

Table S1. Sequences of oligonucleotides used in these studies

Oligonucleotide	Sequence	Position	Used in
FO 2234	CCAGGTAGCAAACCACGAAGCA	-451/-429	<i>SPT2</i> deletion
FO2235	TCGATCAACTGCAACAACCAAGTC	+1295 /+1319	<i>SPT2</i> deletion
FO2823	CGCCACGCAGCAAAGGAGTTCC TACACAATCCGGAACAAGATTGT ACTGAGAGTGCAC	-37/+1	<i>HIR2</i> deletion
FO2824	AAATATAAGAGTTTAAACTATA CATTGTAAAGCCAAACTGTGC GGTATTTACACCG	+2627/+2666	<i>HIR2</i> deletion
FO2618	GCTTGTTACTATGTGCCATCGCG	-145/-122	<i>HIR3</i> deletion
FO2619	GGTCATCAACACACCCGTACACC	+4970/+4992	<i>HIR3</i> deletion
FO2221	CATGAAGAGGAGAAGAGACGCC GTAAGAAGGGCATAACGC CGGATCCCCGGGTTAATTAA	+961/+1000	<i>SPT2</i> tag
FO2223	CATTTACGTCCATATATCAAA ACATATATCAATATTCC GAATTCGAGCTCGTTTAAAC	+1041/+1002	<i>SPT2</i> tag
FO2603	GAGGGAAAAAAGTTGTCCAAAG GAAGGGGTATAAGCTTAGAATT CGAGCTCGTTTAAAC	+2524/+2562	<i>HIR1</i> tag
FO2604	AATGGACTAAGGATCTGGGTTG TTGGTCGTTATATAAACGGATCC CCGGGTTAATTAA	+2483/+2520	<i>HIR1</i> tag
FO2599	AAAATATAAGAGTTTAAACTAT ACATTGTAAAGCCAAAGAATT CGAGCTCGTTTAAAC	+2629/+2667	<i>HIR2</i> tag
FO2600	TACTACACGTTATGCCAAGGA AATGAATATAATATCTCGGATC CCCGGTTAATTAA	+2588/+2625	<i>HIR2</i> tag
FO943	GCAGCTGTCGAAAGAAAAACGG	-1157/-1136	<i>SER3</i> ChIP

FO944	GGTAGCTTCAGGTTGCTTCACGG	-868/-846	<i>SER3</i> ChIP
FO1624	GGCGGAAAAGCGGAACTTTC	-644/-625	<i>SER3</i> ChIP
FO1625	CCCCTTGAATTCCATTTGTGAG	-329/-350	<i>SER3</i> ChIP
FO1799	GAAAATGCAACGCTGCCCGTGC	-234/-213	<i>SER3</i> ChIP
FO1797	GAGCCGCTCATATTCATA GCTTG	+65/+43	<i>SER3</i> ChIP
FO941	CGTTCCACAGCGCTTGAATGCTG	+111/+133	<i>SER3</i> ChIP
FO942	GCTGGTAGCGTAGTCTAAGTCAAC	+429/+406	<i>SER3</i> ChIP
FO1616	CAACTCGCCTTTCTCCAACCTC	+447/+467	<i>SER3</i> ChIP
FO1617	GAAACTTGTCTGGCAGTACCC	+725/+705	<i>SER3</i> ChIP
FO1618	GTGACACTACATGTACCAGC	+760/+779	<i>SER3</i> ChIP
FO1619	GCCACCAATGTGTGGTGTCAAG	+1050/+1029	<i>SER3</i> ChIP
FO1620	CGGTATTGAAGTGGCTACCGC	+1077/+1097	<i>SER3</i> ChIP
FO1621	CGCTTTGGTCAACAGAAGAG	+1342/+1323	<i>SER3</i> ChIP
FO657	TCTGCTAAGATCTCAATTAGATTG	+1378/+1401	<i>SER3</i> ChIP
FO568	GTCCTTAAACATGTGCACTG	-375/-356	<i>SER3</i> ChIP
FO1885	TTCGGGTTTTTTTCCTTCCTTTTCC	-623/-599	<i>PMA1</i> ChIP
FO1886	GTCACCGGTCATAATTGGAATAAC	-390/-413	<i>PMA1</i> ChIP
FO1881	ACGAAACGTGGTCGATGGTGGGTA	-391/-368	<i>PMA1</i> ChIP
FO1849	CTTTTGAATGTGTGTATAAAAGAGAG	-90/-115	<i>PMA1</i> ChIP
FO1810	CGACGACGAAGACAGTGATAACG	+168/+190	<i>PMA1</i> ChIP
FO1811	ATTGAATTGGACCGACGAAAAA CATAAC	+376/+349	<i>PMA1</i> ChIP
FO1812	GTTTGCCAGCTGTCGTTACCACCAC	+1010/+1034	<i>PMA1</i> ChIP/probe
FO1813	GCAGCCAAACAAGCAGTCAACA	+1235/+1209	<i>PMA1</i> ChIP

TCAAG

FO1814	CTATTATTGATGCTTTGAAGAC CTCCAG	+2018/+2045	<i>PMA1</i> ChIP
FO1815	TGCCCAAATAATAGACATACC CCATAA	+2290/+2263	<i>PMA1</i> ChIP/probe
FO1816	CGCTTATTCCTACTAGAGTTTG	+2904/2927	<i>PMA1</i> ChIP
FO1817	GTGTTGTGAATTGTGCTCTATTAG	+3469/3493	<i>PMA1</i> ChIP
FO1231	TGCCCTTTTTCTGTTAGA	-301/-283	<i>PDC1</i> ChIP
FO1453	AACGGTGTTAACGTTGACTTGC	+45/+66	<i>PDC1</i> ChIP
FO1416	AAGTCTTATGGGGTTCCATTGG	+1226/+1247	<i>PDC1</i> ChIP
FO1417	TCTGATCTTAGAGTTGTCG	+1584/+2601	<i>PDC1</i> ChIP
FO1756	GGTAATTAA TCAGCGAAG CGATGA TTT	-190/-163	<i>GALI</i> ChIP
FO1757	TGCGCTAGA ATTGAACTC AGGTAC	+58/+82	<i>GALI</i> ChIP
FO1760	GTTATCATATGTCCAAGCAAAA	+590/+612	<i>GALI</i> ChIP
FO1761	TCCAGAAAGTAAAACAACACC	+877/+898	<i>GALI</i> ChIP
FO1764	ATTTGTTCCATTGCTTTGTC	+1370/+1390	<i>GALI</i> ChIP
FO1765	CTACTCGTTATTATTGCGTATTTT	+1657/+1681	<i>GALI</i> ChIP
FO1326	ATGATCAACTTAGCCAGGACAT CCATA	-246/-220	<i>SCR1</i> probe
FO1327	GTTCAACTAGCGAAGCCGCCAAA TTAA	+441/+468	<i>SCR1</i> probe
FO2308	TGATGCCACTAAGGATGAGAAT AA	+1515/+1538	<i>FLO8</i> probe
FO2309	GGTCTTCAACCATAACCAATATTCC	+2303/+2324	<i>FLO8</i> probe

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