

Suppl. Table S8a: SAM-Result Poor Prognosis Marker

Affy_ID	GeneSymbol	Numerator(r)	Score(d)	Denominator(s+s0)	q-value(%)	264probe_Signature_FDR_25%	26probe_Signature_FDR_3.5%
211506_s_at	IL8	0,224	3,754	0,060	0,000	included	included
211708_s_at	SCD	0,210	3,377	0,062	0,000	included	included
39249_at	AQP3	0,164	3,308	0,050	0,000	included	included
202859_x_at	IL8	0,184	3,299	0,056	0,000	included	included
202627_s_at	SERPINE1	0,132	3,136	0,042	0,000	included	included
212909_at	LYPDC1	0,141	3,118	0,045	0,000	included	included
200737_at	PGK1	0,091	3,090	0,029	0,000	included	included
204344_s_at	SEC23A	0,148	3,075	0,048	0,000	included	included
205810_s_at	WASL	0,138	3,071	0,045	0,000	included	included
217356_s_at	PGK1	0,109	3,031	0,036	0,000	included	included
215779_s_at	HIST1H2BG	0,160	3,017	0,053	0,000	included	included
212344_at	SULF1	0,119	3,008	0,039	0,000	included	included
209875_s_at	SPP1	0,141	3,002	0,047	0,000	included	included
219434_at	TREM1	0,128	2,982	0,043	0,000	included	included
219508_at	GCNT3	0,118	2,966	0,040	0,000	included	included
208881_x_at	IDI1	0,095	2,959	0,032	0,000	included	included
215427_s_at	ZCCHC14	0,106	2,958	0,036	0,000	included	included
214603_at	MAGEA2	0,218	2,956	0,074	0,000	included	included
219875_s_at	PNAS-4	0,143	2,951	0,048	0,000	included	included
204083_s_at	TPM2	0,120	2,948	0,041	0,000	included	included
218468_s_at	GREM1	0,131	2,937	0,045	0,000	included	included
204615_x_at	IDI1	0,096	2,902	0,033	0,000	included	included
212354_at	SULF1	0,108	2,858	0,038	3,454	included	included
218469_at	GREM1	0,106	2,836	0,037	3,454	included	included
212353_at	SULF1	0,104	2,809	0,037	3,454	included	included
202497_x_at	SLC2A3	0,128	2,797	0,046	3,454	included	included
202539_s_at	HMGCR	0,068	2,745	0,025	7,741	included	no
214522_x_at	HIST1H3D	0,149	2,745	0,054	7,741	included	no
212942_s_at	KIAA1199	0,117	2,733	0,043	7,741	included	no
219773_at	NOX4	0,090	2,693	0,033	7,741	included	no
205680_at	MMP10	0,126	2,664	0,047	11,225	included	no
202912_at	ADM	0,106	2,656	0,040	11,225	included	no
201108_s_at	THBS1	0,098	2,654	0,037	11,225	included	no
210387_at	HIST1H2BG	0,146	2,653	0,055	11,225	included	no
205227_at	IL1RAP	0,111	2,648	0,042	11,225	included	no
201695_s_at	NP	0,065	2,645	0,025	11,225	included	no
217073_x_at	APOA1	0,089	2,595	0,034	11,225	included	no
201109_s_at	THBS1	0,107	2,595	0,041	11,225	included	no
215446_s_at	LOX	0,099	2,593	0,038	11,225	included	no
205443_at	SNAPC1	0,067	2,592	0,026	11,225	included	no
209978_s_at	LPA	0,101	2,591	0,039	11,225	included	no
201506_at	TGFBI	0,091	2,571	0,035	11,225	included	no
208577_at	HIST1H3C	0,086	2,569	0,034	11,225	included	no
219927_at	C14orf111	0,130	2,552	0,051	13,815	included	no
212210_at	DKFZP586J0619	0,107	2,539	0,042	13,815	included	no
202134_s_at	WWTR1	0,107	2,527	0,043	13,815	included	no
211162_x_at	SCD	0,140	2,526	0,055	13,815	included	no
218073_s_at	FLJ10407	0,073	2,524	0,029	13,815	included	no
212887_at	SEC23A	0,068	2,524	0,027	13,815	included	no
206354_at	SLCO1B3	0,106	2,522	0,042	13,815	included	no
201324_at	EMP1	0,089	2,520	0,035	13,815	included	no
212902_at	SEC24A	0,064	2,517	0,026	13,815	included	no
214978_s_at	PPFIA4	0,112	2,512	0,045	13,815	included	no
204338_s_at	RGS4	0,137	2,511	0,055	13,815	included	no
210089_s_at	LAMA4	0,117	2,502	0,047	13,815	included	no
214580_x_at	KRT6A	0,180	2,502	0,072	13,815	included	no
39248_at	AQP3	0,131	2,499	0,052	13,815	included	no
214725_at	RPESP	0,119	2,496	0,048	13,815	included	no
219104_at	RNF141	0,077	2,492	0,031	13,815	included	no
211840_s_at	PDE4D	0,098	2,489	0,039	13,815	included	no
200738_s_at	PGK1	0,061	2,482	0,024	13,815	included	no
203099_s_at	CDYL	0,138	2,480	0,056	13,815	included	no
207933_at	ZP2	0,082	2,477	0,033	13,815	included	no
202604_x_at	ADAM10	0,059	2,476	0,024	13,815	included	no
221291_at	ULBP2	0,118	2,472	0,048	13,815	included	no
214455_at	H2BFL	0,148	2,464	0,060	13,815	included	no
210431_at	ALPPL2	0,078	2,463	0,031	13,815	included	no
210845_s_at	PLAUR	0,066	2,454	0,027	14,967	included	no
211160_x_at	ACTN1	0,072	2,440	0,030	14,967	included	no
208547_at	HIST1H2BB	0,099	2,425	0,041	14,967	included	no
219837_s_at	CYTL1	0,112	2,414	0,047	14,967	included	no
201037_at	PFKP	0,092	2,409	0,038	14,967	included	no
200989_at	HIF1A	0,054	2,405	0,022	14,967	included	no
208490_x_at	HIST1H2BF	0,084	2,397	0,035	14,967	included	no
204298_s_at	LOX	0,098	2,387	0,041	16,648	included	no
201888_s_at	IL13RA1	0,069	2,387	0,029	16,648	included	no
220768_s_at	CSNK1G3	0,050	2,386	0,021	16,648	included	no
210405_x_at	TNFRSF10B	0,095	2,383	0,040	16,648	included	no
206315_at	CRLF1	0,163	2,382	0,069	16,648	included	no
217911_s_at	BAG3	0,055	2,378	0,023	16,648	included	no
219886_at	LRRIQ2	0,087	2,369	0,037	16,648	included	no
202628_s_at	SERPINE1	0,106	2,369	0,045	16,648	included	no

Affy_ID	GeneSymbol	Numerator(r)	Score(d)	Denominator(s+s0)	q-value(%)	264probe_Signature_FDR_25%	26probe_Signature_FDR_3.5%
209305_s_at	GADD45B	0,069	2,366	0,029	16,648	included	no
210995_s_at	TRIM23	0,085	2,365	0,036	16,648	included	no
206409_at	TIAM1	0,086	2,362	0,036	16,648	included	no
201325_s_at	EMP1	0,085	2,359	0,036	16,648	included	no
219366_at	AVEN	0,055	2,357	0,023	16,648	included	no
210435_at	NBR2	0,118	2,355	0,050	16,648	included	no
205199_at	CA9	0,097	2,350	0,041	16,648	included	no
209803_s_at	PHLDA2	0,097	2,341	0,042	16,648	included	no
205499_at	SRPX2	0,077	2,334	0,033	16,648	included	no
214216_s_at	KIAA0217	0,135	2,330	0,058	16,648	included	no
217834_s_at	SYNCRIP	0,067	2,321	0,029	16,648	included	no
204883_s_at	HUS1	0,083	2,311	0,036	16,648	included	no
208527_x_at	HIST1H2BE	0,073	2,305	0,032	19,072	included	no
208129_x_at	RUNX1	0,063	2,303	0,027	19,072	included	no
214612_x_at	MAGEA6	0,196	2,302	0,085	19,072	included	no
215488_at	VMD2	0,076	2,299	0,033	19,072	included	no
206512_at	U2AF1L1	0,086	2,298	0,038	19,072	included	no
217428_s_at	COL10A1	0,130	2,292	0,057	19,072	included	no
216796_s_at	- - -	0,066	2,290	0,029	19,072	included	no
208427_s_at	ELAVL2	0,115	2,284	0,051	19,072	included	no
213506_at	F2RL1	0,097	2,283	0,043	19,072	included	no
208180_s_at	H4FH	0,099	2,282	0,043	19,072	included	no
215997_s_at	CUL4B	0,057	2,282	0,025	19,072	included	no
203000_at	STMN2	0,121	2,282	0,053	19,072	included	no
202388_at	RGS2	0,099	2,278	0,044	19,072	included	no
205523_at	HAPLN1	0,157	2,278	0,069	19,072	included	no
217675_at	LOC201501	0,084	2,273	0,037	19,072	included	no
203393_at	HES1	0,072	2,271	0,032	19,072	included	no
214702_at	FN1	0,101	2,268	0,045	19,072	included	no
209601_at	ACOX1	0,063	2,268	0,028	19,072	included	no
204475_at	MMP1	0,187	2,266	0,083	19,072	included	no
220623_s_at	TSGA10	0,089	2,258	0,040	19,072	included	no
216729_at	- - -	0,097	2,252	0,043	19,072	included	no
210990_s_at	LAMA4	0,084	2,251	0,037	19,072	included	no
210257_x_at	CUL4B	0,054	2,246	0,024	19,072	included	no
220759_at	FAM12B	0,088	2,242	0,039	21,089	included	no
212543_at	AIM1	0,066	2,241	0,030	21,089	included	no
219856_at	SARG	0,148	2,240	0,066	21,089	included	no
207696_at	FUT9	0,089	2,238	0,040	21,089	included	no
220031_at	ZA20D1	0,068	2,237	0,030	21,089	included	no
201110_s_at	THBS1	0,088	2,232	0,040	21,089	included	no
208394_x_at	ESM1	0,065	2,225	0,029	21,089	included	no
215649_s_at	MVK	0,084	2,218	0,038	21,089	included	no
206686_at	PDK1	0,117	2,206	0,053	21,089	included	no
206547_s_at	PPEF1	0,083	2,203	0,038	21,089	included	no
221679_s_at	ABHD6	0,085	2,200	0,039	21,089	included	no
204337_at	RGS4	0,076	2,196	0,035	21,089	included	no
210233_at	IL1RAP	0,113	2,191	0,051	21,089	included	no
201170_s_at	BHLHB2	0,062	2,185	0,028	22,601	included	no
211338_at	IFNA2	0,095	2,185	0,044	22,601	included	no
216650_at	LOC442165	0,101	2,184	0,046	22,601	included	no
201903_at	UQCRC1	0,051	2,179	0,023	22,601	included	no
202740_at	ACY1	0,058	2,179	0,027	22,601	included	no
208523_x_at	HIST1H2BI	0,080	2,175	0,037	22,601	included	no
208487_at	LMX1B	0,062	2,173	0,029	22,601	included	no
207370_at	IBSP	0,098	2,171	0,045	22,601	included	no
206113_s_at	RAB5A	0,065	2,171	0,030	22,601	included	no
215733_x_at	CTAG2	0,160	2,169	0,074	22,601	included	no
210805_x_at	RUNX1	0,073	2,168	0,034	22,601	included	no
205924_at	RAB3B	0,091	2,163	0,042	22,601	included	no
216672_s_at	MYT1L	0,075	2,153	0,035	22,601	included	no
206569_at	IL24	0,090	2,152	0,042	22,601	included	no
211617_at	ALDOAP2	0,088	2,151	0,041	22,601	included	no
222219_s_at	TLE6	0,067	2,150	0,031	22,601	included	no
207328_at	ALOX15	0,084	2,149	0,039	22,601	included	no
202363_at	SPOCK	0,082	2,147	0,038	22,601	included	no
202543_s_at	GMFB	0,057	2,147	0,027	22,601	included	no
213563_s_at	GCP2	0,069	2,145	0,032	22,601	included	no
216268_s_at	JAG1	0,069	2,142	0,032	22,601	included	no
210874_s_at	NAT6	0,074	2,139	0,035	22,601	included	no
209101_at	CTGF	0,098	2,139	0,046	22,601	included	no
211758_x_at	TXNDC9	0,047	2,139	0,022	22,601	included	no
201559_s_at	CLIC4	0,079	2,137	0,037	22,601	included	no
213640_s_at	LOX	0,119	2,133	0,056	22,601	included	no
206788_s_at	CBFB	0,076	2,133	0,035	22,601	included	no
219232_s_at	EGLN3	0,075	2,133	0,035	22,601	included	no
219328_at	DDX31	0,094	2,129	0,044	22,601	included	no
202540_s_at	HMGCR	0,053	2,128	0,025	22,601	included	no
209398_at	HIST1H1C	0,092	2,128	0,043	22,601	included	no
215574_at	- - -	0,080	2,124	0,038	22,601	included	no
218036_x_at	NMD3	0,067	2,121	0,031	22,601	included	no
210750_s_at	DLGAP1	0,084	2,118	0,040	22,601	included	no
40524_at	PTPN21	0,075	2,117	0,035	22,601	included	no
201275_at	FDPS	0,058	2,113	0,027	22,601	included	no

Affy_ID	GeneSymbol	Numerator(r)	Score(d)	Denominator(s+s0)	q-value(%)	264probe_Signature_FDR_25%	26probe_Signature_FDR_3.5%
204614_at	SERPINB2	0,116	2,112	0,055	22,601	included	no
201043_s_at	ANP32A	0,095	2,109	0,045	22,601	included	no
201091_s_at	CBX3	0,057	2,108	0,027	22,601	included	no
37512_at	RODH	0,084	2,106	0,040	22,601	included	no
203108_at	GPCR5A	0,149	2,105	0,071	22,601	included	no
219948_x_at	FLJ21934	0,123	2,105	0,059	22,601	included	no
215646_s_at	CSPG2	0,115	2,098	0,055	22,601	included	no
213983_s_at	SCC-112	0,092	2,098	0,044	22,601	included	no
202057_at	KPNA1	0,069	2,098	0,033	22,601	included	no
208613_s_at	FLNB	0,073	2,097	0,035	22,601	included	no
220003_at	FLJ11004	0,086	2,095	0,041	22,601	included	no
201208_s_at	TNFAIP1	0,079	2,095	0,037	22,601	included	no
209822_s_at	VLDLR	0,062	2,095	0,029	22,601	included	no
209909_s_at	TGFB2	0,105	2,092	0,050	22,601	included	no
210467_x_at	MAGEA12	0,114	2,088	0,055	22,601	included	no
215432_at	BUCS1	0,100	2,085	0,048	22,601	included	no
209122_at	ADFP	0,073	2,084	0,035	22,601	included	no
210196_s_at	PSG1	0,103	2,083	0,049	22,601	included	no
210876_at	ANXA2	0,063	2,081	0,030	22,601	included	no
208546_x_at	HIST1H2BH	0,099	2,079	0,048	22,601	included	no
216414_at	- - -	0,068	2,079	0,033	22,601	included	no
212812_at	- - -	0,059	2,078	0,029	22,601	included	no
219733_s_at	SLC27A5	0,087	2,078	0,042	22,601	included	no
215254_at	DSCR1	0,091	2,074	0,044	22,601	included	no
207319_s_at	CDC2L5	0,060	2,070	0,029	22,601	included	no
211668_s_at	PLAU	0,078	2,068	0,038	22,601	included	no
201792_at	AEBP1	0,081	2,066	0,039	22,601	included	no
202238_s_at	NNMT	0,083	2,065	0,040	26,367	included	no
219356_s_at	SNF7DC2	0,060	2,060	0,029	26,367	included	no
210904_s_at	IL13RA1	0,076	2,059	0,037	26,367	included	no
221552_at	ABHD6	0,074	2,059	0,036	26,367	included	no
221009_s_at	ANGPTL4	0,094	2,058	0,046	26,367	included	no
214968_at	DDX51	0,083	2,055	0,041	26,367	included	no
201196_s_at	AMD1	0,063	2,055	0,031	26,367	included	no
212444_at	- - -	0,134	2,054	0,065	26,367	included	no
204845_s_at	ENPEP	0,077	2,053	0,038	26,367	included	no
208496_x_at	HIST1H3G	0,108	2,053	0,053	26,367	included	no
215464_s_at	TAX1BP3	0,050	2,052	0,024	26,367	included	no
212797_at	SORT1	0,099	2,051	0,048	26,367	included	no
204596_s_at	STC1	0,054	2,050	0,026	26,367	included	no
210623_at	LOC51035	0,081	2,050	0,039	26,367	included	no
216915_s_at	PTPN12	0,093	2,050	0,045	26,367	included	no
208144_s_at	PP1345	0,067	2,049	0,033	26,367	included	no
214469_at	HIST1H2AE	0,128	2,049	0,062	26,367	included	no
205479_s_at	PLAU	0,065	2,044	0,032	26,367	included	no
217448_s_at	C14orf92	0,070	2,044	0,034	26,367	included	no
216607_s_at	CYP51A1	0,061	2,043	0,030	26,367	included	no
210619_s_at	HYAL1	0,089	2,042	0,044	26,367	included	no
214540_at	HIST1H2BO	0,099	2,039	0,048	26,367	included	no
216549_s_at	TBC1D22B	0,096	2,039	0,047	26,367	included	no
219478_at	WFDC1	0,111	2,037	0,054	26,367	included	no
209624_s_at	MCCC2	0,078	2,034	0,038	26,367	included	no
212575_at	C19orf6	0,102	2,034	0,050	26,367	included	no
222379_at	KCNE4	0,082	2,031	0,040	26,367	included	no
211924_s_at	PLAUR	0,070	2,030	0,034	26,367	included	no
221933_at	NLGN4X	0,114	2,030	0,056	26,367	included	no
220414_at	CALML5	0,161	2,029	0,079	26,367	included	no
215976_at	DBC1	0,087	2,026	0,043	26,367	included	no
210546_x_at	CTAG1B	0,183	2,025	0,090	26,367	included	no
216712_at	SLC25A30	0,086	2,020	0,042	26,367	included	no
220106_at	NPC1L1	0,074	2,020	0,037	26,367	included	no
212898_at	KIAA0406	0,051	2,019	0,025	26,367	included	no
202213_s_at	CUL4B	0,054	2,019	0,027	26,367	included	no
216830_at	- - -	0,078	2,017	0,039	26,367	included	no
204325_s_at	NF1	0,050	2,017	0,025	26,367	included	no
215423_at	- - -	0,067	2,016	0,033	26,367	included	no
205228_at	RBMS2	0,077	2,013	0,038	26,367	included	no
208083_s_at	ITGB6	0,107	2,011	0,053	26,367	included	no
218182_s_at	CLDN1	0,102	2,011	0,051	26,367	included	no

Suppl. Table S8b: SAM-Result Good Prognosis Marker							
Affy_ID	GeneSymbol	Numerator(r)	Score(d)	Denominator(s+s0)	q-value(%)	264probe_Signature_FDR_25%	26probe_Signature_FDR_3.5%
209591_s_at	BMP7	-0,154	-3,405	0,045	21,089	included	no
221671_x_at	IGKC	-0,144	-3,118	0,046	21,089	included	no
211333_s_at	FASLG	-0,074	-3,107	0,024	21,089	included	no
221651_x_at	IGKC	-0,136	-3,097	0,044	21,089	included	no
211259_s_at	BMP7	-0,135	-3,041	0,044	21,089	included	no
218872_at	TSC	-0,142	-3,019	0,047	21,089	included	no
209460_at	ABAT	-0,122	-3,010	0,041	21,089	included	no
217455_s_at	SSTR2	-0,082	-2,890	0,028	21,089	included	no
208479_at	KCNA1	-0,100	-2,883	0,035	21,089	included	no
205890_s_at	UBD	-0,188	-2,860	0,066	21,089	included	no
221087_s_at	APOL3	-0,076	-2,843	0,027	21,089	included	no
210321_at	GZMH	-0,150	-2,820	0,053	21,089	included	no
214510_at	GPR20	-0,124	-2,777	0,045	26,367	included	no
211430_s_at	IGH@	-0,155	-2,761	0,056	26,367	included	no
214567_s_at	XCL1	-0,171	-2,759	0,062	26,367	included	no
217143_s_at	TRA@	-0,129	-2,722	0,048	26,367	included	no
207796_x_at	KLRD1	-0,086	-2,690	0,032	26,367	included	no
214677_x_at	IGL@	-0,146	-2,688	0,054	26,367	included	no
203915_at	CXCL9	-0,152	-2,681	0,057	26,367	included	no
208498_s_at	AMY2A	-0,141	-2,679	0,052	26,367	included	no
219605_at	ZNF3	-0,086	-2,677	0,032	26,367	included	no
209138_x_at	IGLC2	-0,149	-2,661	0,056	26,367	included	no
209590_at	BMP7	-0,101	-2,654	0,038	26,367	included	no
204781_s_at	FAS	-0,077	-2,653	0,029	26,367	included	no
217378_x_at	LOC391427	-0,212	-2,641	0,080	26,367	included	no
218062_x_at	CDC42EP4	-0,083	-2,631	0,032	26,367	included	no
212272_at	LPIN1	-0,101	-2,624	0,038	26,367	included	no
203608_at	ALDH5A1	-0,108	-2,621	0,041	26,367	included	no
207747_s_at	DOK4	-0,086	-2,619	0,033	26,367	included	no