802	2771	50 µg/ml	0.96	6.54	6.4E-05	1%	proopiomelanocortin (adrenocorticotropin/ beta-lipotropin/ alpha-melanocyte stimulating hormone/ beta-melanocyte stimulating hormone/ beta-endorphin) (POMC), transcript variant 2
IOH54659	15	56191	3746	50 µg/ml	0.85	6.46	6.9E-05	5%	Low molecular weight phosphotyrosine protein phosphatase
IOH41465	44	55008	1250	50 µg/ml	0.69	6.32	7.8E-05	10%	small nuclear ribonucleoprotein polypeptide C (SNRPC)
IOH62570	27	52758	1954	50 µg/ml	0.85	6.04	7.8E-05	5%	zinc finger, MYND domain containing 11, mRNA (cDNA clone MGC:111056 IMAGE:6186814), complete cds
IOH62588	26	51427	2017	50 µg/ml	0.64	5.88	9.0E-05	11%	Outer dense fiber protein 2
IOH42361	67	51217	764	50 µg/ml	0.55	5.85	9.7E-05	15%	complexin 2 (CPLX2), transcript variant 2
IOH11555	41	50977	1259	50 µg/ml	0.99	5.82	7.8E-05	0%	chromosome 1 open reading frame 87 (C1orf87)
IOH39976	34	50752	1515	50 µg/ml	0.79	5.79	8.5E-05	5%	leucine aminopeptidase 3 (LAP3)
IOH40890	26	50593	1984	50 µg/ml	0.62	5.78	9.5E-05	12%	outer dense fiber of sperm tails 2 (ODF2), transcript variant 1
IOH38079	64	49830	779	50 µg/ml	0.70	5.68	8.9E-05	6%	FLJ36874 protein (FLJ36874)
IOH43320	35	49806	1423	50 µg/ml	0.98	5.68	8.2E-05	0%	Smoothelin-like protein 2
IOH27342	33	49792	1509	50 µg/ml	0.79	5.68	9.0E-05	7%	retinaldehyde binding protein 1-like 1 (RLBP1L1)
IOH6528	40	49260	1247	50 µg/ml	0.93	5.61	8.6E-05	2%	forkhead box P1 (FOXP1)
	25	48402	1936	50 µg/ml	0.69	5.51	1.0E-04	9%	MAP/microtubule affinity-regulating kinase 3 (MARK3)
IOH43177	30	47360	1579	50 µg/ml	0.90	5.38	9.5E-05	3%	Calcium channel, voltage-dependent, beta 1 subunit (CACNB1), transcript variant 2, mRNA

IOH40570	32	46289	1469	50 µg/ml	0.89	5.25	9.9E-05	3%	mitochondrial ribosomal protein L28 (MRPL28), nuclear gene encoding mitochondrial protein
IOH13845	30	45228	1508	50 µg/ml	0.71	5.12	1.1E-04	9%	mitochondrial ribosomal protein S6 (MRPS6), nuclear gene encoding mitochondrial protein
IOH27184	33	44561	1350	50 µg/ml	0.85	5.04	1.1E-04	5%	Fanconi anemia, complementation group M (FANCM)
IOH4999	63	44322	704	50 µg/ml	0.81	5.01	1.1E-04	6%	SERPINE1 mRNA binding protein 1 (SERBP1)
IOH5733	34	43877	1290	50 µg/ml	0.88	4.95	1.1E-04	4%	WAS/WASL-interacting protein family member 1
IOH3844	33	43512	1339	50 µg/ml	0.81	4.91	1.2E-04	6%	AKT interacting protein (AKTIP), transcript variant 2
IOH4591	29	43046	1484	50 µg/ml	0.87	4.85	1.1E-04	0%	Coronin-2A
IOH10699	54	41637	778	50 µg/ml	0.75	4.68	1.3E-04	8%	DEAH (Asp-Glu-Ala-His) box polypeptide 40 (DHX40)
IOH12064	53	41634	793	50 µg/ml	0.64	4.68	1.4E-04	10%	cirrhosis, autosomal recessive 1A (cirhin) (CIRH1A)
IOH13703	54	39974	740	50 µg/ml	0.72	4.47	1.5E-04	9%	KIAA1530 protein (KIAA1530)
IOH12380	38	39036	1027	50 µg/ml	0.84	4.36	1.4E-04	5%	homer homolog 2 (Drosophila) (HOMER2)
IOH29576	13	19883	1591	5 µg/ml	0.58	4.35	1.4E-04	14%	protein tyrosine phosphatase type IVA, member 2 (PTP4A2), transcript variant 1
IOH56483	33	37924	1167	50 µg/ml	0.64	4.22	1.7E-04	12%	Nucleoporin NUP53
IOH11180	31	19315	623	5 µg/ml	0.70	4.22	1.4E-04	9%	Zinc finger CCHC domain-containing protein 8
IOH11119	32	37745	1198	50 µg/ml	0.73	4.20	1.6E-04	8%	cDNA clone MGC:17410 IMAGE:4156035, complete cds
IOH29378	35	37230	1064	50 µg/ml	0.52	4.14	1.9E-04	15%	erythrocyte membrane protein band 4.9 (dematin) (EPB49)
IOH21599	56	37133	663	50 µg/ml	0.67	4.12	1.7E-04	10%	cDNA clone IMAGE:5271031
IOH26615	33	36900	1118	50 µg/ml	0.66	4.10	1.7E-04	10%	spermatogenesis associated, serine-rich 2 (SPATS2)
IOH20961	38	36428	959	50 µg/ml	0.56	4.04	1.8E-04	9%	p21(CDKN1A)-activated kinase 6 (PAK6)
IOH43039	70	36066	515	50 µg/ml	0.89	3.99	1.6E-04	3%	cyclin B3 (CCNB3), transcript variant 2
IOH28672	35	35988	1043	50 µg/ml	0.94	3.98	1.6E-04	2%	sulfotransferase family, cytosolic, 1C, member 4 (SULT1C4)
IOH50143	20	35481	1774	50 µg/ml	0.64	3.92	1.9E-04	12%	NCK adaptor protein 2 (NCK2), transcript variant 3, mRNA

IOH39946	24	34112	1421	50 µg/ml	0.65	3.75	2.0E-04	10%	Nuclear protein NP60
IOH12214	33	34039	1047	50 µg/ml	0.76	3.74	2.0E-04	7%	dual specificity phosphatase 13 (DUSP13), transcript variant 6
IOH13466	39	33222	863	50 µg/ml	0.80	3.64	2.0E-04	6%	chromosome 12 open reading frame 52 (C12orf52)
IOH29446	26	31525	1213	50 µg/ml	0.91	3.44	2.1E-04	2%	EMG1 nucleolar protein homolog (S. cerevisiae) (EMG1)
IOH42760	32	31470	983	50 µg/ml	0.74	3.43	2.3E-04	8%	La ribonucleoprotein domain family, member 4 (LARP4), transcript variant 3
IOH45764	31	31421	1014	50 µg/ml	0.63	3.42	2.4E-04	11%	transcription elongation factor A (SII), 2 (TCEA2), transcript variant 2
IOH5070	46	30739	676	50 µg/ml	0.59	3.34	2.6E-04	13%	polo-like kinase 1 (Drosophila) (PLK1)
IOH21130	28	30272	1101	50 µg/ml	0.71	3.28	2.6E-04	9%	dual specificity phosphatase 14 (DUSP14)
IOH40500	49	29698	612	50 µg/ml	0.90	3.21	2.4E-04	3%	Histone deacetylase 7
IOH40120	51	29655	587	50 µg/ml	0.93	3.21	2.4E-04	1%	potassium channel tetramerisation domain containing 18 (KCTD18)
IOH62017	33	28644	868	50 µg/ml	0.56	3.08	3.1E-04	14%	Sin3 histone deacetylase corepressor complex component SDS3
IOH42047	42	28611	681	50 µg/ml	0.70	3.08	2.9E-04	9%	hypothetical LOC401052 (LOC401052)