Supplementary Data

Figure S1

(A) CD147 transcript levels in pancreatic cancer tissues as compared with normal pancreas according to Oncomine database analysis.

(B) Oncomine microarray data analysis of the genes coexpressed with BSG (CD147) in Segara Pancreas by Human Genome U133A Array (data incite from Clin Cancer Res. 2005 May 1;11(9):3587-96). Data was grouped by cancer types and expressed as log2 median-centered intensity.