

Supplementary table: Gene list of promoter-associated hypermethylated genes of OSCC generated by Illumina's Genome Studio software analysis (p 0.000).<

UCSC_REFGENE_NAME	UCSC_REFGENE_ACCESSION	UCSC_CPG_ISLANDS_NAME
DFFB	NM_014704;NM_004402	chr1:3775077-3775369
DUSP1	NM_004417	chr5:172197482-172199606
FEZ2	NM_001042548;NM_005102	chr2:36824792-36825305
RWDD3	NM_001128142;NM_015485	chr1:95699725-95700142
ZNF589	NM_016089	chr3:48282425-48282817
DDAH2	NM_013974	chr6:31695894-31698245
TSC22D3	NM_004089;NM_198057;NM_001015881	chrX:106959378-106959914
LEPRE1	NM_022356;NM_001146289	chr1:43232214-43233359
ORMDL3	NM_139280	chr17:38083226-38084052
HYAL2	NM_003773;NM_033158	chr3:50359839-50360042
MUM1	NM_032853;NR_024247	chr19:1354420-1355350
KIF3C	NM_002254	chr2:26204554-26205649
EPHX2	NM_001979	chr8:27348658-27348883
GEMIN7	NM_001007269;NM_001007270;NM_024707	chr19:45594397-45595007
MIER2	NM_017550	chr19:345260-345590
MAPRE2	NM_001143826	chr18:32556654-32557883
C14orf80	NM_001134877	chr14:105956175-105958197
VEZF1	NM_007146	chr17:56064830-56066051
ZBTB4	NM_001128833;NM_020899	chr17:7382266-7382710
P16	NM_001195132;NM_058195	chr9:21967751-21994490
HYAL2	NM_003773;NM_033158	chr3:50359839-50360042
PPT2	NM_005155;NM_030651;NM_138717	chr6:32121829-32122529
TRIM65	NM_173547	chr17:73892496-73893448
TMEM120A	NM_031925	chr7:75623357-75624164
PFDN5	NM_002624;NM_145897	chr12:53689227-53689570
CCNJ	NM_001134376;NM_001134375;NM_019084	chr10:97802871-97804262
GEMIN7	NM_001007269;NM_001007270;NM_024707	chr19:45594397-45595007
NCKAP5L	NM_001037806;NR_027499;NR_027500;NR_027501	chr12:50221558-50222609
TAF4	NM_003185	chr20:60638612-60638982
C1orf52	NM_198077;NR_024113	chr1:85724701-85725524
TRAPPC5	NM_001042461;NM_174894;NM_001042462	chr19:7745490-7747714
TAF4	NM_003185	chr20:60639272-60639835
TRIM65	NM_173547	chr17:73892496-73893448