

## SUPPLEMENTAL DATA

**Supplemental Table 1. KEGG NF- $\kappa$ B pathway genes correlating with HOXA1 expression in breast cancer datasets.** All KEGG NF- $\kappa$ B pathway genes that show significant correlation with HOXA1 mRNA expression in the 12 breast cancer datasets with significant over-representation of the KEGG NF- $\kappa$ B pathway. The first two columns represent the NCBI gene name and ID. Column 3 shows the dataset (details in Table 1). Columns 4 and 5 show the Correlation ( $R$ ) and Significance ( $P$ ) results of a 2log-Pearson correlation test. Column 6 denotes whether high HOXA1 expression correlates with high (positive) or low (negative) expression of the listed gene. Column 7 shows the array probe selected as in Materials and Methods. Genes with a negative correlation are in red type. Genes consistently correlated in  $\geq 50\%$  of datasets are on green or yellow field (positive or negative correlation, respectively).

**Supplemental Figure 1. NF- $\kappa$ B pathway genes with consistent correlation to HOXA1 expression in breast cancer.** Graphical overview of the NF- $\kappa$ B pathway showing pathway genes with a significant correlation to HOXA1 mRNA expression as in Figure 1, but now for a total of 12 breast cancer datasets. The KEGG NF- $\kappa$ B pathway graph is overlaid with boxes representing positively or negatively correlating genes only if a significant, consistent correlation was found in at least 50% of datasets analysed. Other details are as in Figure 1 and Supplemental Table 1. The overview shows a very consistent positive correlation with HOXA1 mRNA expression, especially for genes acting at the cell surface (activating ligands and receptors; the upstream part of the pathway) and for target genes in the nucleus (the downstream part of the pathway), strongly suggesting that the NF- $\kappa$ B pathway is activated by HOXA1 in breast cancer cells.

**Supplemental Table 2. Primers used to generate HOXA1 deletion mutants.**

**Supplemental Figure 2. Impact of HOXA1 on NF- $\kappa$ B reporter.** Western Blot showing IKK $\alpha$ , IKK $\beta$ , p-IKK $\alpha/\beta$  (Ser176/180), p65 and p-p65 (Ser536) proteins from MCF10A cells transfected with a FLAG-HOXA1 expression plasmid or with a negative control, and treated with TNF $\alpha$  (25 ng/ml) for 0, 10, 15, 20, 30 or 45 minutes. Detection of  $\beta$ -actin was used as loading control. Bands corresponding to detected proteins were cropped for the sake of presentation but all samples were processed in parallel, being run together on single protein gels.

**Supplemental Figure 3. Impact of HOXA1 pentapeptide insertions on activation of the NF- $\kappa$ B reporter.** Activity of the HOXA1 pentapeptide insertion mutants on the NF- $\kappa$ B reporter. Random pentapeptide insertion mutants were called from P1 to P18 as described previously [44], each arrow represents an insertion, with P1 and P18 as the most N-terminal and C-terminal insertions, respectively. Results were obtained as described in the legend to Figure 2.

Gene name	GeneID	Dataset	R	P	Sign	Probe
ATM	472	Bertucci-266	0,136	2,694E-02	Positive	210858_x_at
ATM	472	Black-107	0,250	9,385E-03	Positive	212672_at
ATM	472	Booser-508	0,308	1,176E-12	Positive	210858_x_at
ATM	472	Bos-204	0,242	4,891E-04	Positive	212672_at
ATM	472	Clynes-121	0,310	5,416E-04	Positive	212672_at
ATM	472	Iglehart-123	0,196	2,941E-02	Positive	210858_x_at
ATM	472	Jonsdottir-94	0,238	2,076E-02	Positive	ILMN_1779214
ATM	472	Minn-96	0,291	4,022E-03	Positive	210858_x_at
ATM	472	Smid-210	0,241	4,216E-04	Positive	212672_at
ATM	472	Sotiriou-120	0,238	8,797E-03	Positive	212672_at
BCL10	8915	Bertucci-266	0,145	1,800E-02	Positive	205263_at
BCL10	8915	Bos-204	0,207	2,950E-03	Positive	205263_at
BCL10	8915	Smid-210	0,226	9,764E-04	Positive	205263_at
BCL2	596	Bertucci-266	-0,247	4,781E-05	Negative	203685_at
BCL2	596	Booser-508	-0,186	2,397E-05	Negative	203685_at
BCL2	596	Chin-124	-0,323	2,531E-04	Negative	203685_at
BCL2	596	Minn-96	-0,262	1,000E-02	Negative	203685_at
BCL2	596	Smid-210	-0,172	1,256E-02	Negative	203685_at
BCL2	596	Sotiriou-198	-0,195	5,877E-03	Negative	203685_at
BCL2	596	Wang-286	-0,279	1,677E-06	Negative	203685_at
BCL2A1	597	Bertucci-266	0,289	1,619E-06	Positive	205681_at
BCL2A1	597	Booser-508	0,145	1,025E-03	Positive	205681_at
BCL2A1	597	Bos-204	0,194	5,528E-03	Positive	205681_at
BCL2A1	597	Chin-124	0,184	4,088E-02	Positive	205681_at
BCL2A1	597	Clynes-121	0,306	6,303E-04	Positive	205681_at
BCL2A1	597	EXPO-351	0,186	4,476E-04	Positive	205681_at
BCL2A1	597	Iglehart-123	0,318	3,392E-04	Positive	205681_at
BCL2A1	597	Minn-96	0,206	4,367E-02	Positive	205681_at
BCL2A1	597	Smid-210	0,192	5,302E-03	Positive	205681_at
BCL2A1	597	Wang-286	0,182	1,980E-03	Positive	205681_at
BCL2L1	598	Bos-204	-0,236	6,841E-04	Negative	212312_at
BCL2L1	598	Clynes-121	-0,182	4,549E-02	Negative	212312_at
BCL2L1	598	EXPO-351	-0,163	2,239E-03	Negative	212312_at
BCL2L1	598	Minn-96	-0,385	1,065E-04	Negative	212312_at
BCL2L1	598	Smid-210	-0,211	2,145E-03	Negative	212312_at
BCL2L1	598	Sotiriou-198	-0,206	3,629E-03	Negative	212312_at
BCL2L1	598	Wang-286	-0,150	1,124E-02	Negative	212312_at
BIRC2		Bertucci-266	0,224	2,270E-04	Positive	202076_at
BIRC2	329	Booser-508	0,115	9,386E-03	Positive	202076_at
BIRC2	329	Bos-204	0,255	2,286E-04	Positive	202076_at
BIRC2	329	Clynes-121	0,264	3,428E-03	Positive	202076_at
BIRC2	329	Jonsdottir-94	0,244	1,758E-02	Positive	ILMN_2182704
BIRC2	329	Smid-210	0,260	1,407E-04	Positive	202076_at
BIRC2	329	Sotiriou-198	0,258	2,471E-04	Positive	202076_at
BIRC2	329	Wang-286	0,238	4,788E-05	Positive	202076_at
BIRC3	330	Bertucci-266	0,340	1,243E-08	Positive	210538_s_at
BIRC3	330	Booser-508	0,162	2,526E-04	Positive	210538_s_at
BIRC3	330	Bos-204	0,260	1,710E-04	Positive	210538_s_at
BIRC3	330	Chin-124	0,254	4,416E-03	Positive	210538_s_at
BIRC3	330	Clynes-121	0,321	3,287E-04	Positive	210538_s_at
BIRC3	330	EXPO-351	0,166	1,786E-03	Positive	210538_s_at
BIRC3	330	Iglehart-123	0,313	4,315E-04	Positive	210538_s_at
BIRC3	330	Smid-210	0,257	1,679E-04	Positive	210538_s_at
BIRC3	330	Wang-286	0,145	1,427E-02	Positive	210538_s_at
BLNK	29760	Minn-96	-0,211	3,914E-02	Negative	207655_s_at
BLNK	29760	Sotiriou-120	0,222	1,473E-02	Positive	207655_s_at
BTK	695	Bertucci-266	0,189	1,998E-03	Positive	205504_at
BTK	695	Black-107	0,270	4,840E-03	Positive	205504_at

Gene name	GeneID	Dataset	R	P	Sign	Probe
BTK	695	Booser-508	0,291	2,118E-11	Positive	205504_at
BTK	695	Bos-204	0,202	3,800E-03	Positive	205504_at
BTK	695	Chin-124	0,248	5,400E-03	Positive	205504_at
BTK	695	Clynes-121	0,329	2,280E-04	Positive	205504_at
BTK	695	EXPO-351	0,165	1,928E-03	Positive	205504_at
BTK	695	Iglehart-123	0,268	2,779E-03	Positive	205504_at
BTK	695	Jonsdottir-94	0,237	2,119E-02	Positive	ILMN_1662026
BTK	695	Smid-210	0,169	1,445E-02	Positive	205504_at
BTK	695	Sotiriou-120	0,281	1,890E-03	Positive	205504_at
BTK	695	Wang-286	0,217	2,178E-04	Positive	205504_at
CARD11	84433	Bertucci-266	0,204	8,395E-04	Positive	223514_at
CARD11	84433	Bos-204	0,140	4,540E-02	Positive	223514_at
CARD11	84433	Clynes-121	0,319	3,634E-04	Positive	223514_at
CARD11	84433	Jonsdottir-94	0,216	3,624E-02	Positive	ILMN_1721978
CARD11	84433	Smid-210	0,166	1,617E-02	Positive	223514_at
CCL13	6357	Bertucci-266	0,281	3,189E-06	Positive	206407_s_at
CCL13	6357	Booser-508	0,203	3,874E-06	Positive	206407_s_at
CCL13	6357	Bos-204	0,163	2,002E-02	Positive	206407_s_at
CCL13	6357	Chin-124	0,259	3,644E-03	Positive	206407_s_at
CCL13	6357	EXPO-351	0,193	2,812E-04	Positive	206407_s_at
CCL13	6357	Jonsdottir-94	0,230	2,593E-02	Positive	ILMN_1783593
CCL13	6357	Minn-96	0,401	5,131E-05	Positive	206407_s_at
CCL13	6357	Smid-210	0,174	1,144E-02	Positive	206407_s_at
CCL13	6357	Sotiriou-120	0,181	4,769E-02	Positive	206407_s_at
CCL13	6357	Wang-286	0,155	8,582E-03	Positive	206407_s_at
CCL19	6363	Bertucci-266	0,288	1,769E-06	Positive	210072_at
CCL19	6363	Black-107	0,224	2,020E-02	Positive	210072_at
CCL19	6363	Booser-508	0,145	1,075E-03	Positive	210072_at
CCL19	6363	Bos-204	0,249	3,268E-04	Positive	210072_at
CCL19	6363	Chin-124	0,234	8,760E-03	Positive	210072_at
CCL19	6363	Clynes-121	0,236	9,123E-03	Positive	210072_at
CCL19	6363	EXPO-351	0,164	2,083E-03	Positive	210072_at
CCL19	6363	Iglehart-123	0,308	5,374E-04	Positive	210072_at
CCL19	6363	Jonsdottir-94	0,204	4,873E-02	Positive	ILMN_1769129
CCL19	6363	Smid-210	0,215	1,753E-03	Positive	210072_at
CCL19	6363	Wang-286	0,205	4,788E-04	Positive	210072_at
CCL21	6366	Bertucci-266	0,314	1,634E-07	Positive	204606_at
CCL21	6366	Booser-508	0,155	4,450E-04	Positive	204606_at
CCL21	6366	Bos-204	0,240	5,452E-04	Positive	204606_at
CCL21	6366	Chin-124	0,282	1,494E-03	Positive	204606_at
CCL21	6366	Clynes-121	0,319	3,571E-04	Positive	204606_at
CCL21	6366	Jonsdottir-94	0,282	5,812E-03	Positive	ILMN_1677505
CCL21	6366	Smid-210	0,195	4,569E-03	Positive	204606_at
CCL21	6366	Sotiriou-120	0,239	8,622E-03	Positive	204606_at
CCL21	6366	Wang-286	0,120	4,316E-02	Positive	204606_at
CCL4	6351	Bertucci-266	0,242	6,475E-05	Positive	204103_at
CCL4	6351	Black-107	0,249	9,806E-03	Positive	204103_at
CCL4	6351	Booser-508	0,130	3,236E-03	Positive	204103_at
CCL4	6351	Bos-204	0,207	2,917E-03	Positive	204103_at
CCL4	6351	Clynes-121	0,337	1,563E-04	Positive	204103_at
CCL4	6351	Smid-210	0,217	1,555E-03	Positive	204103_at
CCL4L2	9560	Jonsdottir-94	0,209	4,339E-02	Positive	ILMN_1716276
CD14	929	Bertucci-266	0,227	1,949E-04	Positive	201743_at
CD14	929	Black-107	0,279	3,588E-03	Positive	201743_at
CD14	929	Booser-508	0,213	1,344E-06	Positive	201743_at
CD14	929	Bos-204	0,230	9,327E-04	Positive	201743_at
CD14	929	Clynes-121	0,308	6,006E-04	Positive	201743_at
CD14	929	EXPO-351	0,188	3,889E-04	Positive	201743_at

Gene name	GeneID	Dataset	R	P	Sign	Probe
CD14	929	Iglehart-123	0,180	4,644E-02	Positive	201743_at
CD14	929	Minn-96	0,386	1,042E-04	Positive	201743_at
CD14	929	Servant-343	0,147	6,433E-03	Positive	ILMN_2396444
CD14	929	Smid-210	0,220	1,348E-03	Positive	201743_at
CD14	929	Sotiriou-120	0,360	5,354E-05	Positive	201743_at
CD14	929	Wang-286	0,216	2,400E-04	Positive	201743_at
CD40	958	Bertucci-266	0,238	8,927E-05	Positive	35150_at
CD40	958	Black-107	0,267	5,487E-03	Positive	35150_at
CD40	958	Booser-508	0,319	1,625E-13	Positive	35150_at
CD40	958	Chin-124	0,309	4,854E-04	Positive	215346_at
CD40	958	Clynes-121	0,284	1,608E-03	Positive	215346_at
CD40	958	EXPO-351	0,148	5,403E-03	Positive	35150_at
CD40	958	Jonsdottir-94	0,398	7,202E-05	Positive	ILMN_1779257
CD40	958	Minn-96	0,330	1,044E-03	Positive	35150_at
CD40LG	959	Bertucci-266	0,219	3,195E-04	Positive	207892_at
CD40LG	959	Booser-508	0,210	1,711E-06	Positive	207892_at
CD40LG	959	Bos-204	0,179	1,057E-02	Positive	207892_at
CD40LG	959	Iglehart-123	0,334	1,618E-04	Positive	207892_at
CD40LG	959	Jonsdottir-94	0,290	4,536E-03	Positive	ILMN_1659077
CD40LG	959	Smid-210	0,136	4,970E-02	Positive	207892_at
CFLAR	8837	Bertucci-266	0,256	2,407E-05	Positive	239629_at
CFLAR	8837	Booser-508	0,302	3,661E-12	Positive	211316_x_at
CFLAR	8837	Bos-204	0,236	6,795E-04	Positive	209939_x_at
CFLAR	8837	Chin-124	0,265	2,963E-03	Positive	209939_x_at
CFLAR	8837	EXPO-351	0,211	6,555E-05	Positive	239629_at
CFLAR	8837	Iglehart-123	0,221	1,387E-02	Positive	209939_x_at
CFLAR	8837	Smid-210	0,220	1,336E-03	Positive	209939_x_at
CFLAR	8837	Wang-286	0,286	8,733E-07	Positive	209939_x_at
<b>CHUK</b>	<b>1147</b>	<b>Bos-204</b>	<b>-0,143</b>	<b>4,151E-02</b>	<b>Negative</b>	<b>209666_s_at</b>
CSNK2A2	1459	Bertucci-266	0,390	4,229E-11	Positive	224922_at
CSNK2A2	1459	Black-107	0,210	3,013E-02	Positive	224922_at
CSNK2A2	1459	Booser-508	0,116	9,078E-03	Positive	203575_at
CSNK2A2	1459	EXPO-351	0,180	7,147E-04	Positive	224922_at
CSNK2A2	1459	Jonsdottir-94	0,218	3,452E-02	Positive	ILMN_1723843
CSNK2A2	1459	Sotiriou-120	0,189	3,858E-02	Positive	224922_at
CSNK2A2	1459	Sotiriou-198	0,252	3,355E-04	Positive	203575_at
CSNK2A2	1459	Wang-286	0,294	4,262E-07	Positive	203575_at
<b>CSNK2B</b>	<b>1460</b>	<b>Booser-508</b>	<b>-0,137</b>	<b>2,027E-03</b>	<b>Negative</b>	<b>201390_s_at</b>
CSNK2B	1460	Jonsdottir-94	0,329	1,222E-03	Positive	ILMN_1800461
CXCL12	6387	Bertucci-266	0,259	1,892E-05	Positive	209687_at
CXCL12	6387	Black-107	0,391	3,056E-05	Positive	209687_at
CXCL12	6387	Bos-204	0,220	1,543E-03	Positive	209687_at
CXCL12	6387	Chin-124	0,210	1,946E-02	Positive	209687_at
CXCL12	6387	Clynes-121	0,304	6,909E-04	Positive	209687_at
CXCL12	6387	EXPO-351	0,186	4,631E-04	Positive	209687_at
CXCL12	6387	Servant-343	0,114	3,523E-02	Positive	ILMN_1791447
CXCL12	6387	Smid-210	0,170	1,378E-02	Positive	209687_at
CXCL12	6387	Sotiriou-120	0,421	1,673E-06	Positive	209687_at
CXCL12	6387	Sotiriou-198	0,155	2,954E-02	Positive	209687_at
CXCL2	2920	Bertucci-266	0,333	2,569E-08	Positive	209774_x_at
CXCL2	2920	Black-107	0,206	3,361E-02	Positive	209774_x_at
CXCL2	2920	Booser-508	0,198	6,920E-06	Positive	209774_x_at
CXCL2	2920	Bos-204	0,442	3,600E-11	Positive	209774_x_at
CXCL2	2920	Chin-124	0,345	8,711E-05	Positive	209774_x_at
CXCL2	2920	Clynes-121	0,213	1,915E-02	Positive	209774_x_at
CXCL2	2920	EXPO-351	0,195	2,290E-04	Positive	209774_x_at
CXCL2	2920	Iglehart-123	0,268	2,766E-03	Positive	209774_x_at
CXCL2	2920	Minn-96	0,284	5,104E-03	Positive	209774_x_at

Gene name	GeneID	Dataset	R	P	Sign	Probe
CXCL2	2920	Smid-210	0,407	8,978E-10	Positive	209774_x_at
CXCL2	2920	Sotiriou-198	0,290	3,360E-05	Positive	209774_x_at
CXCL2	2920	Wang-286	0,247	2,319E-05	Positive	209774_x_at
DDX58	23586	Bos-204	0,211	2,490E-03	Positive	218943_s_at
DDX58	23586	Smid-210	0,226	9,475E-04	Positive	218943_s_at
ERC1	23085	Bertucci-266	0,138	2,420E-02	Positive	215606_s_at
ERC1	23085	Booser-508	0,223	3,864E-07	Positive	215606_s_at
ERC1	23085	EXPO-351	0,135	1,139E-02	Positive	226049_at
GADD45B	4616	Bertucci-266	0,207	7,000E-04	Positive	207574_s_at
GADD45B	4616	Bos-204	0,153	2,927E-02	Positive	207574_s_at
GADD45B	4616	Smid-210	0,139	4,396E-02	Positive	207574_s_at
GADD45B	4616	Sotiriou-120	0,269	2,929E-03	Positive	207574_s_at
GADD45B	4616	Wang-286	0,214	2,673E-04	Positive	207574_s_at
ICAM1	3383	Bertucci-266	0,357	1,958E-09	Positive	202637_s_at
ICAM1	3383	Booser-508	0,194	1,052E-05	Positive	202637_s_at
ICAM1	3383	Bos-204	0,295	1,859E-05	Positive	202637_s_at
ICAM1	3383	Chin-124	0,207	2,092E-02	Positive	202637_s_at
ICAM1	3383	Clynes-121	0,350	8,457E-05	Positive	202637_s_at
ICAM1	3383	EXPO-351	0,120	2,485E-02	Positive	202637_s_at
ICAM1	3383	Minn-96	0,337	7,788E-04	Positive	202637_s_at
ICAM1	3383	Smid-210	0,280	3,882E-05	Positive	202637_s_at
ICAM1	3383	Sotiriou-198	0,151	3,378E-02	Positive	202637_s_at
ICAM1	3383	Wang-286	0,270	3,489E-06	Positive	202637_s_at
IKBKB	3351	Bertucci-266	-0,154	1,181E-02	Negative	209341_s_at
IKBKB	3351	Booser-508	-0,199	6,332E-06	Negative	209341_s_at
IKBKB	3351	Chin-124	-0,182	4,360E-02	Negative	209341_s_at
IKBKB	3351	EXPO-351	-0,152	4,388E-03	Negative	209341_s_at
IKBKB	3351	Minn-96	-0,219	3,180E-02	Negative	209341_s_at
IKBKB	3351	Wang-286	-0,222	1,524E-04	Negative	209341_s_at
IKBKG	8517	Booser-508	0,229	1,824E-07	Positive	36004_at
IKBKG	8517	Servant-343	0,115	3,386E-02	Positive	ILMN_1707308
IL1B	3553	Bertucci-266	0,233	1,260E-04	Positive	205067_at
IL1B	3553	Black-107	0,232	1,605E-02	Positive	205067_at
IL1B	3553	Booser-508	0,245	2,212E-08	Positive	205067_at
IL1B	3553	Bos-204	0,310	6,314E-06	Positive	205067_at
IL1B	3553	Clynes-121	0,282	1,696E-03	Positive	205067_at
IL1B	3553	Minn-96	0,219	3,207E-02	Positive	205067_at
IL1B	3553	Smid-210	0,299	1,008E-05	Positive	205067_at
IL1B	3553	Wang-286	0,129	2,975E-02	Positive	205067_at
IL1R1	3554	Bertucci-266	0,273	6,353E-06	Positive	202948_at
IL1R1	3554	Black-107	0,355	1,732E-04	Positive	202948_at
IL1R1	3554	Booser-508	0,128	3,826E-03	Positive	202948_at
IL1R1	3554	Bos-204	0,188	7,044E-03	Positive	202948_at
IL1R1	3554	Chin-124	0,316	3,548E-04	Positive	202948_at
IL1R1	3554	EXPO-351	0,254	1,438E-06	Positive	202948_at
IL1R1	3554	Minn-96	0,301	2,921E-03	Positive	202948_at
IL1R1	3554	Sotiriou-120	0,410	3,255E-06	Positive	202948_at
IL1R1	3554	Sotiriou-198	0,265	1,589E-04	Positive	202948_at
IL1R1	3554	Wang-286	0,169	4,134E-03	Positive	202948_at
IL8 (CXCL8)	3576	Bertucci-266	0,237	9,382E-05	Positive	202859_x_at
IL8 (CXCL8)	3576	Booser-508	0,175	7,428E-05	Positive	202859_x_at
IL8 (CXCL8)	3576	Bos-204	0,245	4,071E-04	Positive	202859_x_at
IL8 (CXCL8)	3576	Chin-124	0,187	3,757E-02	Positive	202859_x_at
IL8 (CXCL8)	3576	Clynes-121	0,280	1,838E-03	Positive	202859_x_at
IL8 (CXCL8)	3576	Jonsdottir-94	0,279	6,495E-03	Positive	ILMN_2184373
IL8 (CXCL8)	3576	Minn-96	0,269	8,062E-03	Positive	202859_x_at
IL8 (CXCL8)	3576	Smid-210	0,220	1,360E-03	Positive	202859_x_at
IL8 (CXCL8)	3576	Wang-286	0,270	3,649E-06	Positive	202859_x_at

Gene name	GeneID	Dataset	R	P	Sign	Probe
IRAK1	3654	Bertucci-266	0,196	1,302E-03	Positive	201587_s_at
IRAK1	3654	Bos-204	0,175	1,212E-02	Positive	201587_s_at
IRAK1	3654	Chin-124	0,208	2,026E-02	Positive	201587_s_at
IRAK1	3654	EXPO-351	0,131	1,422E-02	Positive	201587_s_at
IRAK1	3654	Smid-210	0,197	4,136E-03	Positive	201587_s_at
IRAK1	3654	Sotiriou-198	0,146	4,059E-02	Positive	201587_s_at
IRAK1	3654	Wang-286	0,242	3,398E-05	Positive	201587_s_at
IRAK4	51135	Bertucci-266	-0,135	2,822E-02	Negative	219618_at
IRAK4	51135	Booser-508	0,155	4,524E-04	Positive	219618_at
LAT	27040	Booser-508	0,091	3,942E-02	Positive	209881_s_at
LAT	27040	Bos-204	0,151	3,128E-02	Positive	209881_s_at
LAT	27040	Chin-124	0,302	6,518E-04	Positive	211005_at
LAT	27040	Jonsdottir-94	0,204	4,834E-02	Positive	ILMN_2404625
LAT	27040	Servant-343	0,114	3,542E-02	Positive	ILMN_2404625
LAT	27040	Smid-210	0,171	1,334E-02	Positive	209881_s_at
LBP	3929	Bertucci-266	0,253	2,889E-05	Positive	214461_at
LBP	3929	Booser-508	0,147	8,971E-04	Positive	214461_at
LBP	3929	Bos-204	0,188	6,998E-03	Positive	214461_at
LBP	3929	Chin-124	0,202	2,421E-02	Positive	214461_at
LBP	3929	EXPO-351	0,119	2,529E-02	Positive	211652_s_at
LBP	3929	Smid-210	0,205	2,800E-03	Positive	214461_at
LBP	3929	Wang-286	0,189	1,303E-03	Positive	214461_at
LCK	3932	Bertucci-266	0,374	3,078E-10	Positive	204891_s_at
LCK	3932	Black-107	0,263	6,149E-03	Positive	204891_s_at
LCK	3932	Booser-508	0,165	1,802E-04	Positive	204890_s_at
LCK	3932	Bos-204	0,220	1,607E-03	Positive	204891_s_at
LCK	3932	Chin-124	0,236	8,320E-03	Positive	204890_s_at
LCK	3932	Clynes-121	0,324	2,841E-04	Positive	204891_s_at
LCK	3932	EXPO-351	0,143	7,094E-03	Positive	204891_s_at
LCK	3932	Iglehart-123	0,333	1,688E-04	Positive	204891_s_at
LCK	3932	Smid-210	0,201	3,406E-03	Positive	204891_s_at
LTA	4049	Bertucci-266	0,183	2,722E-03	Positive	206975_at
LTA	4049	Booser-508	0,257	4,286E-09	Positive	206975_at
LTA	4049	Jonsdottir-94	0,232	2,427E-02	Positive	ILMN_1795464
LTB	4050	Bertucci-266	0,240	7,892E-05	Positive	207339_s_at
LTB	4050	Booser-508	0,141	1,424E-03	Positive	207339_s_at
LTB	4050	Bos-204	0,236	6,945E-04	Positive	207339_s_at
LTB	4050	Chin-124	0,252	4,697E-03	Positive	207339_s_at
LTB	4050	Clynes-121	0,276	2,217E-03	Positive	207339_s_at
LTB	4050	Iglehart-123	0,371	2,431E-05	Positive	207339_s_at
LTB	4050	Jonsdottir-94	0,216	3,654E-02	Positive	ILMN_2376205
LTB	4050	Minn-96	0,207	4,266E-02	Positive	207339_s_at
LTB	4050	Smid-210	0,235	5,906E-04	Positive	207339_s_at
LTB	4050	Sotiriou-120	0,220	1,575E-02	Positive	207339_s_at
LTB	4050	Wang-286	0,227	1,101E-04	Positive	207339_s_at
LTBR	4055	EXPO-351	0,117	2,779E-02	Positive	203005_at
LTBR	4055	Jonsdottir-94	0,292	4,288E-03	Positive	ILMN_1667476
LTBR	4055	Servant-343	-0,124	2,113E-02	Negative	ILMN_1667476
LTBR	4055	Sotiriou-120	-0,225	1,358E-02	Negative	203005_at
LTBR	4055	Wang-286	0,198	7,666E-04	Positive	203005_at
LY96	23643	Bertucci-266	0,299	6,633E-07	Positive	206584_at
LY96	23643	Black-107	0,379	5,622E-05	Positive	206584_at
LY96	23643	Booser-508	0,253	7,049E-09	Positive	206584_at
LY96	23643	Bos-204	0,342	5,643E-07	Positive	206584_at
LY96	23643	Chin-124	0,307	5,218E-04	Positive	206584_at
LY96	23643	Clynes-121	0,464	8,023E-08	Positive	206584_at
LY96	23643	EXPO-351	0,249	2,432E-06	Positive	206584_at
LY96	23643	Iglehart-123	0,282	1,585E-03	Positive	206584_at

Gene name	GeneID	Dataset	R	P	Sign	Probe
LY96	23643	Jonsdottir-94	0,289	4,714E-03	Positive	ILMN_1724533
LY96	23643	Smid-210	0,313	3,796E-06	Positive	206584_at
LY96	23643	Sotiriou-120	0,312	5,233E-04	Positive	206584_at
LY96	23643	Sotiriou-198	0,142	4,618E-02	Positive	206584_at
LY96	23643	Wang-286	0,231	7,930E-05	Positive	206584_at
LYN	4067	Bertucci-266	0,320	9,757E-08	Positive	210754_s_at
LYN	4067	Black-107	0,220	2,302E-02	Positive	202625_at
LYN	4067	Booser-508	0,261	2,245E-09	Positive	210754_s_at
LYN	4067	Bos-204	0,395	4,825E-09	Positive	210754_s_at
LYN	4067	Chin-124	0,259	3,684E-03	Positive	202625_at
LYN	4067	Clynes-121	0,387	1,151E-05	Positive	202625_at
LYN	4067	EXPO-351	0,223	2,478E-05	Positive	210754_s_at
LYN	4067	Iglehart-123	0,346	8,992E-05	Positive	210754_s_at
LYN	4067	Jonsdottir-94	0,215	3,727E-02	Positive	ILMN_1781155
LYN	4067	Minn-96	0,334	8,722E-04	Positive	210754_s_at
LYN	4067	Smid-210	0,405	1,086E-09	Positive	210754_s_at
LYN	4067	Sotiriou-198	0,202	4,267E-03	Positive	202626_s_at
LYN	4067	Wang-286	0,336	5,434E-09	Positive	202626_s_at
MALT1	10892	Bertucci-266	0,322	7,818E-08	Positive	210017_at
MALT1	10892	Booser-508	0,220	5,416E-07	Positive	210018_x_at
MALT1	10892	Bos-204	0,227	1,094E-03	Positive	210017_at
MALT1	10892	Chin-124	0,232	9,663E-03	Positive	210017_at
MALT1	10892	EXPO-351	0,112	3,593E-02	Positive	208309_s_at
MALT1	10892	Iglehart-123	0,225	1,229E-02	Positive	208309_s_at
MALT1	10892	Smid-210	0,167	1,550E-02	Positive	210017_at
MALT1	10892	Sotiriou-198	0,194	6,091E-03	Positive	210018_x_at
MALT1	10892	Wang-286	0,168	4,366E-03	Positive	210018_x_at
MAP3K14	9020	Bertucci-266	0,247	4,636E-05	Positive	205192_at
MAP3K14	9020	Black-107	0,209	3,074E-02	Positive	205192_at
MAP3K14	9020	Booser-508	0,182	3,794E-05	Positive	205192_at
MAP3K14	9020	Bos-204	0,265	1,313E-04	Positive	205192_at
MAP3K14	9020	EXPO-351	0,119	2,595E-02	Positive	205192_at
MAP3K14	9020	Jonsdottir-94	0,397	7,368E-05	Positive	ILMN_1724070
MAP3K14	9020	Smid-210	0,249	2,646E-04	Positive	205192_at
MAP3K14	9020	Wang-286	0,116	4,924E-02	Positive	205192_at
MAP3K7	6885	Bertucci-266	0,226	2,000E-04	Positive	206854_s_at
MAP3K7	6885	Booser-508	0,163	2,165E-04	Positive	206853_s_at
MAP3K7	6885	Bos-204	0,182	9,069E-03	Positive	206854_s_at
MAP3K7	6885	EXPO-351	0,133	1,291E-02	Positive	206854_s_at
MAP3K7	6885	Iglehart-123	0,194	3,187E-02	Positive	211537_x_at
MAP3K7	6885	Smid-210	0,203	3,081E-03	Positive	206854_s_at
MYD88	4615	Bertucci-266	0,255	2,587E-05	Positive	209124_at
MYD88	4615	Booser-508	0,189	1,838E-05	Positive	209124_at
MYD88	4615	Bos-204	0,247	3,590E-04	Positive	209124_at
MYD88	4615	Chin-124	0,193	3,146E-02	Positive	209124_at
MYD88	4615	Jonsdottir-94	0,217	3,598E-02	Positive	ILMN_1738523
MYD88	4615	Smid-210	0,235	5,801E-04	Positive	209124_at
MYD88	4615	Wang-286	0,147	1,296E-02	Positive	209124_at
NFKB1	4790	Bertucci-266	0,127	3,857E-02	Positive	209239_at
NFKB1	4790	Booser-508	0,136	2,059E-03	Positive	209239_at
NFKB1	4790	Servant-343	0,126	1,961E-02	Positive	ILMN_1714965
NFKB1	4790	Wang-286	-0,155	8,856E-03	Negative	209239_at
NFKB2	4791	Bertucci-266	0,128	3,647E-02	Positive	207535_s_at
NFKB2	4791	Booser-508	0,121	6,352E-03	Positive	207535_s_at
NFKB2	4791	Servant-343	-0,118	2,859E-02	Negative	ILMN_1799062
NFKBIA	4792	EXPO-351	0,110	3,875E-02	Positive	201502_s_at
NFKBIA	4792	Servant-343	0,121	2,480E-02	Positive	ILMN_1773154
PARP1	142	Booser-508	-0,128	3,840E-03	Negative	208644_at

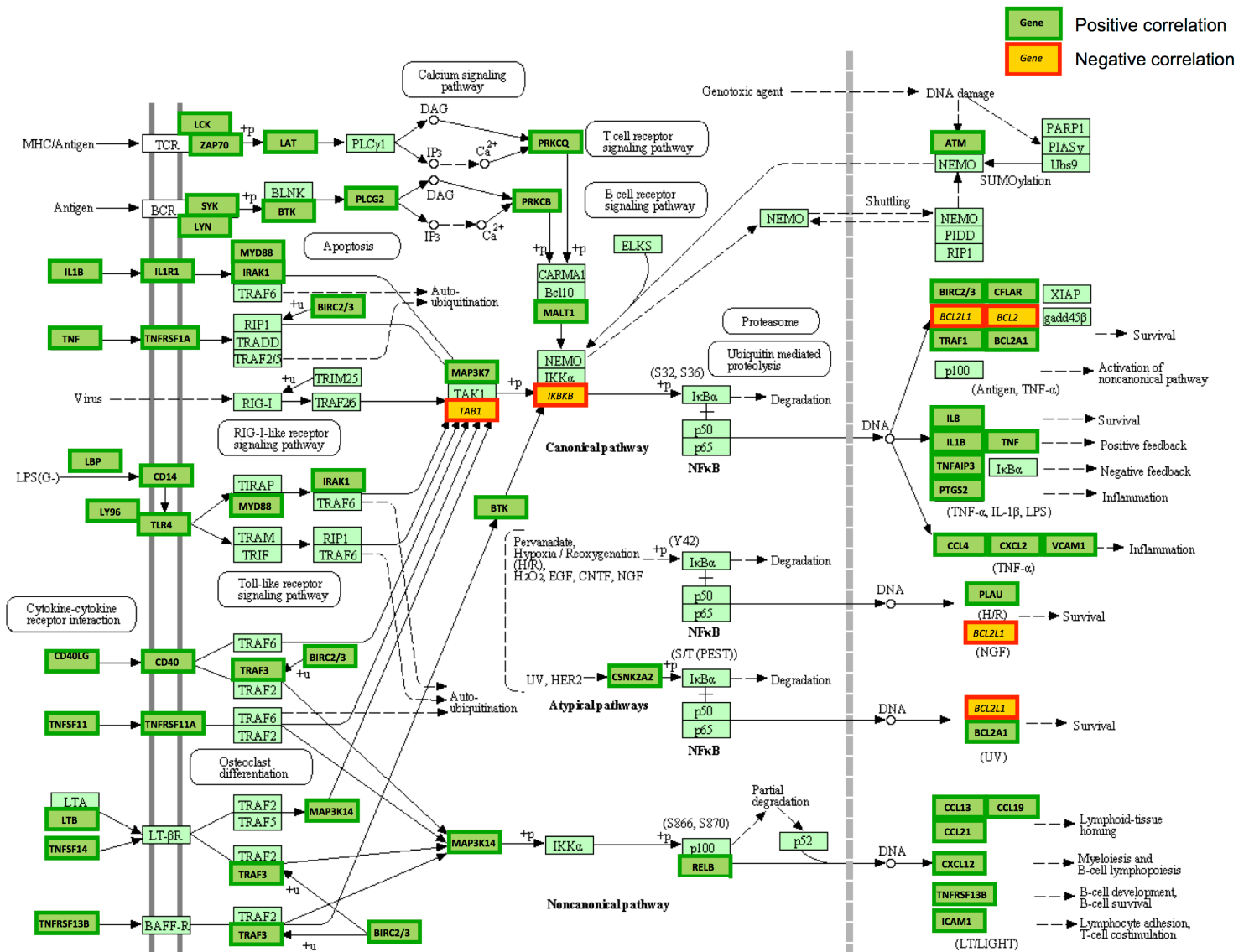
Gene name	GeneID	Dataset	R	P	Sign	Probe
PARP1	142	Iglehart-123	-0,277	1,958E-03	Negative	208644_at
PARP1	142	Minn-96	-0,219	3,191E-02	Negative	208644_at
PARP1	142	Sotiriou-120	-0,224	1,410E-02	Negative	208644_at
PARP1	142	Wang-286	-0,121	4,086E-02	Negative	208644_at
PIAS4	51588	Bertucci-266	-0,252	3,164E-05	Negative	212879_x_at
PIAS4	51588	Clynes-121	-0,184	4,282E-02	Negative	212879_x_at
PIAS4	51588	Iglehart-123	-0,194	3,164E-02	Negative	212879_x_at
PIAS4	51588	Minn-96	-0,267	8,502E-03	Negative	212881_at
PIAS4	51588	Servant-343	0,110	4,218E-02	Positive	ILMN_1802905
PIAS4	51588	Sotiriou-198	-0,183	9,822E-03	Negative	212879_x_at
PIDD	55367	Wang-286	-0,192	1,087E-03	Negative	221640_s_at
PLAU	5328	Bertucci-266	0,234	1,150E-04	Positive	205479_s_at
PLAU	5328	Black-107	0,235	1,498E-02	Positive	205479_s_at
PLAU	5328	Booser-508	0,179	4,962E-05	Positive	205479_s_at
PLAU	5328	Bos-204	0,308	7,430E-06	Positive	205479_s_at
PLAU	5328	Clynes-121	0,204	2,490E-02	Positive	205479_s_at
PLAU	5328	EXPO-351	0,118	2,711E-02	Positive	205479_s_at
PLAU	5328	Smid-210	0,289	2,159E-05	Positive	205479_s_at
PLAU	5328	Sotiriou-120	0,313	5,091E-04	Positive	205479_s_at
PLAU	5328	Wang-286	0,199	7,279E-04	Positive	205479_s_at
PLCG1	5335	Bertucci-266	0,208	6,537E-04	Positive	202789_at
PLCG1	5335	Booser-508	0,099	2,624E-02	Positive	202789_at
PLCG1	5335	Smid-210	0,157	2,309E-02	Positive	202789_at
PLCG2	5336	Bertucci-266	0,417	1,313E-12	Positive	204613_at
PLCG2	5336	Black-107	0,258	7,262E-03	Positive	204613_at
PLCG2	5336	Booser-508	0,352	2,852E-16	Positive	204613_at
PLCG2	5336	Bos-204	0,264	1,323E-04	Positive	204613_at
PLCG2	5336	Chin-124	0,285	1,321E-03	Positive	204613_at
PLCG2	5336	Clynes-121	0,302	7,620E-04	Positive	204613_at
PLCG2	5336	EXPO-351	0,280	9,536E-08	Positive	204613_at
PLCG2	5336	Iglehart-123	0,262	3,468E-03	Positive	204613_at
PLCG2	5336	Jonsdottir-94	0,376	1,870E-04	Positive	ILMN_1815719
PLCG2	5336	Minn-96	0,294	3,601E-03	Positive	204613_at
PLCG2	5336	Smid-210	0,228	8,692E-04	Positive	204613_at
PLCG2	5336	Sotiriou-120	0,207	2,325E-02	Positive	204613_at
PLCG2	5336	Sotiriou-198	0,173	1,483E-02	Positive	204613_at
PLCG2	5336	Wang-286	0,307	1,176E-07	Positive	204613_at
PRKCB	5579	Bertucci-266	0,316	1,394E-07	Positive	209685_s_at
PRKCB	5579	Black-107	0,255	8,052E-03	Positive	209685_s_at
PRKCB	5579	Booser-508	0,278	1,717E-10	Positive	209685_s_at
PRKCB	5579	Chin-124	0,249	5,316E-03	Positive	209685_s_at
PRKCB	5579	Clynes-121	0,390	9,629E-06	Positive	209685_s_at
PRKCB	5579	EXPO-351	0,183	5,789E-04	Positive	209685_s_at
PRKCB	5579	Iglehart-123	0,353	6,358E-05	Positive	209685_s_at
PRKCB	5579	Minn-96	0,235	2,113E-02	Positive	209685_s_at
PRKCB	5579	Sotiriou-120	0,260	4,063E-03	Positive	209685_s_at
PRKCB	5579	Wang-286	0,220	1,781E-04	Positive	209685_s_at
PRKCQ	5588	Bertucci-266	0,372	3,851E-10	Positive	210038_at
PRKCQ	5588	Black-107	0,220	2,290E-02	Positive	210038_at
PRKCQ	5588	Booser-508	0,300	4,843E-12	Positive	210038_at
PRKCQ	5588	Bos-204	0,181	9,694E-03	Positive	210038_at
PRKCQ	5588	Chin-124	0,269	2,502E-03	Positive	210038_at
PRKCQ	5588	EXPO-351	0,158	2,938E-03	Positive	210038_at
PRKCQ	5588	Jonsdottir-94	0,243	1,807E-02	Positive	ILMN_1733421
PRKCQ	5588	Minn-96	0,203	4,710E-02	Positive	210038_at
PRKCQ	5588	Smid-210	0,146	3,420E-02	Positive	210038_at
PRKCQ	5588	Wang-286	0,162	6,176E-03	Positive	210038_at
PTGS2	5743	Bertucci-266	0,260	1,758E-05	Positive	204748_at



Gene name	GeneID	Dataset	R	P	Sign	Probe
PTGS2	5743	Booser-508	0,211	1,649E-06	Positive	204748_at
PTGS2	5743	Bos-204	0,363	9,280E-08	Positive	204748_at
PTGS2	5743	Chin-124	0,282	1,503E-03	Positive	204748_at
PTGS2	5743	Clynes-121	0,264	3,445E-03	Positive	204748_at
PTGS2	5743	EXPO-351	0,105	4,833E-02	Positive	204748_at
PTGS2	5743	Minn-96	0,380	1,369E-04	Positive	204748_at
PTGS2	5743	Smid-210	0,320	2,171E-06	Positive	204748_at
PTGS2	5743	Sotiriou-198	0,275	8,921E-05	Positive	204748_at
PTGS2	5743	Wang-286	0,321	2,748E-08	Positive	204748_at
RELA	5970	Bertucci-266	0,135	2,746E-02	Positive	201783_s_at
RELA	5970	Jonsdottir-94	0,362	3,325E-04	Positive	ILMN_1705266
RELB	5971	Bertucci-266	0,319	1,038E-07	Positive	205205_at
RELB	5971	Booser-508	0,131	3,142E-03	Positive	205205_at
RELB	5971	Jonsdottir-94	0,418	2,721E-05	Positive	ILMN_1811258
RELB	5971	Minn-96	0,269	8,158E-03	Positive	205205_at
RELB	5971	Smid-210	0,139	4,379E-02	Positive	205205_at
RELB	5971	Wang-286	0,129	2,946E-02	Positive	205205_at
SYK	6850	Bertucci-266	0,278	4,136E-06	Positive	226068_at
SYK	6850	Black-107	0,239	1,315E-02	Positive	226068_at
SYK	6850	Booser-508	0,156	4,171E-04	Positive	207540_s_at
SYK	6850	Bos-204	0,212	2,380E-03	Positive	226068_at
SYK	6850	Clynes-121	0,220	1,550E-02	Positive	207540_s_at
SYK	6850	EXPO-351	0,160	2,696E-03	Positive	226068_at
SYK	6850	Iglehart-123	0,194	3,156E-02	Positive	226068_at
SYK	6850	Jonsdottir-94	0,311	2,265E-03	Positive	ILMN_2059549
SYK	6850	Minn-96	0,260	1,037E-02	Positive	207540_s_at
SYK	6850	Servant-343	0,111	4,029E-02	Positive	ILMN_2059549
SYK	6850	Smid-210	0,183	7,684E-03	Positive	226068_at
SYK	6850	Sotiriou-120	0,238	8,926E-03	Positive	207540_s_at
TAB1	10454	Bertucci-266	-0,137	2,494E-02	Negative	203901_at
TAB1	10454	Bos-204	-0,175	1,207E-02	Negative	203901_at
TAB1	10454	Clynes-121	-0,251	5,447E-03	Negative	203901_at
TAB1	10454	Minn-96	-0,203	4,741E-02	Negative	203901_at
TAB1	10454	Smid-210	-0,223	1,119E-03	Negative	203901_at
TAB1	10454	Wang-286	-0,149	1,188E-02	Negative	203901_at
TAB2	23118	Bertucci-266	0,236	1,040E-04	Positive	212184_s_at
TAB2	23118	Booser-508	0,154	5,011E-04	Positive	210284_s_at
TAB2	23118	Clynes-121	0,205	2,413E-02	Positive	212184_s_at
TAB2	23118	Minn-96	0,344	6,096E-04	Positive	210284_s_at
TAB2	23118	Wang-286	0,145	1,415E-02	Positive	212184_s_at
TIRAP	114609	Bos-204	-0,149	3,391E-02	Negative	1554091_a_at
TIRAP	114609	Smid-210	-0,142	3,913E-02	Negative	1554091_a_at
TLR4	7099	Bertucci-266	0,209	6,175E-04	Positive	224341_x_at
TLR4	7099	Black-107	0,256	7,810E-03	Positive	224341_x_at
TLR4	7099	Booser-508	0,266	1,045E-09	Positive	221060_s_at
TLR4	7099	Clynes-121	0,363	4,167E-05	Positive	224341_x_at
TLR4	7099	Iglehart-123	0,222	1,354E-02	Positive	221060_s_at
TLR4	7099	Jonsdottir-94	0,220	3,296E-02	Positive	ILMN_1706217
TLR4	7099	Minn-96	0,227	2,613E-02	Positive	221060_s_at
TLR4	7099	Sotiriou-120	0,299	9,182E-04	Positive	224341_x_at
TNF	7124	Bertucci-266	0,241	7,314E-05	Positive	207113_s_at
TNF	7124	Booser-508	0,244	2,670E-08	Positive	207113_s_at
TNF	7124	Bos-204	0,162	2,063E-02	Positive	207113_s_at
TNF	7124	Jonsdottir-94	0,263	1,057E-02	Positive	ILMN_1728106
TNF	7124	Smid-210	0,152	2,740E-02	Positive	207113_s_at
TNF	7124	Sotiriou-120	0,261	4,025E-03	Positive	207113_s_at
TNFAIP3	7128	Bertucci-266	0,405	6,297E-12	Positive	202644_s_at
TNFAIP3	7128	Black-107	0,219	2,352E-02	Positive	202644_s_at

Gene name	GeneID	Dataset	R	P	Sign	Probe
TNFAIP3	7128	Booser-508	0,214	1,134E-06	Positive	202644_s_at
TNFAIP3	7128	Bos-204	0,247	3,638E-04	Positive	202644_s_at
TNFAIP3	7128	Chin-124	0,305	5,821E-04	Positive	202644_s_at
TNFAIP3	7128	Clynes-121	0,329	2,306E-04	Positive	202644_s_at
TNFAIP3	7128	EXPO-351	0,157	3,274E-03	Positive	202644_s_at
TNFAIP3	7128	Iglehart-123	0,269	2,583E-03	Positive	202644_s_at
TNFAIP3	7128	Jonsdottir-94	0,212	4,008E-02	Positive	ILMN_1702691
TNFAIP3	7128	Minn-96	0,281	5,520E-03	Positive	202644_s_at
TNFAIP3	7128	Smid-210	0,252	2,286E-04	Positive	202644_s_at
TNFAIP3	7128	Wang-286	0,301	2,172E-07	Positive	202644_s_at
TNFRSF11A	8792	Bertucci-266	0,198	1,152E-03	Positive	238846_at
TNFRSF11A	8792	Booser-508	0,143	1,259E-03	Positive	207037_at
TNFRSF11A	8792	Bos-204	0,167	1,703E-02	Positive	238846_at
TNFRSF11A	8792	Chin-124	0,279	1,721E-03	Positive	207037_at
TNFRSF11A	8792	EXPO-351	0,256	1,225E-06	Positive	238846_at
TNFRSF11A	8792	Wang-286	0,168	4,371E-03	Positive	207037_at
TNFRSF13C	115650	Clynes-121	0,236	9,016E-03	Positive	1552892_at
TNFRSF13C	115650	Jonsdottir-94	0,219	3,367E-02	Positive	ILMN_1731742
TNFRSF13C	115650	Servant-343	0,109	4,326E-02	Positive	ILMN_1731742
TNFRSF1A	7132	Bertucci-266	0,210	5,788E-04	Positive	207643_s_at
TNFRSF1A	7132	Black-107	0,331	5,065E-04	Positive	207643_s_at
TNFRSF1A	7132	Bos-204	0,237	6,534E-04	Positive	207643_s_at
TNFRSF1A	7132	EXPO-351	0,210	7,105E-05	Positive	207643_s_at
TNFRSF1A	7132	Jonsdottir-94	0,251	1,479E-02	Positive	ILMN_1685005
TNFRSF1A	7132	Minn-96	0,259	1,091E-02	Positive	207643_s_at
TNFRSF1A	7132	Smid-210	0,214	1,808E-03	Positive	207643_s_at
TNFRSF1A	7132	Sotiriou-198	0,253	3,277E-04	Positive	207643_s_at
TNFRSF1A	7132	Wang-286	0,238	4,924E-05	Positive	207643_s_at
TNFSF11	8600	Bertucci-266	0,139	2,343E-02	Positive	211153_s_at
TNFSF11	8600	Booser-508	0,180	4,635E-05	Positive	211153_s_at
TNFSF11	8600	Clynes-121	0,233	1,019E-02	Positive	210643_at
TNFSF11	8600	EXPO-351	0,136	1,061E-02	Positive	210643_at
TNFSF11	8600	Jonsdottir-94	0,294	4,060E-03	Positive	ILMN_1673175
TNFSF11	8600	Servant-343	0,156	3,816E-03	Positive	ILMN_1673175
TNFSF11	8600	Sotiriou-120	0,342	1,336E-04	Positive	210643_at
TNFSF11	8600	Sotiriou-198	0,163	2,180E-02	Positive	210643_at
TNFSF13B	10673	Bertucci-266	0,273	6,454E-06	Positive	223502_s_at
TNFSF13B	10673	Black-107	0,242	1,218E-02	Positive	223502_s_at
TNFSF13B	10673	Bos-204	0,258	1,999E-04	Positive	223502_s_at
TNFSF13B	10673	Clynes-121	0,302	7,655E-04	Positive	223502_s_at
TNFSF13B	10673	EXPO-351	0,161	2,508E-03	Positive	223502_s_at
TNFSF13B	10673	Iglehart-123	0,270	2,552E-03	Positive	223502_s_at
TNFSF13B	10673	Smid-210	0,276	4,936E-05	Positive	223502_s_at
TNFSF13B	10673	Sotiriou-120	0,201	2,735E-02	Positive	223502_s_at
TNFSF14	8740	Bertucci-266	0,297	8,297E-07	Positive	207907_at
TNFSF14	8740	Black-107	0,197	4,149E-02	Positive	233935_at
TNFSF14	8740	Booser-508	0,181	4,234E-05	Positive	207907_at
TNFSF14	8740	Bos-204	0,162	2,026E-02	Positive	207907_at
TNFSF14	8740	Chin-124	0,254	4,485E-03	Positive	207907_at
TNFSF14	8740	Servant-343	0,111	4,040E-02	Positive	ILMN_2363392
TNFSF14	8740	Sotiriou-120	0,269	2,987E-03	Positive	207907_at
TRADD	8717	Bertucci-266	0,189	2,009E-03	Positive	1729_at
TRADD	8717	Black-107	0,232	1,598E-02	Positive	1729_at
TRADD	8717	Wang-286	0,197	7,945E-04	Positive	205641_s_at
TRAF1	7185	Bertucci-266	0,236	1,036E-04	Positive	205599_at
TRAF1	7185	Black-107	0,191	4,833E-02	Positive	205599_at
TRAF1	7185	Booser-508	0,276	2,377E-10	Positive	205599_at
TRAF1	7185	Bos-204	0,162	2,043E-02	Positive	205599_at

Gene name	GeneID	Dataset	R	P	Sign	Probe
TRAF1	7185	Chin-124	0,251	4,950E-03	Positive	205599_at
TRAF1	7185	Clynes-121	0,253	5,055E-03	Positive	205599_at
TRAF1	7185	Iglehart-123	0,297	8,427E-04	Positive	205599_at
TRAF1	7185	Jonsdottir-94	0,398	7,245E-05	Positive	ILMN_1698218
TRAF1	7185	Smid-210	0,156	2,406E-02	Positive	205599_at
TRAF1	7185	Wang-286	0,167	4,733E-03	Positive	205599_at
TRAF2	7186	Booser-508	-0,118	7,533E-03	Negative	204413_at
TRAF2	7186	EXPO-351	-0,113	3,361E-02	Negative	204413_at
TRAF2	7186	Servant-343	-0,163	2,416E-03	Negative	ILMN_1691487
TRAF3	7187	Bertucci-266	0,233	1,225E-04	Positive	221571_at
TRAF3	7187	Bos-204	0,190	6,407E-03	Positive	221571_at
TRAF3	7187	Clynes-121	0,229	1,155E-02	Positive	221571_at
TRAF3	7187	Jonsdottir-94	0,209	4,352E-02	Positive	ILMN_2383774
TRAF3	7187	Smid-210	0,187	6,659E-03	Positive	221571_at
TRAF3	7187	Sotiriou-120	0,185	4,255E-02	Positive	221571_at
TRAF5	7188	Bertucci-266	-0,130	3,471E-02	Negative	204352_at
TRAF5	7188	Booser-508	-0,109	1,405E-02	Negative	204352_at
TRAF5	7188	EXPO-351	-0,107	4,539E-02	Negative	204352_at
TRAF5	7188	Minn-96	-0,216	3,492E-02	Negative	204352_at
TRAF5	7188	Smid-210	-0,162	1,913E-02	Negative	204352_at
TRAF5	7188	Sotiriou-120	0,225	1,363E-02	Positive	204352_at
TRAF6	7189	Bertucci-266	0,173	4,555E-03	Positive	205558_at
TRAF6	7189	Booser-508	0,111	1,268E-02	Positive	205558_at
TRAF6	7189	Jonsdottir-94	0,262	1,073E-02	Positive	ILMN_2392143
TRIM25	7706	Minn-96	0,225	2,719E-02	Positive	206911_at
UBE2I	7329	Booser-508	-0,276	2,300E-10	Negative	213535_s_at
UBE2I	7329	Minn-96	-0,290	4,179E-03	Negative	213535_s_at
UBE2I	7329	Sotiriou-198	-0,147	3,877E-02	Negative	213535_s_at
UBE2I	7329	Wang-286	-0,119	4,518E-02	Negative	213535_s_at
VCAM1	7412	Bertucci-266	0,314	1,777E-07	Positive	203868_s_at
VCAM1	7412	Black-107	0,215	2,645E-02	Positive	203868_s_at
VCAM1	7412	Booser-508	0,205	3,104E-06	Positive	203868_s_at
VCAM1	7412	Bos-204	0,208	2,894E-03	Positive	203868_s_at
VCAM1	7412	Chin-124	0,318	3,229E-04	Positive	203868_s_at
VCAM1	7412	Clynes-121	0,419	1,681E-06	Positive	203868_s_at
VCAM1	7412	EXPO-351	0,165	1,971E-03	Positive	203868_s_at
VCAM1	7412	Iglehart-123	0,325	2,446E-04	Positive	203868_s_at
VCAM1	7412	Jonsdottir-94	0,284	5,596E-03	Positive	ILMN_2307903
VCAM1	7412	Minn-96	0,204	4,637E-02	Positive	203868_s_at
VCAM1	7412	Smid-210	0,198	3,934E-03	Positive	203868_s_at
VCAM1	7412	Sotiriou-198	0,215	2,407E-03	Positive	203868_s_at
VCAM1	7412	Wang-286	0,308	1,113E-07	Positive	203868_s_at
XIAP	331	Booser-508	0,173	9,246E-05	Positive	206536_s_at
XIAP	331	Servant-343	0,109	4,376E-02	Positive	ILMN_1815361
XIAP	331	Wang-286	-0,178	2,508E-03	Negative	206536_s_at
ZAP70	7535	Bertucci-266	0,252	3,242E-05	Positive	214032_at
ZAP70	7535	Booser-508	0,157	3,898E-04	Positive	214032_at
ZAP70	7535	Bos-204	0,151	3,093E-02	Positive	1555613_a_at
ZAP70	7535	Chin-124	0,289	1,128E-03	Positive	214032_at
ZAP70	7535	Iglehart-123	0,213	1,826E-02	Positive	1555613_a_at
ZAP70	7535	Jonsdottir-94	0,254	1,360E-02	Positive	ILMN_1719756
ZAP70	7535	Smid-210	0,164	1,739E-02	Positive	1555613_a_at
ZAP70	7535	Wang-286	0,169	4,245E-03	Positive	214032_at



Primer	Sequence
1	GGGGACAACCTTTGTACAAAAAAGTTGGCATGAACTCCTTTCTGG
2	GGGGACAACCTTTGTACAAAAAAGTTGGCATGTGCGGCGGCGACGACCGC
3	GCGGCAGCTGCCAAGGAGGGGCTCCTG
4	GCGGCAGCTGCCACCTTTGACTGGATG
5	GGGGACAACCTTTGTACAAGAAAGTTGGGTAGTGGGAGGTAGTCAGAGTGTC
6	GGCAGCTGCCGCTTGACCCACGTAGCC
7	GGCAGCTGCCGCGTGGTGATGGTG
8	GGCAGCTGCCGCGGGCGAGCTGATCTGCACCCC
9	GCGGCAGCTGCCCCCCAGACGGCTACTTACCAG
10	TTAACTGCAGTCACTCACGCTTCTTCTG

HOXA1 mutants	First PCR		Second PCR	
	Primer F	Primer R	Primer F	Primer R
HOXA1 <sup>ΔNter</sup>	2	5	x	x
HOXA1 <sup>ΔCenter</sup>	1	7	4	5
HOXA1 <sup>Δ11His</sup>	1	8	9	5
HOXA1 <sup>ΔHD</sup>	1	6	3	5
HOXA1 <sup>ΔCter</sup>	1	10	x	x

# Supplemental Figure 2

Taminiau *et al.*

