

**Correlation coefficients of the most differentially regulated tumor genes  
and clinicopathological parametres**

Gene symbol	Postoperative survival		Tumor size		Intestinal subtype		Lymph node metastasis	
	Pearson	Spearman	Pearson	Spearman	Pearson	Spearman	Pearson	Spearman
ADA	-0.633	-0.604						
ADH1C				-0.578				
AGXT2I1	-0.684	-0.595						
ANGPTL3	-0.702	-0.656						
APLP1	-0.740	-0.635						
AQP4	-0.692	-0.582						
B3GAT1	-0.724	-0.721						
C6ORF58				-0.569				
CAPN9						-0.651		
CCDC121		-0.563						
CHIA	-0.608	-0.565	0.566					
CKM	-0.626	-0.604						
CLCNKA	-0.673	-0.611						
DPCR1						-0.572		
CLDN1	-0.58	-0.693						
FGA	-0.626	-0.677						
FLJ42875	-0.592							
GKN1					-0.573	-0.606		
KCNJ16		-0.580						
KLK6					0.575	0.574		
KLK11		-0.709						
MFSD4	-0.652							
RAP1GAP								0.574
REP15						-0.577		
RPRM	-0.680	-0.719	0.579					
SCGB2A1					-0.587			
SH3GL2	-0.732	-0.629	0.572					
SOSTDC1					-0.571	-0.594		
SOX21					-0.606	-0.596		
TMED6	-0.578	-0.616						
TRIM50	-0.708	-0.693						