

**Supplementary Table 2.** List of hypermethylated genes in stage I-III and stage IV CRCs

	GeneName	Location*
Frequently methylated only in stage I-III CRCs (291 genes)		
	<i>MFSD7</i>	chr4:673998-674058
	<i>AKAP12</i>	chr6:151688516-151719602
	<i>TMEM46</i>	chr13:25516735-25523198
	<i>NFATC1</i>	chr18:75388107-75390311
	<i>ZNF382</i>	chr19:41788061-41811339
	<i>CCDC8</i>	chr19:51605929-51608681
	<i>CDO1</i>	chr5:115168329-115180304
	<i>IRF4</i>	chr6:336752-347637
	<i>OPRK1</i>	chr8:54300829-54326747
	<i>VIM</i>	chr10:17311281-17319598
	<i>CCNT1</i>	chr12:47373019-47397048
	<i>WDR81</i>	chr17:1577613-1588629
	<i>LHX1</i>	chr17:32369870-32372804
	<i>LIMD2</i>	chr17:59126981-59131777
	<i>GALR1</i>	chr18:73090996-73111084
	<i>ARHGEF18</i>	chr19:7417929-7443363
	<i>NTSR1</i>	chr20:60810634-60864568
	<i>SCARF2</i>	chr22:19108875-19122146
	<i>KLHDC7B</i>	chr22:49333328-49335409
	<i>IGSF21</i>	chr1:18560448-18577563
	<i>C1orf61</i>	chr1:154662053-154667117
	<i>MIXL1</i>	chr1:224478006-224480136
	<i>SPDYA</i>	chr2:28892367-28917179
	<i>SIX2</i>	chr2:45085847-45090047
	<i>ARHGEF4</i>	chr2:131404995-131521306
	<i>FNI</i>	chr2:215943262-215956451
	<i>WWTR1</i>	chr3:150720722-150903805
	<i>RNF150</i>	chr4:142006175-142274066
	<i>ADAMTS16</i>	chr5:5193443-5373412
	<i>PACSIN1</i>	chr6:34590642-34610978
	<i>KCNK9</i>	chr8:140699431-140784481
	<i>BC027908:289</i>	chr10:125757497-125757497
	<i>FANK1</i>	chr10:127575098-127688151
	<i>BCAT1</i>	chr12:24855546-24993575
	<i>COL4A1</i>	chr13:109599442-109757497
	<i>LBXCOR1</i>	chr15:65904995-65913228
	<i>SNTB2</i>	chr16:67778551-67900456
	<i>ALOX12</i>	chr17:6853532-6854779
	<i>TRMT1</i>	chr19:13076715-13088563
	<i>NPAS1</i>	chr19:52215983-52240857

	<i>ZNF671</i>	chr19:62922931-62930807
	<i>EEF1A2</i>	chr20:61589810-61600949
	<i>GALNT14</i>	chr2:31008438-31231572
	<i>GAD2</i>	chr10:26545600-26548253
	<i>FGF14</i>	chr13:101171206-101852125
	<i>HBQ1</i>	chr16:170335-171178
	<i>SLC6A2</i>	chr16:54247523-54295201
	<i>RAB34</i>	chr17:24065430-24069314
	<i>RNF126</i>	chr19:598526-614227
	<i>ZNF154</i>	chr19:62903622-62912391
	<i>BC057764:-301</i>	chr2:132621639-132635995
	<i>EVC2</i>	chr4:5615053-5762176
	<i>ANXA5</i>	chr4:122808602-122837597
	<i>POU4F2</i>	chr4:147779495-147783073
	<i>BVES</i>	chr6:105651392-105691742
	<i>CDKN2A</i>	chr9:21964476-21964826
	<i>DCLRE1C</i>	chr10:14988877-15036100
	<i>KCNA1</i>	chr12:4890806-4892293
	<i>RNF113B</i>	chr13:97626040-97627522
	<i>RASGRF1</i>	chr15:77057777-77143996
	<i>RPRML</i>	chr17:42410521-42411613
	<i>SDCBP2</i>	chr20:1238621-1257838
	<i>MAFB</i>	chr20:38747931-38751290
	<i>GNAS</i>	chr20:56899820-56919645
	<i>PDXK</i>	chr21:43963406-44006616
	<i>DHRS3</i>	chr1:12561571-12600407
	<i>RSPO1</i>	chr1:37850008-37873078
	<i>ATP8B2</i>	chr1:152567899-152576644
	<i>MFSD4</i>	chr1:203804735-203838669
	<i>PTPRN</i>	chr7:157024516-158073179
	<i>EDEM1</i>	chr3:5204433-5236642
	<i>DALRD3</i>	chr3:49027439-49033471
	<i>MINA</i>	chr3:99143351-99173985
	<i>MSX1</i>	chr4:4912293-4916561
	<i>C6orf114</i>	chr6:13578081-13578491
	<i>TRAF3IP2</i>	chr6:112000613-112034014
	<i>IGFBP3</i>	chr7:45918369-45927396
	<i>EPHB4</i>	chr7:100238123-100263079
	<i>LBX1</i>	chr10:102976723-102978707
	<i>ACY3</i>	chr11:67166602-67174706
	<i>ZIC5</i>	chr13:99413276-99422179
	<i>MESP1</i>	chr15:88094110-88095545
	<i>CIDEA</i>	chr18:12244628-12267594

	<i>ST8SIA5</i>	chr18:42525411-42591037
	<i>MATK</i>	chr19:3728967-3737415
	<i>ICAM1</i>	chr19:10242517-10258291
	<i>C20orf117</i>	chr20:34845599-34878124
	<i>ZNF335</i>	chr20:44027674-44032271
	<i>CDH4</i>	chr20:59608229-59945694
	<i>FOXD3</i>	chr1:63561318-63563385
	<i>NTRK1</i>	chr1:155097295-155118266
	<i>HLX1</i>	chr1:219119366-219125023
	<i>ITPKB</i>	chr1:224961524-224993499
	<i>FKBP1B</i>	chr2:24130276-24140054
	<i>BOC</i>	chr3:114414065-114473142
	<i>MCF2L2</i>	chr3:184428511-184628549
	<i>GSH2</i>	chr4:54661005-54662879
	<i>TCF7</i>	chr5:133507182-133511819
	<i>MGC13017</i>	chr2:114073075-114075623
	<i>FOXC1</i>	chr6:1555680-1559128
	<i>BMP6</i>	chr6:7672010-7826960
	<i>TRIM26</i>	chr6_qbl_hap2:1399094-1428021
	<i>DDR1</i>	chr6_qbl_hap2:2097628-2114869
	<i>HTR1B</i>	chr6:78228667-78229839
	<i>OLIG3</i>	chr6:137855029-137857224
	<i>SNX10</i>	chr7:26370470-26380474
	<i>LIMK1</i>	chr7:73136092-73174790
	<i>SLC25A40</i>	chr7:87303354-87343604
	<i>DGKI</i>	chr7:136724925-137182149
	<i>ADRA1A</i>	chr8:26669830-26778839
	<i>DUSP4</i>	chr8:29249537-29264104
	<i>PSKH2</i>	chr8:87129807-87150967
	<i>LAPTM4B</i>	chr8:98856985-98934006
	<i>C9orf93</i>	chr9:15542872-15734775
	<i>ODF2</i>	chr9:130258550-130287574
	<i>HPSE2</i>	chr10:100208867-100985609
	<i>SLC18A2</i>	chr10:118990706-119027085
	<i>MGMT</i>	chr10:131155456-131455358
	<i>STK32C</i>	chr10:133888649-133996053
	<i>C10orf93</i>	chr10:134600613-134606054
	<i>BTG4</i>	chr11:110872450-110888274
	<i>PRPF40B</i>	chr12:48311910-48324716
	<i>FOXO1</i>	chr13:39946131-40083264
	<i>PHF11</i>	chr13:48968529-49001118
	<i>FARP1</i>	chr13:97593435-97900024
	<i>KDELC1</i>	chr13:102234632-102249405

	<i>TRIM9</i>	chr14:50532535-50632172
	<i>PUNC</i>	chr15:63407664-63408988
	<i>ZNF771</i>	chr16:30326437-30348421
	<i>PRR14</i>	chr16:30572931-30574036
	<i>PARD6A</i>	chr16:66252636-66254182
	<i>KIAA0182</i>	chr16:84254451-84267313
	<i>NEUROD2</i>	chr17:35015230-35016378
	<i>HSPB9</i>	chr17:37528395-37528874
	<i>CYB561</i>	chr17:58863397-58877277
	<i>SLC16A5</i>	chr17:70596628-70613843
	<i>FLJ35220</i>	chr17:76003710-76018737
	<i>CCDC57</i>	chr17:77704775-77736165
	<i>BRUNOL4</i>	chr18:33077006-33399998
	<i>REEP6</i>	chr19:1442165-1448924
	<i>GIPC3</i>	chr19:3536569-3541434
	<i>ZNF557</i>	chr19:7020471-7038978
	<i>ELAVL3</i>	chr19:11423143-11452803
	<i>CACNA1A</i>	chr19:13178257-13478274
	<i>GGN</i>	chr19:43567901-43570508
	<i>RELB</i>	chr19:50196552-50233292
	<i>ZNF586</i>	chr19:62972837-62983796
	<i>SRC</i>	chr20:35407971-35467235
	<i>PLTP</i>	chr20:43960804-43974193
	<i>RBM38</i>	chr20:55399870-55417792
	<i>STX16</i>	chr20:56659734-56687988
	<i>DERL3</i>	chr22:22506911-22511201
	<i>WDR45</i>	chrX:48819036-48845003
	<i>PLP2</i>	chrX:48915217-48918382
	<i>NKAP</i>	chrX:118943051-118961763
	<i>MTM1</i>	chrX:149487705-149592274
	<i>PAX7</i>	chr1:18830087-18935219
	<i>SFPQ</i>	chr1:35421788-35431322
	<i>UROD</i>	chr1:45250417-45252835
	<i>KCND3</i>	chr1:112119977-112333300
	<i>SEMA6C</i>	chr1:149370787-149385728
	<i>SLC39A1</i>	chr1:152198212-152206812
	<i>HNRPU</i>	chr1:243080225-243094450
	<i>MYCN</i>	chr2:15999497-16004580
	<i>MSH2</i>	chr2:47526191-47555913
	<i>MAP4K4</i>	chr2:101847824-101877584
	<i>LYPD6</i>	chr2:149932738-150038384
	<i>GAD1</i>	chr2:171413823-171425905
	<i>OBFC2A</i>	chr2:192251961-192259906

	<i>FLJ37953</i>	chr2:200515053-200528704
	<i>PTHR2</i>	chr2:209061982-209413063
	<i>IRS1</i>	chr2:227367970-227371698
	<i>VHL</i>	chr3:10158319-10168746
	<i>C3orf60</i>	chr3:49034078-49035930
	<i>RRP9</i>	chr3:51942486-51950962
	<i>RPN1</i>	chr3:129821503-129852409
	<i>FNDC3B</i>	chr3:173241080-173601181
	<i>RNF168</i>	chr3:197683025-197714979
	<i>WDR53</i>	chr3:197765456-197779810
	<i>SLIT2</i>	chr4:19864333-20229886
	<i>KCNIP4</i>	chr4:20339337-21559472
	<i>APBB2</i>	chr4:40510996-40641847
	<i>RAP1GDS1</i>	chr4:99401550-99584035
	<i>BC033172:-620</i>	chr4:183299799-183301557
	<i>PDLIM3</i>	chr4:186659845-186693706
	<i>ANKRD43</i>	chr5:132176932-132180388
	<i>LOC153328</i>	chr5:135198264-135252223
	<i>NRG2</i>	chr5:139207444-139403063
	<i>ADRA1B</i>	chr5:159276318-159332595
	<i>NKX2-5</i>	chr5:172591744-172594868
	<i>SERPINB6</i>	chr6:2893392-2917398
	<i>SLC35D3</i>	chr6:137285095-137288469
	<i>BC014953:110</i>	chr6:139391512-139406132
	<i>TTYH3</i>	chr7:2653578-2670962
	<i>DAGLB</i>	chr7:6415285-6490374
	<i>KIAA1706</i>	chr7:36159361-36307677
	<i>ASL</i>	chr7:65178269-65186106
	<i>TMEM130</i>	chr7:98283077-98305573
	<i>PTCD1</i>	chr7:98854682-98874355
	<i>ZNF789</i>	chr7:98908451-98923153
	<i>CUTL1</i>	chr7:101247602-101680064
	<i>NOM1</i>	chr7:156447606-156449580
	<i>SOX7</i>	chr8:10618688-10625432
	<i>CEBPD</i>	chr8:48812029-48813279
	<i>CA13</i>	chr8:86344968-86383554
	<i>OSGIN2</i>	chr8:90983935-91009270
	<i>FBXO43</i>	chr8:101218297-101227252
	<i>DPYS</i>	chr8:105461917-105500258
	<i>ZHX2</i>	chr8:124032932-124035445
	<i>NIBP</i>	chr8:140811770-141537860
	<i>PCSK5</i>	chr9:77986134-77998155
	<i>FSD1L</i>	chr9:107250146-107333065

	<i>FAM78A</i>	chr9:133123286-133141727
	<i>BC037281:-910</i>	chr10:11693309-11694231
	<i>CACNB2</i>	chr10:18669620-18847351
	<i>KIAA1217</i>	chr10:24548561-24876778
	<i>PAPSS2</i>	chr10:89410692-89477226
	<i>AVP11</i>	chr10:99427171-99437005
	<i>HMX2</i>	chr10:124897628-124900178
	<i>RPLP2</i>	chr11:800185-802876
	<i>CCKBR</i>	chr11:6248582-6249151
	<i>KCNJ11</i>	chr11:17363983-17367440
	<i>NRXN2</i>	chr11:64130222-64247236
	<i>RASGRP2</i>	chr11:64250959-64269504
	<i>FGF19</i>	chr11:69222187-69228287
	<i>ARHGEF17</i>	chr11:72697317-72757784
	<i>ESAM</i>	chr11:124128302-124137433
	<i>FLI1</i>	chr11:128069199-128186099
	<i>ITPR2</i>	chr12:26431634-26538540
	<i>PTHLH</i>	chr12:28006521-28016183
	<i>KCNH3</i>	chr12:48219207-48238344
	<i>NDUFA4L2</i>	chr12:55914953-55920742
	<i>AVPR1A</i>	chr12:61826483-61832857
	<i>NFYB</i>	chr12:103049040-103056170
	<i>MSI1</i>	chr12:119263516-119291341
	<i>DYNLL1</i>	chr12:119418242-119420681
	<i>ATP8A2</i>	chr13:25039271-25399421
	<i>HS6ST3</i>	chr13:95541094-96289813
	<i>LRP10</i>	chr11:46876685-46896652
	<i>TM9SF1</i>	chr14:23730782-23732315
	<i>C14orf104</i>	chr14:49161642-49171698
	<i>GALNTL1</i>	chr14:68796668-68889141
	<i>ZFYVE1</i>	chr14:72505912-72563592
	<i>CINP</i>	chr14:101884369-101899006
	<i>SIVA1</i>	chr14:104290515-104297041
	<i>GABRG3</i>	chr15:24799263-25451729
	<i>KIAA1370</i>	chr15:50664234-50758245
	<i>ETFA</i>	chr15:74295684-74390865
	<i>ZNF75A</i>	chr16:3298361-3308577
	<i>C16orf45</i>	chr16:15503624-15589418
	<i>SULT1A1</i>	chr16:28524417-28542367
	<i>YPEL3</i>	chr16:30012584-30015022
	<i>ZNF688</i>	chr16:30488524-30491229
	<i>GPT2</i>	chr16:45515246-45522699
	<i>CDH11</i>	chr16:63538184-63596445

	<i>MON1B</i>	chr16:75785366-75791044
	<i>CYB5D2</i>	chr17:3993211-4007740
	<i>HES7</i>	chr17:7965024-7968127
	<i>NTN1</i>	chr17:8865584-9088042
	<i>ULK2</i>	chr17:19618058-19711822
	<i>TNFAIP1</i>	chr17:23686913-23698160
	<i>GJC1</i>	chr17:40237521-40238711
	<i>SOCS3</i>	chr17:73866094-73866771
	<i>DSC3</i>	chr18:26824050-26876779
	<i>CBLN2</i>	chr18:68354895-68362703
	<i>SALL3</i>	chr18:74854042-74859181
	<i>FAM108A1</i>	chr19:1827975-1836432
	<i>PLEKHJ1</i>	chr19:2184155-2187951
	<i>TLE6</i>	chr19:2928562-2946176
	<i>ATCAY</i>	chr19:3860484-3879080
	<i>SH3GL1</i>	chr19:4311367-4351471
	<i>CLPP</i>	chr19:6317251-6319915
	<i>DNAJB1</i>	chr19:14486582-14490201
	<i>MDS032</i>	chr19:17187155-17191638
	<i>FAM125A</i>	chr19:17393714-17397140
	<i>FOSB</i>	chr19:50663093-50670277
	<i>HIF3A</i>	chr19:51498716-51538530
	<i>SLC17A7</i>	chr19:54628328-54633383
	<i>PNKP</i>	chr19:55056273-55062630
	<i>SYT5</i>	chr19:60378352-60382266
	<i>ZNF419A</i>	chr19:62690891-62697860
	<i>C20orf42</i>	chr20:6003492-6052191
	<i>ATP6V1E1</i>	chr22:16454903-16491588
	<i>AIFM3</i>	chr22:19658802-19665649
	<i>PHF21B</i>	chr22:43655706-43784245
	<i>BRD1</i>	chr22:48552941-48604456
	<i>CTPS2</i>	chrX:16516043-16640980
	<i>APOO</i>	chrX:23761398-23835955
	<i>OTUD5</i>	chrX:48664432-48699837
	<i>LOC203547</i>	chrX:150316435-150324185
	<i>TMSB4Y</i>	chrY:14324841-14327296
Frequently methylated only in stage IV CRCs (49 genes)		
	<i>ADAM12</i>	chr10:127719904-128067055
	<i>AGTR1</i>	chr3:149930657-149943480
	<i>ALAD</i>	chr9:115188413-115203439
	<i>ANTXR1</i>	chr2:69093993-69329806
	<i>BC058917:-87</i>	chr12:77782972-78367943
	<i>C16orf46</i>	chr16:79644603-79668373

	<i>CARM1</i>	chr19:10843253-10894448
	<i>CCDC12</i>	chr3:46938225-46993243
	<i>CNIH3</i>	chr1:222870802-222994872
	<i>DNAJB2</i>	chr2:219855708-219859866
	<i>DYX1C1</i>	chr15:53509798-53587724
	<i>EML3</i>	chr11:62130111-62135392
	<i>FBXO44</i>	chr1:11637501-11645971
	<i>FHL2</i>	chr2:105343715-105382113
	<i>FLJ10916</i>	chr2:88250950-88267261
	<i>FLJ39237</i>	chr7:100600498-100610277
	<i>FLJ44815</i>	chr17:29925255-29930501
	<i>FLT4</i>	chr5:179967359-180009230
	<i>GPR137</i>	chr11:63794110-63812262
	<i>GRAMD2</i>	chr15:70241659-70277180
	<i>HOXD4</i>	chr2:176724359-176726195
	<i>INTS3</i>	chr1:151967173-152013179
	<i>KCNK2</i>	chr1:213245508-213477059
	<i>KHK</i>	chr2:27163115-27177123
	<i>LINGO1</i>	chr15:75692424-75775530
	<i>LOC375748</i>	chr9:130078246-130088634
	<i>LOC554235</i>	chr19:55706669-55709759
	<i>LPPR2</i>	chr19:11327107-11337372
	<i>LRRC20</i>	chr10:71728735-71812388
	<i>MAN2C1</i>	chr15:73435186-73447994
	<i>MRPS25</i>	chr3:15065023-15081820
	<i>MSC</i>	chr8:72916331-72919285
	<i>NUP98</i>	chr11:3689635-3775468
	<i>PPIF</i>	chr10:80777226-80785095
	<i>SIAH1</i>	chr16:46952992-46953840
	<i>SLC37A3</i>	chr7:139680021-139744780
	<i>TIFA</i>	chr4:113418467-113419021
	<i>TMEM139</i>	chr7:142692166-142695263
	<i>VGf</i>	chr7:100592517-100595572
	<i>VGLL3</i>	chr3:87069813-87122947
	<i>VPS16</i>	chr20:2792595-2795380
	<i>ZNF19</i>	chr16:70065477-70080718
	<i>ZNF207</i>	chr17:27711730-27732088
	<i>ZNF259</i>	chr11:116154486-116163949
	<i>ZNF526</i>	chr19:47420396-47422408
	<i>ZNF550</i>	chr19:62745020-62763040
	<i>ZNF563</i>	chr19:12291005-12305502
	<i>ZNF790</i>	chr19:42000171-42021124
	<i>ZBPB</i>	chr7:49947585-50103372



Frequently methylated in both stage I-III and stage IV CRCs (290 genes)		
	<i>C9orf50</i>	chr9:131414325-131422876
	<i>SNF1LK</i>	chr21:43658827-43671430
	<i>C2orf32</i>	chr2:68373646-68400687
	<i>ITGA4</i>	chr2:182095156-182110713
	<i>TMEFF2</i>	chr2:192749714-192767889
	<i>FGF12</i>	chr3:193342422-193928066
	<i>CCDC4</i>	chr4:41807714-41849652
	<i>MDF1</i>	chr6:41714176-41729960
	<i>RSPO2</i>	chr8:108980720-109164742
	<i>DMRT1</i>	chr9:831690-959090
	<i>NRG3</i>	chr10:84549478-84735341
	<i>PHOX2A</i>	chr11:71627769-71632868
	<i>TMEM132D</i>	chr12:128122224-128954165
	<i>LRFN5</i>	chr14:41238380-41443499
	<i>SALL1</i>	chr16:49728524-49742684
	<i>AK124155:-10</i>	chr19:12064078-12071475
	<i>AK124155:112</i>	chr19:12064078-12071475
	<i>ZNF667</i>	chr19:61643018-61680555
	<i>TBX15</i>	chr1:119227189-119243271
	<i>ALK</i>	chr2:29269144-29997936
	<i>ZIC1</i>	chr3:148609871-148617196
	<i>KCNAB1</i>	chr3:157491470-157739621
	<i>NPY2R</i>	chr4:156354542-156355687
	<i>CYP26C1</i>	chr10:94811011-94818444
	<i>MYOD1</i>	chr11:17697686-17700254
	<i>NELL1</i>	chr11:20647712-21553577
	<i>CHST11</i>	chr12:103374879-103676013
	<i>CUTL2</i>	chr12:109956212-110272739
	<i>SLC13A5</i>	chr17:6538140-6557388
	<i>C17orf46</i>	chr17:40687543-40695262
	<i>VSX1</i>	chr20:25000132-25010767
	<i>GRIK1</i>	chr21:29875616-30234153
	<i>DSCR6</i>	chr21:37300320-37313828
	<i>FRMPD4</i>	chrX:12066506-12650450
	<i>PAP2D</i>	chr1:99128389-99243037
	<i>LOC388969</i>	chr2:85690667-85692663
	<i>GATA2</i>	chr3:129680960-129694718
	<i>GATA4</i>	chr8:11599126-11603846
	<i>XKR4</i>	chr8:56177571-56601264
	<i>FLJ46831</i>	chr10:129425945-129429440
	<i>HNT</i>	chr11:131586861-131711926
	<i>ST6GALNAC5</i>	chr1:77105774-77302325

	<i>GDNF</i>	chr5:37851510-37870767
	<i>AK023251:-177</i>	chr6:72180870-72187169
	<i>PRDM14</i>	chr8:71126577-71146116
	<i>GFRA1</i>	chr10:117812943-118022966
	<i>NTRK3</i>	chr15:86321599-86377280
	<i>CA10</i>	chr17:47065836-47592376
	<i>NETO1</i>	chr18:68652255-68683914
	<i>KCNG1</i>	chr20:49058624-49073024
	<i>CSPG5</i>	chr3:47578732-47595363
	<i>MATR3</i>	chr5:138637691-138686555
	<i>PENK</i>	chr8:57520358-57521788
	<i>SYT9</i>	chr11:7229757-7446846
	<i>PTPN5</i>	chr11:18706051-18769965
	<i>GABRA5</i>	chr15:24663151-24776749
	<i>DKFZp666G057</i>	chr15:80342207-80364320
	<i>NRXN1</i>	chr2:51003297-51113178
	<i>TMEM132E</i>	chr17:29931881-29990450
	<i>TDRD10</i>	chr1:152741319-152787247
	<i>TFPI2</i>	chr7:93353681-93358001
	<i>FBN1</i>	chr15:46487797-46547591
	<i>PAX1</i>	chr20:21634297-21644620
	<i>hsa-mir-124a-3:-454</i>	chr20:61280297-61280383
	<i>HTR2C</i>	chrX:113724807-114050880
	<i>GRIK3</i>	chr1:37042806-37272431
	<i>ESR1</i>	chr6:152243483-152423952
	<i>WDR86</i>	chr7:150709140-150738673
	<i>NRG1</i>	chr8:32624071-32720312
	<i>AK094650:-294</i>	chr8:69406011-69522799
	<i>DBC1</i>	chr9:121015719-121171560
	<i>URM1</i>	chr9:130173461-130192834
	<i>MYO3A</i>	chr10:26264679-26541471
	<i>TRPC6</i>	chr11:100829591-100959869
	<i>FZD10</i>	chr12:129212985-129216238
	<i>CCNA1</i>	chr13:35904409-35915019
	<i>CLDN10</i>	chr13:95002961-95029907
	<i>SLC24A4</i>	chr14:91975432-92032349
	<i>ABHD9</i>	chr19:15198730-15204768
	<i>ZNF415</i>	chr19:58302944-58327957
	<i>CHST2</i>	chr3:144322349-144323941
	<i>NXPH1</i>	chr7:8440110-8759118
	<i>TWIST1</i>	chr7:19122861-19123469
	<i>WBSCR17</i>	chr7:70403664-70816520
	<i>DBX1</i>	chr11:20134336-20138446

	<i>FADS1</i>	chr11:61323677-61340886
	<i>THY1</i>	chr11:118794098-118799064
	<i>FAIM2</i>	chr12:48546947-48584032
	<i>TRHDE</i>	chr12:70952730-71345689
	<i>FBXO39</i>	chr17:6630823-6641144
	<i>SLC4A11</i>	chr20:3156063-3167836
	<i>CBLN4</i>	chr20:54005903-54013419
	<i>STOX2</i>	chr4:185167305-185175869
	<i>COL9A1</i>	chr6:70982464-71069507
	<i>RSPO3</i>	chr6:127481741-127560603
	<i>SLC46A2</i>	chr9:114681021-114692866
	<i>SOX5</i>	chr12:23587412-23995132
	<i>ACAN</i>	chr15:87200760-87219589
	<i>FLJ40125</i>	chr19:50692354-50697604
	<i>PPP1R16B</i>	chr20:36897882-36985081
	<i>BHLHB4</i>	chr20:61107846-61108787
	<i>TMEM61</i>	chr1:55219053-55230554
	<i>FBXL7</i>	chr5:15553305-15992900
	<i>KIAA1688</i>	chr8:145725371-145809696
	<i>PVRL1</i>	chr11:119050309-119104645
	<i>ADCY4</i>	chr14:23871301-23874102
	<i>PRIMA1</i>	chr14:93254397-93324519
	<i>BNC1</i>	chr15:81715659-81744472
	<i>HBM</i>	chr16:155973-156767
	<i>CBLN1</i>	chr16:47870200-47873184
	<i>SLFN13</i>	chr17:30786228-30799885
	<i>NCAM2</i>	chr21:21292504-21833085
	<i>FGF13</i>	chrX:138113887-138114818
	<i>SOX3</i>	chrX:139412818-139414891
	<i>C14orf39</i>	chr14:59972427-60022517
	<i>ZNF347</i>	chr19:58334460-58354096
	<i>SETDB1</i>	chr1:149165581-149202945
	<i>CADM3</i>	chr1:157408001-157433474
	<i>CD34</i>	chr1:206126506-206151306
	<i>EGFLAM</i>	chr5:38498449-38501338
	<i>HTR1E</i>	chr6:87781772-87782869
	<i>ADCY1</i>	chr7:45580650-45729239
	<i>AUTS2</i>	chr7:68701841-69895821
	<i>ADRB3</i>	chr8:37939673-37943341
	<i>ITGA8</i>	chr10:15599094-15801776
	<i>ZNF606</i>	chr19:63180253-63206526
	<i>C2orf40</i>	chr2:106048545-106061041
	<i>FAM19A4</i>	chr3:68863607-69064401

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	<i>BARHL1</i>	chr9:134447814-134455461
	<i>FLJ21963</i>	chr12:79995989-80172990
	<i>MEGF11</i>	chr15:63978120-64333129
	<i>HS3ST3A1</i>	chr17:13339731-13445969
	<i>CLDN5</i>	chr22:17891122-17891778
	<i>LIN28</i>	chr1:26609856-26628806
	<i>TRIB2</i>	chr2:12774449-12800309
	<i>SCTR</i>	chr2:119913889-119998498
	<i>KY</i>	chr3:135801455-135852554
	<i>HIST1H4J</i>	chr6:27899882-27900237
	<i>NPTX2</i>	chr7:98086915-98094592
	<i>SDC2</i>	chr8:97575058-97693213
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	<i>SNAP25</i>	chr20:10147477-10236065
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	<i>NID1</i>	chr1:234205755-234295104
	<i>ERBB4</i>	chr2:212134732-213111597
	<i>TSP50</i>	chr3:46728610-46829068
	<i>ZFP42</i>	chr4:189160956-189161888
	<i>TLX3</i>	chr5:170668893-170671743
	<i>HOXA6</i>	chr7:27151641-27153893
	<i>CYP7B1</i>	chr8:65671083-65873902
	<i>SHC3</i>	chr9:90817889-90983502
	<i>DFNB31</i>	chr9:116280422-116307551
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	<i>SLC1A2</i>	chr11:35229328-35398100
	<i>GPC6</i>	chr13:92995517-93282343
	<i>SUMO2</i>	chr17:70675420-70690693
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	<i>TBX19</i>	chr1:166533486-166550288
	<i>ACTA1</i>	chr1:227633616-227636466
	<i>LRRN1</i>	chr3:3861326-3863476
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	<i>POU4F3</i>	chr5:145698869-145700200
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	<i>WNT3</i>	chr17:42196857-42251081
	<i>RAX</i>	chr18:55085247-55091605
	<i>CACNG8</i>	chr19:59158106-59177951
	<i>PLCB1</i>	chr20:8061296-8578636
	<i>PREX1</i>	chr20:46674200-46877827
	<i>C20orf58</i>	chr20:61342584-61356335
	<i>CLIC6</i>	chr21:34963558-35012389
	<i>ESX1</i>	chrX:103381375-103386244
	<i>FIGN</i>	chr2:164172364-164300759
	<i>OSMR</i>	chr5:38881893-38970159
	<i>NR0B1</i>	chrX:30232460-30237416
	<i>TDGF1</i>	chr3:46594217-46598956
	<i>PDE8B</i>	chr5:76542462-76758984
	<i>HOXA5</i>	chr7:27147521-27149812
	<i>EPO</i>	chr7:100156359-100159259
	<i>BC058917:118</i>	chr12:77783089-77783089
	<i>COL24A1</i>	chr1:86283399-86395034
	<i>PLD5</i>	chr1:240318895-240754621
	<i>CCK</i>	chr5:43412505-43448245
	<i>SOX14</i>	chr3:138966269-138967086
	<i>ADRB2</i>	chr5:148186349-148188381
	<i>VWC2</i>	chr7:49783803-49922684
	<i>TNFRSF10C</i>	chr8:23016379-23030895
	<i>ADARB2</i>	chr10:1218073-1769718
	<i>NKX6-2</i>	chr10:134448310-134449527
	<i>CALCB</i>	chr11:15051722-15060464
	<i>PRMT8</i>	chr12:3360776-3573399
	<i>CACNG3</i>	chr16:24174377-24281238
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	<i>MN1</i>	chr22:26474265-26517871
	<i>HSPA6</i>	chr1:159760660-159763311
	<i>CACNA1D</i>	chr3:53504071-53821532
	<i>TMEM108</i>	chr3:134581251-134597747
	<i>OSAP</i>	chr4:140406767-140420942
	<i>MEOX2</i>	chr7:15617362-15692833
	<i>PDE1C</i>	chr7:31795772-32077516
	<i>FIGNL1</i>	chr7:50480455-50482479
	<i>CNTNAP2</i>	chr7:147667486-147749019
	<i>ASAH3L</i>	chr9:19398925-19442500
	<i>ITIH5</i>	chr10:7653374-7748940

	<i>GATA3</i>	chr10:8136673-8157170
	<i>C10orf53</i>	chr10:50557690-50586962
	<i>SLC16A12</i>	chr10:91180036-91285293
	<i>EMX2</i>	chr10:119291946-119299047
	<i>UTF1</i>	chr10:134893768-134895052
	<i>SCT</i>	chr11:616313-617173
	<i>SOX21</i>	chr13:94159880-94162390
	<i>MYCBPAP</i>	chr17:45940811-45963861
	<i>IL11</i>	chr19:60567569-60573626
	<i>SCUBE1</i>	chr22:41963526-42069299
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	<i>GNB4</i>	chr3:180596570-180652065
	<i>D4S234E</i>	chr4:4440213-4471686
	<i>DDIT4L</i>	chr4:101326050-101330636
	<i>GRIA2</i>	chr4:158361844-158502254
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	<i>LY6K</i>	chr8:143778533-143782613
	<i>COL5A1</i>	chr9:136673473-136876509
	<i>WIT1</i>	chr11:32413861-32418196
	<i>FADS2</i>	chr11:61352289-61391401
	<i>COL2A1</i>	chr12:46653015-46668521
	<i>C13orf21</i>	chr13:43846159-43862587
	<i>FAM100A</i>	chr16:4598885-4604928
	<i>PTPRM</i>	chr18:7557314-8396859
	<i>THBD</i>	chr20:22976594-22978141
	<i>ANKRD53</i>	chr2:71059083-71066137
	<i>FEZF2</i>	chr3:62330387-62334230
	<i>SERP1</i>	chr3:151742470-151803705
	<i>CXCL14</i>	chr5:134934274-134942868
	<i>ANKRD6</i>	chr6:90387654-90400124
	<i>EYA4</i>	chr6:133825452-133894951
	<i>LY6H</i>	chr8:144310706-144313428
	<i>SYCP1</i>	chr1:115198978-115339513
	<i>BACH2</i>	chr6:90716596-91063182
	<i>DFNA5</i>	chr7:24704499-24764164
	<i>MMP16</i>	chr8:89150577-89408833
	<i>GDF6</i>	chr8:97223734-97242196
	<i>DMRTA1</i>	chr9:22436840-22442472
	<i>UAP1L1</i>	chr9:139091774-139098811
	<i>KCNIP2</i>	chr10:103575721-103593667
	<i>LOC643923</i>	chr11:106967681-106969159
	<i>LITAF</i>	chr16:11557642-11588290
	<i>COMMD7</i>	chr20:30754154-30795475

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	<i>EDARADD</i>	chr1:234625339-234714631
	<i>FIGLA</i>	chr2:70857950-70871283
	<i>LRPIB</i>	chr2:140705466-142605740
	<i>VEGFC</i>	chr4:177841685-177950889
	<i>CMYA5</i>	chr5:79021415-79131805
	<i>TCF21</i>	chr6:134251970-134255086
	<i>FLNC</i>	chr7:128257719-128286564
	<i>TMEM176B</i>	chr7:150119309-150129381
	<i>UNQ9433</i>	chr8:53609253-53640565
	<i>MKX</i>	chr10:28001809-28074784
	<i>ARMC4</i>	chr10:28141103-28327797
	<i>TMEM26</i>	chr10:62842659-62883214
	<i>GRID1</i>	chr10:87349292-88116230
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	<i>CALCA</i>	chr11:14945796-14950408
	<i>RAB39</i>	chr11:107304487-107339418
	<i>KCNA5</i>	chr12:5023346-5026210
	<i>ATP5G2</i>	chr12:52345211-52356779
	<i>C13orf33</i>	chr13:30378312-30397709
	<i>BIVM</i>	chr13:102249400-102291514
	<i>NFATC4</i>	chr14:23908199-23916235
	<i>KCNK10</i>	chr14:87720999-87863004
	<i>PYDC1</i>	chr16:31135581-31135850
	<i>HSF4</i>	chr16:65755845-65761349
	<i>FLJ10847</i>	chr17:19391890-19422938
	<i>FAM20A</i>	chr17:64044860-64108690
	<i>NOVA2</i>	chr19:51134631-51168497
	<i>KCNJ14</i>	chr19:53656318-53661179
	<i>SLCO4A1</i>	chr20:60744242-60774092
	<i>TIAMI</i>	chr21:31506818-31854161

\* Locations are corresponding to the UCSC Genome Database Assembly Version: May 2004 (<http://genome.ucsc.edu/index.html>). † Average values (cy5/cy3) of 10 cancerous tissues. ‡ Average values (cy5/cy3) of 10 adjacent tissues.