	Cosmic ID	Cosmic Coverage	Broad-Novartis Cancer Cell line Encyclopedia	GEO datasets ID	Description	Туре
CP-D	n/a		n/a	n/a		
ESO26	1503366	partial (1071 genes)	n/a	n/a		
ESO51	1503367	partial (1102 genes)	n/a	n/a		
Flo-1	1503361	partial (1053 genes)	n/a	GSE43150	Association between Hypomethylation of Noncoding Regions and Overexpression of the Inc-RNA, AFAP1-AS1, in Barrett's Esophagus	Methylation profiling by high throughput sequencing
Jh-Eso- Ad1	n/a		Expression arrays (source: Achilles)	n/a		
OACM5.1 C	1503363	partial (836 genes)	n/a	n/a		
OACP4 C	1503362	partial (621 genes)	n/a	n/a		
OE33	910549	partial (954 genes)	Expression Arrays, SNP arrays (source EACC)	GSE36841 GSE44120 GSE57130 GSE47763 GSE36133 GSE36138 GSE366728, GSE66729 GSE43150	Expression data from OE33 oesophageal adenocarcinoma tumour cells following 24 hour co-culture with human adipocytes, adipose tissue explants or control M199 medium miR-223 in esophageal adenocarcinoma carcinogenesis Combination of HDAC inhibitors and Azacytidine for Cancer Cell Selective Targeting of Esophageal Cancer Cells STAT3 expression, activity and functional consequences of STAT3 inhibition in esophageal squamous cell carcinomas and Barrett's adenocarcinomas Expression data from the Cancer Cell Line Encyclopedia (CCLE) SNP array data from the Cancer Cell Line Encyclopedia (CCLE) Gene expression data ARID1A knockdown OE33 cells versus mockOE33 cells The Landscape of Antisense Gene Expression in Human Cancer Association between Hypomethylation of Noncoding Regions and Overexpression of the Inc-RNA, AFAP1-AS1, in Barrett's Esophagus	Expression profiling by array Genome variation profiling by SNP array Expression profiling by array Expression profiling by high throughput sequencing Methylation profiling by high throughput sequencing
SK-GT-4	SK-GT-4	partial (977 genes)	n/a	GSE2144 GSE13376 GSE43150	Esophageal cell response to low pH: time course Exposure of Barrett's associated adenocarcinoma cell lines SKGT4to deoxycholic acid (DCA) Association between Hypomethylation of Noncoding Regions and Overexpression of the Inc-RNA, AFAP1-AS1, in Barrett's Esophagus	Expression profiling by array Expression profiling by array Methylation profiling by high throughput sequencing