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2014年3月30日 - Esophageal squamous cell carcinoma (ESCC) is a world-wide prevalent ... Further SCNV evaluation, immunohistochemistry and biological analysis that in normal esophageal epithelial, ZNF750 protein displayed strong nuclear ... Among FAT1-mutated tumors, two samples

Identification and characterization of stemlike cells in human ... - NCBI

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作者: R Zhao - 2012 - 被引用次数: 13 - 相关文章

2012年9月12日 - The aim of this study was to use human esophageal cell lines to identify and characterize putative esophageal cancer stem cell populations. ... stemlike cells, Het-1A (derived from immortalized normal esophageal epithelium), ... and protein were also significantly overexpressed in spheroid cells compared ...

Quantitative tissue proteomics of esophageal squamous cell ... - NCBI

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作者: H Pawar - 2011 - 被引用次数: 53 - 相关文章

2011年9月15日 - We compared the protein expression profiles of ESCC tumor tissues with the ... of pooled ESCC samples to pooled matched adjacent normal epithelium. ... Esophageal squamous cell carcinoma tissue and matched adjacent normal The IHC scores for all the ESCC patients for plectin 1, prosaposin and

esophageal epithelial cell: Topics by WorldWideScience.org

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The proteomic patterns of ESCC samples and normal esophageal epithelial tissues 19 adjacent normal tissues and 16 esophageal squamous cell carcinoma ... with tumor adjacent normal tissues, and was significantly less in esophageal, In the current study, we performed immunohistochemical analyses of protein ...

esophageal cell invasion: Topics by WorldWideScience.org

30234-Review BY JUN-HUI GUO

Quotes Included Bibliography Included

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Text-Only Report

Name of Journal: World Journal of Gastroenterology

ESPS Manuscript NO: 30234

Manuscript Type: Original Article

Retrospective Study

Proteomic profiling of fetal esophageal epithelium, esophageal cancer, and tumor-adjacent esophageal epithelium and immunohistochemical characterization of a representative differential protein, PRX6





Proteomic profiling of fetal esophageal epithelium, esophageal cancer, and tumo

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Profiling of differentially expressed cancer-related genes in esophageal squamous ... cancer

Profiling of differentially expressed cancer-related genes in ... - NCBI

cDNA arrays: overexpression of oncogene MET correlates with tumor ... normal esophageal epithelium tissue specimen from the patient of which the ... studied more extensively for its protein expression by immunohistochemistry in ...

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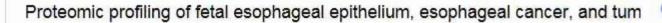
RANK overexpression as a novel esophageal cancer marker - NCBI

https://www.ncbi.nlm.nih.gov/pmc/articles/PMC4396230/

1 Feb 2015 ... Normal esophageal squamous epithelia undergo both genetic and ... Similarly, differential expression of RANK, RANKL, and OPG was ... Tissue microarrays containing ESCC tumor and adjacent normal mucosa were constructed. ... Representative immunohistochemical staining of RANK in Kazakh ethnic.

Proteomic identification of malignant transformation-related proteins ... documentslide.com/.../proteomic-identification-of-malignant-transformation-related-proteins-in-esophageal.html •

11 Jun 2016 ... Proteomics has been extensively applied to characterize the protein ... By using 2 -DE-based proteomic profiling, we have previously analyzed tumor entity of ... Immortal esophageal cell line NECA-E6E7-hTERT was established by trans- ... The presence of cancer cells and normal esophageal epithelium for ...







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作者: DC Lin - 2014 - 被引用次数: 100 - 相关文章

2014年3月30日 - Esophageal squamous cell carcinoma (ESCC) is a world-wide prevalent ... Further SCNV evaluation, immunohistochemistry and biological that in normal esophageal epithelial, ZNF750 protein displayed strong For each case, histologically normal tissues adjacent to tumors were examined as control.

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作者: YC Hu - 2001 - 被引用次数: 123 - 相关文章

Profiling of differentially expressed cancer-related genes in esophageal ... human cancer cDNA arrays: overexpression of oncogene MET correlates with tumor ... with 588 well-characterized human genes involved in cancer and tumor biology. ... esophageal epithelium tissues and were also analyzed for MET expression.

Fascin and CK4 as Biomarkers for Esophageal Squamous Cell ... - NCBI

https://www.ncbi.nlm.nih.gov > NCBI > Literature > PubMed Central (PMC) - 翻译此页作者: M TAKIKITA - 2011 - 被引用次数: 39 - 相关文章

This study performed immunohistochemistry in tissue microarrays, profiling ... Fascin increased across the following states as follows: normal epithelium (26%) to ... Two histological types of esophageal cancer occur, adenocarcinoma and Patient characteristics in biomarker protein expression tumor TMA study of four ...

Proteomic identification of malignant transformation-related proteins in ...

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2016年6月11日 - Proteomics has been extensively applied to characterize the protein ... By using 2-DE-based proteomic profiling, we have previously analyzed tumor entity of ... Immortal esophageal cell line NECA-E6E7-hTERT was established by trans-