

**bc-GenExMiner 3.0: new mining module computes breast cancer gene-expression correlation analyses**

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Supplementary Method:

**Multi-correlation score (MCS) calculation (eigenvalue decomposition)**

Given a matrix M, an eigenvalue is a scalar  $\lambda$  such that  $Mu=\lambda u$  for some non-zero vector v, which is called an eigenvector. As the correlation matrix C of a given set of genes is diagonalizable, we can find a matrix U of linearly independent eigenvectors such that  $C=U\Lambda U^{-1}$ , where U is the matrix of eigenvectors and  $\Lambda$  is a diagonal matrix whose diagonal elements are the different eigenvalues. Each eigenvalue is equal to the variance explained by the corresponding eigenvector. As the eigenvectors are linearly independent, taking the largest eigenvalue permits to quantify the variance that is the most important within the data, and when divided by the sum of all eigenvalues and multiply by 100 to get the multi-correlation score, to know the proportion of this “information” compared to the whole “information” contained in the data.

**Supplementary Table 1** Screening of continuous clusters of correlated ( $r \geq 0.30$  and  $p \leq 0.05$ ) co-expressed genes ( $n \geq 3$ ) based on Buiness results as input genes by means of bc-GenExMiner v3.0 gene expression correlation analysis by chromosomal location [15]

N <sup>a</sup>	Cytoband	Molecular subtype	bc-GenExMiner continuous cluster of correlated co-expressed genes		MCS <sup>b</sup> (%)
			No.	genes	
1	17q12-q21	All patients	7	<i>STARD3, TCAP, PNMT, PGAP3, ERBB2, C17orf37, GRB7</i>	59.84
		All patients	3	<i>FBXL20, MED1, CDK12</i>	78.06
		HER2			83.53
		luminal A			60.34
		luminal B			68.43
		All patients	7	<i>MED24, THRA, NR1D1, MSL1, CASC3, RAPGEFL1, WIPF2</i>	54.33
		HER2			59.97
		HER2	8	<i>MED24, THRA, NR1D1, MSL1, CASC3, RAPGEFL1, WIPF2, CDC6</i>	58.05
		All patients	7	<i>SOC5T, ARHGAP23, SRCIN1, C17orf96, MLLT6, CISD3, PCGF2</i>	58.11
		HER2	5	<i>SRCIN1, C17orf96, MLLT6, CISD3, PCGF2</i>	76.67
		Luminal B	3	<i>LOC100287808, PSMB3, PIP4K2B</i>	71.99
		All patients	3	<i>PSMB3, PIP4K2B, CWC25</i>	70.52
		HER2			90.65
		Luminal B	3	<i>SYNRG, DDX52, HNF1B</i>	60.89
		All patients	4	<i>TADA2A, DUSP14, SYNRG, DDX52</i>	56.36
		HER2			68.94
		Luminal B	3	<i>ARL4D, DHX8, RPL19</i>	62.50
		All patients	3	<i>GSDMB, ORMDL3, PSMD3</i>	71.50
		HER2			68.85
2	17q11-q12	All patients	6	<i>TRAF4, C17orf63, ERA1, FLOT2, DHRS13, PHF12</i>	57.08
		HER2			80.20
		Basal-like	4	<i>TRAF4, C17orf63, ERA1, FLOT2</i>	53.87
		Luminal B			70.20
		Luminal A	3	<i>ERA1, FLOT2, DHRS13</i>	61.04
		All patients	5	<i>SPAG5, SGK494, KIAA0100, SDF2, SUPT6H</i>	58.85
		Basal-like	3	<i>KIAA0100, SDF2, SUPT6H</i>	69.66
		HER2			90.45
		Luminal B			68.79
		All patients	5	<i>TMEM97, IFT20, TNFAIP1, POLDIP2, TMEM199</i>	61.17
		Basal-like			55.78
		HER2	4	<i>TMEM97, IFT20, TNFAIP1, POLDIP2</i>	81.07
		Luminal B	4	<i>IFT20, TNFAIP1, POLDIP2, TMEM199</i>	66.58
		Basal-like, HER2	3	<i>BLMH, CPD, GOSR1</i>	56.55
		All patients	3	<i>TMEM199, SARMI, SLC46A1</i>	68.42
					58.26
3	8q24	All patients	7	<i>FBXL6, GPR172A, ADCK5, CPSF1, SLC39A4, VPS28, CYHRI</i>	58.77
		basal-like			64.15
		HER2	4	<i>FBXL6, GPR172A, ADCK5, CPSF1</i>	70.30
		Luminal A			50.46
		Luminal B			72.15
		HER2	4	<i>CPSF1, SLC39A4, VPS28, CYHRI</i>	61.90
		All patients	7	<i>GRINA, OPLAH, EXOSC4, GPA1, CYC1, SHARPIN, MAF1</i>	59.64
		Luminal B			78.04
		Basal-like	6	<i>GRINA, OPLAH, EXOSC4, GPA1, CYC1, SHARPIN</i>	61.34
		HER2	5	<i>EXOSC4, GPA1, CYC1, SHARPIN, MAF1</i>	68.28
		Luminal A	4	<i>EXOSC4, GPA1, CYC1, SHARPIN</i>	59.72
		All patients	5	<i>ZNF707, MAPK15, FAM83H, SCRIB, PUF60</i>	59.48
		Basal-like	4	<i>FAM83H, SCRIB, PUF60, NRBP2</i>	65.45
		HER2	3	<i>FAM83H, SCRIB, PUF60</i>	77.83
		Luminal A			65.77
		Luminal B	5	<i>FAM83H, SCRIB, PUF60, NRBP2, PLEC</i>	68.82
		All patients	5	<i>MFSD3, RECOL4, LRRC14, C8orf82, ARHGAP39</i>	57.25
		Basal-like	3	<i>RECOL4, LRRC14, C8orf82</i>	60.14
		Luminal B	4	<i>RECOL4, LRRC14, C8orf82, ARHGAP39</i>	68.71
		All patients	10	<i>C8orf51, RHPN1, ZC3H3, GSDMD, C8orf73, NAPRTI, EEF1D, TIGD5, PYCRL, TSTA3</i>	49.02
		Basal-like	4	<i>EEF1D, TIGD5, PYCRL, TSTA3</i>	55.80
		Luminal B	4	<i>TIGD5, PYCRL, TSTA3, ZNF623</i>	64.65
		All patients	7	<i>ZNF517, ZNF7, COMM5, ZNF250, ZNF16, ZNF252, C8orf33</i>	56.16
		Basal-like	6	<i>ZNF7, COMM5, ZNF250, ZNF16, ZNF252, C8orf33</i>	63.65
		HER2	4	<i>ZNF7, COMM5, ZNF250, ZNF16</i>	73.41
		Luminal B			79.67
		Luminal A	3	<i>COMM5, ZNF250, ZNF16</i>	56.09

		All patients	10	<i>MAPK15, FAM83H, SCRIB, PUF60, NRBP2, PLEC, PARP10, GRINA, OPLAH, EXOSC4</i>	54.45
		Basal-like	6	<i>GRINA, OPLAH, EXOSC4, GPAA1, CYC1, SHARPIN</i>	61.34
		HER2	5	<i>EXOSC4, GPAA1, CYC1, SHARPIN, MAF1</i>	68.28
		Luminal A	4	<i>EXOSC4, GPAA1, CYC1, SHARPIN</i>	59.72
		Luminal B	7	<i>GRINA, OPLAH, EXOSC4, GPAA1, CYC1, SHARPIN, MAF1</i>	78.04
4	6p21	All patients	4	<i>HLA-DOB, TAP2, PSMB8, TAPI</i>	59.93
		Basal-like	9	<i>HLA-DOB, TAP2, PSMB8, TAPI, HLA-DRB1, PSMB9, HLA-DQB1, HLA-DMB, HLA-DMA</i>	68.48
		All patients	7	<i>PSMB8, TAPI, HLA-DRB1, PSMB9, HLA-DQB1, HLA-DMB, HLA-DMA</i>	65.73
		HER2			66.42
		Luminal B	5	<i>PSMB8, TAPI, HLA-DRB1, PSMB9, HLA-DQB1</i>	65.77
		Basal-like	4	<i>HLA-DRA, HLA-DOA1, HLA-DRB6, HLA-DOB</i>	60.39
		All patients	4	<i>HLA-DQB2, HLA-DPA1, HLA-DPB1, HLA-DPB2</i>	60.91
		Basal-like	3	<i>HLA-DQB2, HLA-DPA1, HLA-DPB1</i>	63.85
		HER2			68.96
		Luminal B	4	<i>HLA-DOA, HLA-DQB2, HLA-DPA1, HLA-DPB1</i>	61.76
		All patients	3	<i>HLA-DOA, HLA-DQB2, HLA-DPA1</i>	59.30
		All patients	3	<i>HSD17B8, RING1, VPS52</i>	58.70
		All patients	4	<i>PRICKLE4, USP49, MED20, BYSL</i>	52.47
5	17q24-q25	All patients	7	<i>SUMO2, NUP85, GGA3, MRPS7, MIF4GD, SLC25A19, GRB2</i>	53.47
		Basal-like	3	<i>NUP85, GGA3, MRPS7</i>	61.04
		HER2	12	<i>CDR2L, ICT1, ATP5H, KCTD2, SLC16A5, ARMC7, NT5C, HNI, SUMO2, NUP85, GGA3, MRPS7</i>	74.89
		Luminal B	3	<i>NUP85, GGA3, MRPS7</i>	68.95
		HER2	4	<i>SAP30BP, ITGB4, GALK1, H3F3B</i>	67.43
		All patients	4	<i>ARL16, HGS, MRPL12, SLC25A10</i>	65.17
		Basal-like			62.98
		HER2	3	<i>HGS, MRPL12, SLC25A10</i>	73.10
		Luminal A			58.43
		Luminal B			69.81
		All patients	5	<i>ARMC7, NT5C, HNI, SUMO2, NUP85</i>	53.89
		Basal-like	3	<i>HNI, SUMO2, NUP85</i>	71.29
		HER2	10	<i>C17orf28, CDR2L, ICT1, ATP5H, KCTD2, SLC16A5, ARMC7, NT5C, HNI, SUMO2</i>	76.06
		HER2	4	<i>SAP30BP, ITGB4, GALK1, H3F3B</i>	67.43
		All patients	3	<i>SIRT7, PCYT2, MAFG</i>	63.88
		HER2			66.05
		Basal-like	4	<i>SIRT7, PCYT2, MAFG, PYCR1</i>	55.43
		All patients	8	<i>ASPScri, STRA13, LRRK45, RAC3, DCXR, RFNG, GPS1, DUSIL</i>	55.40
		Basal-like	5	<i>ASPScri, STRA13, LRRK45, RAC3, DCXR</i>	57.41
		HER2			71.89
		All patients	4	<i>KIAA0195, CASKIN2, TSEN54, LLGL2</i>	58.62
		HER2	6	<i>SLC25A19, GRB2, KIAA0195, CASKIN2, TSEN54, LLGL2</i>	69.04
		Luminal B	4	<i>SLC25A19, GRB2, KIAA0195, CASKIN2</i>	58.89
		Basal-like	3	<i>CASKIN2, TSEN54, LLGL2</i>	56.87
		HER2	3	<i>RECQL5, SAP30BP, ITGB4</i>	67.00
		Basal-like	4	<i>LLGL2, MYO15B, RECQL5, SAP30BP</i>	36.39
		All patients	4	<i>ARL16, HGS, MRPL12, SLC25A10</i>	65.17
		Basal-like			62.98
		All patients	4	<i>P4HB, ARHGDI, ANAPC11, THOC4</i>	58.47
		Basal-like			67.44
		HER2			66.34
		Luminal B			67.01
6	11q12-q13	All patients	3	<i>MRPL49, SYVNI, CAPN1</i>	61.39
		All patients	3	<i>DPF2, TIGD3, SLC25A45</i>	57.45
		HER2	3	<i>POLA2, CDC42EP2, DPF2</i>	61.86
		All patients	5	<i>MAP3K11, PCNXL3, SIPAI, RELA, KAT5</i>	49.49
		Luminal B	3	<i>SCYLI, LTBP3, SSSCA1</i>	65.63
		HER2	3	<i>PACSI, EEF1G, TUT1</i>	56.61
		All patients	5	<i>KLC2, RAB1B, CNIH2, YIF1A, TMEM151A</i>	51.89
7	1q21-q22	All patients	3	<i>ADAM15, EFNA4, EFNA3</i>	55.87
		HER2			57.33
		Basal-like	3	<i>DCST2, DCST1, ADAM15</i>	77.24
		HER2	4	<i>PYGO2, SHC1, CKS1B, FLAD1</i>	62.77
		Luminal B	3	<i>SHC1, CKS1B, FLAD1</i>	56.03
		All patients	4	<i>FAM189B, SCAMP3, CLK2, HCN3</i>	53.88
		Basal-like	3	<i>FAM189B, SCAMP3, CLK2</i>	61.79
		HER2			70.37
		Luminal B			57.88
		Luminal A	4	<i>GBA, FAM189B, SCAMP3, CLK2</i>	59.26
		All patients	4	<i>MSTO1, YY1API, DAP3, GON4L</i>	56.76

		Luminal A		60.28
		Basal-like	3	MSTO1, YY1API, DAP3
		HER2	3	YY1API, DAP3, GON4L
		HER2	3	EFNA1, DPM3, KRTCAP2
		Luminal B	3	EFNA4, EFNA3, EFNA1
		All patients	3	FDPS, Clorf104, RUSC1
		All patients	4	PMVK, PBXIP1, PYGO2, SHC1
		HER2		
		All patients	4	PSMD4, ZNF687, PI4KB, RFX5
		Basal-like	3	PSMD4, ZNF687, PI4KB
		Luminal A		
		HER2	4	VPS72, PIP5K1A, PSMD4, ZNF687
		All patients	4	POGZ, CGN, TUFT1, SNX27
		HER2	3	POGZ, CGN, TUFT1
		Luminal A		
8	17q21-q23	All patients	6	DHX40, CLTC, PTRH2, TMEM49, TUBD1, RPS6KB1
		Basal-like	3	CLTC, PTRH2, TMEM49
		HER2	8	DHX40, CLTC, PTRH2, TMEM49, TUBD1, RPS6KB1, RNFT1, HEATR6
		Luminal B	3	DHX40, CLTC, PTRH2
		HER2	7	CLTC, PTRH2, TMEM49, TUBD1, RPS6KB1, RNFT1, HEATR6
		Luminal B	4	TUBD1, RPS6KB1, RNFT1, HEATR6
		All patients	6	DHX40, CLTC, PTRH2, TMEM49, TUBD1, RPS6KB1
		HER2	3	ACSF2, CHAD, RSAD1
		HER2	4	TOB1, SPAG9, NME1, NME1-NME2
9	10p11-p12	All patients	5	UPF2, DHTKD1, SEC61A2, NUDT5, CDC123
		Luminal B	4	DHTKD1, SEC61A2, NUDT5, CDC123
		Basal-like	3	RPP38, NMT2, FAMI171A1
		All patients	3	SEPHS1, BEND7, PRPF18
		Basal-like		
		All patients		UPF2, DHTKD1, SEC61A2, NUDT5, CDC123
		Basal-like	3	UPF2, DHTKD1, SEC61A2
		Basal-like	3	HSPA14, SUV39H2, DCLRE1C
		Luminal B		
		HER2	3	FAM188A, PTER, RSU1
10	19q13	All patients	7	ZNF579, FIZ1, ZNF524, ZNF784, ZNF580, ZNF581, CCDC106
		Basal-like	3	TMEM190, RPL28, UBE2S
		All patients	5	UBE2S, SHISA7, ISOC2, ZNF628, NAT14
		All patients	5	ZNF416, ZIK1, ZNF530, ZNF134, ZNF211
		Luminal B	3	ZNF530, ZNF134, ZNF211
		All patients	3	ZNF135, ZSCAN18, ZNF329
		Basal-like		
		HER2		
		Luminal A		
		Luminal B		
		All patients	4	ZSCAN18, ZNF329, ZNF274, ZNF544
		Basal-like	3	ZSCAN18, ZNF329, ZNF274
		Luminal B	3	ZNF132, KIR2DS3, KIR2DS2
		Basal-like	3	ZBTB45, TRIM28, CHMP2A
		Luminal B		
11	16p13	All patients	3	SNHG9, NTHL1, TSC2
		All patients	3	SLC9A3R2, SNHG9, NTHL1
		All patients	3	NME4, DECR2, RAB11FIP3
		Luminal A		
		Basal-like	3	DECR2, RAB11FIP3, SOLH
		All patients	4	RAB11FIP3, SOLH, PIGO, RAB40C
		Luminal A	3	SOLH, PIGO, RAB40C
		All patients	8	FAM195A, WDR90, RHOT2, RHBTL1, STUB1, JMJD8, WDR24, FBXL16
		Basal-like	3	FAM195A, WDR90, RHOT2
		Luminal A		
		Luminal B	3	RHOT2, RHBTL1, STUB1
		All patients	3	CCDC154, CLCN7, TELO2
		All patients	3	SNHG9, NTHL1, TSC2
12	8p11-q13	All patients	5	LSM1, BAG4, DDHD2, PPAPDC1B, WHSC1L1
		Basal-like		
		HER2		
		Luminal B		
		All patients	3	FGFR1, TACCI, PLEKHA2
		All patients	3	GTF2E2, GSR, UBXN8
		HER2	4	GTF2E2, GSR, UBXN8, PPP2CB
		Luminal B	7	LEPROTL1, DCTN6, RBPM3, GTF2E2, GSR, UBXN8, PPP2CB
13	19p13	All patients	5	GCDH, SYCE2, FARSA, CALR, RAD23A
		Basal-like	4	FARSA, CALR, RAD23A, GADD45GIP1

		Luminal B		57.17
		Luminal A	3	62.97
		All patients	4	56.36
		All patients	5	51.63
		Basal-like	4	59.77
		HER2	3	55.65
		Luminal B		64.19
		All patients	4	59.03
		Basal-like	3	71.77
14	1q42-q44	All patients	4	54.39
		Basal-like		59.05
		All patients	3	67.48
		Basal-like		62.60
		HER2	3	60.64
		Luminal B		70.81
		All patients	4	58.40
		Basal-like		57.22
		All patients	3	57.49
		Luminal A		62.57
		All patients	6	55.14
		Basal-like	4	68.04
		Luminal B		71.47
		Luminal A	5	57.94
15	16q11-q13	All patients	6	62.75
		Luminal A		76.39
		Basal-like	8	70.68
		HER2	3	77.14
		Luminal B	4	71.87
		All patients	3	64.71
		Luminal A	3	64.44
		Luminal B		75.91
		All patients	4	53.95
		Luminal B	3	76.68
16	16q21-q24	All patients	5	49.87
		All patients	4	58.44
		Basal-like	3	61.58
		Luminal A		63.03
		Luminal B		65.08
		All patients	3	57.13
		All patients	7	51.69
		Basal-like	3	57.63
		Luminal B		64.69
		All patients	3	57.94
17	6p21-p22	All patients	4	63.22
		Basal-like		74.94
		HER2		65.95
		Luminal A		55.79
		Luminal A	3	62.75
		All patients	6	55.39
		Basal-like	3	72.44
		HER2	3	68.75
		All patients	3	67.40
		Basal-like		73.73
		Luminal A		74.45
		Luminal B		74.40
		All patients	5	65.05
		HER2		67.52
		Basal-like	4	64.89
		HER2	3	68.63
		All patients	4	64.34
18	11q13-q14	Basal-like		60.20
		HER2		69.29
		Luminal A		60.81
		All patients	4	65.37
		Luminal B	3	85.43
		All patients	5	64.33
		Luminal B		74.46
		HER2	3	84.24
		Luminal A		62.95
		All patients	3	67.11
		Luminal A		59.57
		Luminal B		81.35

		Luminal A	3	<i>PRCP, C11orf82, RAB30</i>	47.99
		All patients	3	<i>OR7E126P, MAP6, MOGAT2</i>	60.79
19	1p32-p35	All patients	3	<i>IPO13, ATP6V0B, B4GALT2</i>	58.45
		Basal-like	4	<i>DPH2, IPO13, ATP6V0B, B4GALT2</i>	53.90
		Basal-like	3	<i>EIF2B3, HECTD3, UROD</i>	56.72
		HER2	3	<i>PRDX1, AKR1A1, NASP</i>	61.19
		All patients	3	<i>DMAPI, ERI3, RNF220</i>	59.11
		Basal-like			61.91
		Luminal B			62.15
20	19q13	All patients	3	<i>INPP5B, LOC100128093, SF3A3</i>	59.11
		All patients	4	<i>SCAF1, IRF3, BCL2L12, PRMT1</i>	57.03
		Basal-like	3	<i>IRF3, BCL2L12, PRMT1</i>	66.60
		Luminal B			77.90
		All patients	3	<i>NOSIP, PRRG2, PRR12</i>	58.27
		Luminal B			65.38
		All patients	5	<i>MED25, DKFZP434J0226, PTOV1, PNKP, AKT1S1</i>	54.13
		Luminal A	3	<i>DKFZp434J0226, PTOV1, PNKP</i>	61.32
		Luminal B	3	<i>PTOV1, PNKP, AKT1S1</i>	70.23
21	19q13	Luminal B	3	<i>VRK3, ZNF473, MYH14</i>	71.39
		All patients	3	<i>ZNF383, HKR1, ZNF527</i>	61.05
		All patients	3	<i>WDR87, SIPA1L3, DPF1</i>	59.42
		All patients	3	<i>PSMD8, GGN, FAM98C</i>	58.38
22	3p21	HER2	3	<i>ACTN4, CAPN12, LGALS7</i>	68.01
		Luminal B	3	<i>GPX1, RHOA, TCTA</i>	55.52
		All patients	7	<i>P4HTM, WDR6, DALRD3, NDUFAF3, IMPDH2, QRICH1, QARS</i>	50.05
		HER2	3	<i>IMPDH2, QRICH1, QARS</i>	62.24
23	16p11-p13	All patients	3	<i>QARS, USP19, LAMB2</i>	58.44
		All patient	3	<i>BCKDK, MYST1, PRSS8</i>	63.11
		Luminal A			72.57
		Luminal B			68.34
		Luminal A	3	<i>FUS, PYCARD, PYDC1</i>	65.44
		All patients	3	<i>FBXL19, ORAI3, SETD1A</i>	65.50
		All patients	7	<i>PRRI4, FBRS, SRCAP, PHKG2, C16orf93, RNF40, ZNF629</i>	54.88
		Luminal A	4	<i>PRRI4, FBRS, SRCAP, PHKG2</i>	63.22
		Luminal A	4	<i>PHKG2, C16orf93, RNF40, ZNF629</i>	61.39
		Luminal B	3	<i>PRRI4, FBRS, SRCAP</i>	65.51
		All patients	5	<i>BOLA2, SULT1A3, LOC595101, CD2BP2, TBC1D10B</i>	51.47
		All patients	5	<i>ZNF768, ZNF747, ZNF764, ZNF688, ZNF785</i>	55.52
24	14q13-q21	Luminal B	3	<i>KLHD2, SDCCAG1, ARF6</i>	60.88
		All patients	3	<i>SOS2, L2HGDH, ATP5S</i>	54.12
		All patients	3	<i>RPL36AL, MGAT2, C14orf104</i>	57.30
		HER2			56.13
		Basal-like	3	<i>PPIL5, RPL36AL, MGAT2</i>	61.89
		All patients	3	<i>MAP4K5, ATL1, SAV1</i>	60.16
		All patient	3	<i>FAM179B, PRPF39, FKBP3</i>	60.88
		Basal-like			59.81
		Luminal B			65.65
25	12q13-q15	Luminal B	3	<i>KLHL28, FAM179B, PRPF39</i>	72.22
		HER2	3	<i>YEATS4, FRS2, CCT2</i>	66.61
		Luminal B			73.98
		All patients	4	<i>MDM1, RAP1B, NUP107, SLC35E3</i>	57.47
		HER2			75.43
		Luminal A	3	<i>MDM1, RAP1B, NUP107</i>	58.61
		All patients	3	<i>NUP107, SLC35E3, MDM2</i>	57.57
26	6q16-q21	HER2	4	<i>RAP1B, NUP107, SLC35E3, MDM2</i>	66.94
		Luminal B			75.13
		All patients	7	<i>ATG5, AIM1, RTN4IP1, QRSL1, C6orf203, BEND3, PDSS2</i>	51.52
		Basal-like	4	<i>ATG5, AIM1, RTN4IP1, QRSL1</i>	59.51
		HER2	4	<i>AIM1, RTN4IP1, QRSL1, C6orf203</i>	69.08
		Luminal B	3	<i>ATG5, AIM1, RTN4IP1</i>	69.82
		Luminal A	3	<i>RTN4IP1, QRSL1, C6orf203</i>	57.74
27	8q21-23	HER2	3	<i>SMPD2, MICAL1, ZBTB24</i>	62.37
		Luminal B			71.04
		Luminal B	3	<i>WASF1, CDC40, DDO</i>	57.97
		All patients	6	<i>RAD54B, KIAA1429, ESRP1, DPY19L4, INTS8, CCNE2</i>	57.87
		Basal-like	3	<i>RAD54B, KIAA1429, ESRP1</i>	65.96
		Basal-like	4	<i>KIAA1429, ESRP1, DPY19L4, INTS8</i>	55.24
		Luminal B			72.40
28	12q13-q15	HER2	5	<i>RAD54B, KIAA1429, ESRP1, DPY19L4, INTS8</i>	68.10
		HER2	7	<i>ESRP1, DPY19L4, INTS8, CCNE2, TP53INP1, C8orf38, PLEKHF2</i>	61.95
		All patients	3	<i>UQCRCB, MTERFD1, PTDSSI</i>	67.67
		Basal-like			69.88

		HER2			72.53
		Luminal A			56.98
		Luminal B			74.31
		Luminal A	3	<i>SDC2, PGCP, TSPYL5</i>	60.86
		Luminal B	4	<i>MTERFD1, PTDSSI, SDC2, PGCP</i>	64.49
		Basal-like	3	<i>TSPYL5, MTDH, LAPTM4B</i>	59.82
		Luminal B	3	<i>MTDH, LAPTM4B, MATN2</i>	60.76
		All patients	3	<i>FBXO43, POLR2K, SPAG1</i>	56.99
		HER2	4	<i>POLR2K, SPAG1, RNF19A, ANKRD46</i>	63.19
28	3q26-q27	All patients	3	<i>DVL3, AP2M1, ABCF3</i>	54.60
		Basal-like	4	<i>EIF2B5, DVL3, AP2M1, ABCF3</i>	52.75
		Basal-like	3	<i>PSMD2, EIF4G1, FAM131A</i>	63.68
		Luminal B			62.97
29	11p11-p13	Basal-like	3	<i>APIP, PDHX, CD44</i>	66.50
		Luminal B	3	<i>FJXI, TRIM44, LDLRAD3</i>	69.21
		Luminal B	3	<i>CAPRINI, NAT10, ABTB2</i>	62.31
		Basal-like	3	<i>TRIM44, LDLRAD3, COMMD9</i>	63.38
		Luminal B			65.13
30	19p11-p13	All patients	5	<i>LASS1, COPE, DDX49, HOMER3, ARMC6</i>	51.35
		Basal-like	4	<i>COPE, DDX49, HOMER3, ARMC6</i>	59.16
		All patients	3	<i>TMEM161A, RFXANK, NR2C2AP</i>	67.26
		Basal-like			77.37
		Luminal A			56.15
		All patients	3	<i>C19orf50, UBA52, C19orf60</i>	59.75
31	20q13	All patients	4	<i>OSBPPL2, ADRM1, LAMA5, LOC100287547</i>	53.01
		HER2	3	<i>OSBPPL2, ADRM1, LAMA5</i>	57.59
		Luminal B	3	<i>SYCP2, PPP1R3D, C20orf177</i>	68.54
		All patients	3	<i>TCFL5, DIDO1, C20orf11</i>	61.26
		Luminal B			62.65
32	9q33-q34	Luminal B	3	<i>GAPVD1, MAPKAP1, PBX3</i>	56.09
		HER2	3	<i>LMX1B, ZBTB43, ZBTB34</i>	82.42
		All patients	3	<i>TTC16, TOR2A, SH2D3C</i>	56.53

a: Buness order of frequency; b: multi-correlation score

**Supplementary Table 2** Screening of continuous clusters of correlated ( $r \geq 0.30$  and  $p \leq 0.05$ ) co-expressed genes ( $n \geq 2$ ) according to ER status of genes overexpressed in gained regions based on Han's results as input genes by means of bc-GenExMiner v3.0 gene expression correlation analysis by chromosomal location [21]

Cytoband	Input gene	Test results			ER- specificity
		ER <sup>a</sup> - cluster	ER+ cluster	Basal-like cluster	
9p24.3-p21.1	<i>KANK1</i>	<i>KANK1, DMRT1, DMRT3, DMRT2</i> <b>MCS<sup>b</sup> = 46.7</b>	<i>DOCK8, KANK1, DMRT1, DMRT3, DMRT2</i> <b>MCS = 52.5</b>	No cluster	Discordant
	<i>KIAA0020</i>	Discontinuous		Discontinuous	Tendency
	<i>RCL1</i>	<i>PPAPDC2, CDC37L1, AK3, RCL1, JAK2</i> <b>MCS = 55.3</b>	<i>PPAPDC2, CDC37L1, AK3, RCL1</i> <b>MCS = 54</b>	<i>PPAPDC2, CDC37L1, AK3, RCL1</i>	Yes
	<i>UHRF2</i>	No cluster	No cluster	Discontinuous	No
	<i>GLDC</i>	No cluster	No cluster	No cluster	No
	<i>KDM4C</i>	No cluster	No cluster	Discontinuous	No
	<i>C9orf123</i>	Discontinuous		Discontinuous	Tendency
	<i>NFIB</i>	<i>MPDZ, NFIB, ZDHHC21</i> <b>MCS = 58.8</b>	<i>MPDZ, NFIB</i> <b>MCS = 57.8</b>	<i>MPDZ, NFIB, ZDHHC21</i>	Yes
	<i>PSIP1</i>	<i>TTC39B, SNAPC3, PSIP1, C9orf93</i> <b>MCS = 60.4</b>	<i>TTC39B, SNAPC3, PSIP1, C9orf93</i> <b>MCS = 61</b>	<i>TTC39B, SNAPC3, PSIP1</i>	Discordant
	<i>PLIN2</i>	No cluster	No cluster	No cluster	No
	<i>RPS6</i>	No cluster	No cluster	No cluster	No
	<i>MTAP</i>	No cluster	No cluster	No cluster	No
	<i>PLAA</i>	<i>TUSCI, C9orf82, PLAA</i> <b>MCS = 66.7</b>	<i>TUSCI, C9orf82, PLAA</i> <b>MCS = 62.7</b>	<i>TUSCI, C9orf82, PLAA</i>	Yes
	<i>MOBKL2B</i>	<i>C9orf11, MOBKL2B, IFNK</i> <b>MCS = 63.1</b>	<i>C9orf11, MOBKL2B</i> <b>MCS = 56.8</b>	No cluster	Yes
10p15.3-p14	<i>CNTLN</i>	No cluster	No cluster	No cluster	No
	<i>HAUS6</i>	Discontinuous		Discontinuous	Tendency
	<i>CDKN2A</i>	<i>CDKN2A, CDKN2B</i> <b>MCS = 52.4</b>	<i>CDKN2A, CDKN2B, DMRTA1</i> <b>MCS = 54.1</b>	<i>CDKN2A, CDKN2B</i>	Discordant
	<i>GTPBP4</i>	<i>LARP4B, GTPBP4, IDI2</i> <b>MCS = 52</b>	<i>LARP4B, GTPBP4, IDI2, IDII</i> <b>MCS = 54.7</b>	<i>LARP4B, GTPBP4, IDI2</i>	Discordant
	<i>PFKP</i>	Discontinuous			Tendency
	<i>IL15RA</i>	<i>IL15RA, IL2RA</i> <b>MCS = 49.7</b>	<i>ANKRD16, FBXO18, IL15RA, IL2RA</i> <b>MCS = 50.2</b>	<i>IL15RA, IL2RA</i>	Discordant
	<i>RBM17</i>	No cluster	No cluster	No cluster	No
	<i>CELF2</i>	No cluster <b>MCS = 64</b>	<i>SFTA1P, LOC254312, CELF2</i> <b>MCS = 59</b>	No cluster	Yes
	<i>NUDT5</i>	<i>UPF2, DHTKD1, SEC61A2, NUDT5, CDC123</i> <b>MCS = 55.6</b>	<i>UPF2, DHTKD1, SEC61A2, NUDT5, CDC123</i> <b>MCS = 53.6</b>	Discontinuous	Yes
	<i>IDII</i>	No cluster <b>MCS = 52</b>	<i>LARP4B, GTPBP4, IDI2, IDII</i> <b>MCS = 54.7</b>	No cluster	Discordant
12p13.33-p13.31	<i>WDR37</i>	No cluster	No cluster	No cluster	No
	<i>CALML5</i>	No cluster	No cluster	No cluster	No
	<i>GDI2</i>	<i>C10orf18, GDI2</i> <b>MCS = 67.7</b>	No cluster <b>MCS = 61.5</b>	<i>C10orf18, GDI2, FBXO18</i>	Yes
	<i>KIN</i>	<i>KIN, ATP5C1, TAF3</i> <b>MCS = 64.9</b>	<i>KIN, ATP5C1</i> <b>MCS = 55.1</b>	<i>KIN, ATP5C1</i>	Yes
	<i>USP6NL</i>	<i>USP6NL, ECHDC3, C10orf47</i> <b>MCS = 61</b>	No cluster <b>MCS = 55.9</b>	<i>USP6NL, ECHDC3</i>	Yes
	<i>CCDC77</i>	<i>KDM5A, CCDC77</i> <b>MCS = 83.6</b>	<i>KDM5A, CCDC77</i> <b>MCS = 73.3</b>	<i>KDM5A, CCDC77, B4GALNT3</i>	Yes
	<i>NINJ2</i>	<i>NINJ2, LOC100049716</i> <b>MCS = 71.3</b>	No cluster <b>MCS = 63.8</b>	<i>NINJ2, LOC100049716, WNK1</i>	Yes
	<i>WNT5B</i>	No cluster	No cluster	<i>FBXL14, WNT5B</i>	No
13q31.3-q34	<i>FOXMI</i>	<i>FOXM1, C12orf32, TULP3, TEAD4</i> <b>MCS = 61.7</b>	<i>FOXM1, C12orf32</i> <b>MCS = 54.4</b>	<i>FOXM1, C12orf32, TULP3, TEAD4</i>	Yes
	<i>C12orf4</i>	<i>C12orf4, RAD51API, DYRK4</i> <b>MCS = 62.9</b>	<i>C12orf4, RAD51API</i> <b>MCS = 54.3</b>	<i>C12orf4, RAD51API, DYRK4</i>	Yes
	<i>NDUFA9</i>	No cluster	No cluster	No cluster	No
	<i>DZIP1</i>	No cluster	No cluster	No cluster	No
	<i>RAP2A</i>	<i>MBNL2, RAP2A, IPO5</i> <b>MCS = 61.3</b>	<i>RAP2A, IPO5</i> <b>MCS = 56.6</b>	<i>MBNL2, RAP2A, IPO5, FARPI, STK24</i>	Yes
	<i>TNFSF13B</i>	<i>TNFSF13B, SP3P</i>	No cluster	No cluster	Yes

		<b>MCS = 64.8</b>		
18q12.1-12.2	<i>ANKRD10</i>	<i>CARKD, CARS2, ING1, ANKRD10, ARHGEF7, C13orf16</i> <b>MCS = 50.6</b>	<i>CARKD, CARS2, ING1, ANKRD10, ARHGEF7, C13orf16</i> <b>MCS = 50.8</b>	<i>CARKD, CARS2, ING1, ANKRD10, ARHGEF7</i> Discordant
	<i>GPR180</i>	<i>TGDS, GPR180</i> <b>MCS = 67.6</b>	<i>TGDS, GPR180</i> <b>MCS = 74.8</b>	<i>TGDS, GPR180</i> Discordant
	<i>CLYBL</i>	No cluster <b>MCS = 60.2</b>	<i>TM9SF2, CLYBL</i> <b>MCS = 67.6</b>	Discontinuous Discordant
18q21.33-q23	<i>DSC2</i>	<i>DSC3, DSC2</i> <b>MCS = 73</b>	<i>DSC3, DSC2</i> <b>MCS = 66.5</b>	<i>DSC3, DCS2</i> Yes
	<i>DSG2</i>	<i>DSG3, DSG2</i> <b>MCS = 66.1</b>	No cluster <b>MCS = 60.8</b>	<i>DSG3, DSG2</i> Yes
	<i>RNF138</i>	No cluster	No cluster	No cluster No
	<i>MAPRE2</i>	No cluster	No cluster	No cluster No
21q22.3	<i>SERPINB5</i>	No cluster	No cluster	No cluster No
	<i>MBP</i>	No cluster	No cluster	No cluster No
	<i>NFATC1</i>	No cluster <b>MCS = 60.6</b>	<i>ATP9B, NFATC1</i> <b>MCS = 65.6</b>	No cluster Discordant
	<i>CDH19</i>	No cluster	No cluster	No cluster No
	<i>PQLC1</i>	<i>PQLC1, FLJ37638</i> <b>MCS = 74.6</b>	<i>PQLC1, FLJ37638</i> <b>MCS = 73.4</b>	Discontinuous Yes
	<i>WDR4</i>	No cluster	No cluster	No cluster No
	<i>CBS</i>	<i>CBS, U2AF1</i> <b>MCS = 67.1</b>	No cluster <b>MCS = 61.8</b>	<i>CBS, U2AF1</i> Yes
	<i>PDXK</i>	<i>RRP1B, PDXK, CSTB, RRP1</i> <b>MCS = 62.6</b>	<i>RRP1B, PDXK</i> <b>MCS = 50.9</b>	<i>RRP1B, PDXK, CSTB, RRP1</i> Yes
	<i>C21orf70</i>	No cluster	No cluster	No cluster No
	<i>SI100B</i>	No cluster	No cluster	<i>SI100B, PRMT2</i> No
	<i>TMPRSS3</i>	No cluster	No cluster	No cluster No
	<i>PDE9A</i>	No cluster	No cluster	No cluster No
	<i>AGPAT3</i>	No cluster	No cluster	<i>RRP1, AGPAT3</i> No
	<i>SUMO3</i>	<i>UBE2G2, SUMO3</i> <b>MCS = 66.2</b>	<i>UBE2G2, SUMO3</i> <b>(MCS = 65.7)</b>	<i>UBE2G2, SUMO3, PTTGIIP</i> Yes
	<i>SUMO3</i>	<i>SUMO3, PTTGIIP</i> <b>MCS = 75.1</b>	<i>SUMO3, PTTGIIP</i> <b>MCS = 73.2</b>	<i>UBE2G2, SUMO3, PTTGIIP</i> Yes

a: oestrogen receptor; b: multi-correlation score

Supplementary Figure 1: Level of expression genes belonging to *ESR1* cluster (*RMND1*, *C6orf211*, *C6orf97*, *ESR1*) in breast cancer patients according to oestrogen receptor (ER) status, based on bcGenExMiner v3.0 data.

