

**Name of Journal:** *World Journal of Gastroenterology*

**Manuscript NO:** 41325

**Manuscript Type:** ORIGINAL ARTICLE

### *Basic Study*

**Prognostic value of sorting nexin 10 weak expression in stomach adenocarcinoma revealed by weighted gene co-expression network analysis**

Zhang J *et al.* Prognostic value of SNX10 in SA

### **Abstract**

#### *AIM*

To detect significant clusters of co-expressed genes associated with tumorigenesis that might help to predict stomach adenocarcinoma (SA) prognosis.

#### *METHODS*

### Match Overview

1	<b>Internet</b> 54 words crawled on 25-Jul-2018 <a href="http://www.wjgnet.com">www.wjgnet.com</a>	1%
2	<b>Crossref</b> 12 words Mengjiao Han, Dongjun Dai, Neelum Aziz Yousafzai, Falian... Wang et al. "CXXC4 activates apoptosis through up-regulatin	<1%

[All](#)[Images](#)[News](#)[Videos](#)[Shopping](#)[More](#)[Settings](#)[Tools](#)

About 35 results (0.52 seconds)

### [Integrated network analysis and logistic regression modeling identify ...](#)

<https://www.ncbi.nlm.nih.gov/pmc/articles/PMC4502639/> ▼

by V Randhawa - 2015 - Cited by 5 - Related articles

Jul 16, 2015 - Keywords: **Coexpression network analysis**, Gene module, Hub gene, Microarray, ... Unfortunately, only 35 % of cases of oral **cancer** are **detected** at the earliest stage ... developed to identify biomarkers for oral **cancer prognosis** and diagnosis [8–10]. ..... **WGCNA**, Weighted gene correlation **network analysis** ...

### [Prognostic value of tumor size in gastric cancer: an analysis of 2,379 ...](#)

<https://www.ncbi.nlm.nih.gov/pubmed/23319074>

by P Guo - 2013 - Cited by 14 - Related articles

**Prognostic value** of tumor size in **gastric cancer**: an **analysis** of 2,379 patients. ... Tumor size has been included into the staging systems of many solid tumors, such ... Multivariate **analysis revealed** that tumor size as well as age, tumor location, ...

Missing: `snx10 wgcna ee expression network`

### [\(PDF\) Integrated network analysis and logistic... - ResearchGate](#)

[https://www.researchgate.net/.../280302751\\_Integrated\\_network\\_analysis\\_and\\_logistic...](https://www.researchgate.net/.../280302751_Integrated_network_analysis_and_logistic...)

by V Randhawa - 2015 - Cited by 5 - Related articles

Jul 23, 2015 - **Gene expression** profiles were classified into **cancer** stage-related ... Figure 1: The steps involved in systems **level analysis** of data on oral .... of oral **cancer** are **detected** at the earliest stage (without .... stage is an effective **prognostic** factor, to the best of our .... **WGCNA** defined the **network** into color-coded.

[全部](#)[图片](#)[新闻](#)[视频](#)[购物](#)[更多](#)[设置](#)[工具](#)

找到约 4,490 条结果 (用时 0.57 秒)

### Weighted gene co-expression network analysis reveals potential ...

<https://www.ncbi.nlm.nih.gov/pubmed/29180012> - 翻译此页

作者: Y Li - 2018 - 被引用次数: 1 - 相关文章

2017年11月24日 - **Weighted gene co-expression network analysis reveals** potential genes involved in early metamorphosis process in sea cucumber ...

### Weighted gene co-expression network analysis in identification of ...

<https://www.ncbi.nlm.nih.gov/pubmed/28203405> ▾ 翻译此页

作者: F Tian - 2017 - 被引用次数: 6 - 相关文章

2017年1月4日 - **Weighted gene co-expression network analysis** in identification of metastasis- related genes of lung squamous cell **carcinoma** based on the ...

### Weighted gene co-expression network analysis of colorectal cancer ...

<https://www.ncbi.nlm.nih.gov/pubmed/27571956> ▾ 翻译此页

作者: B Gao - 2016 - 被引用次数: 6 - 相关文章

2016年6月30日 - **Weighted gene co-expression network analysis** of colorectal **cancer** liver ... metastatic **tumor** in the liver, gene ontology (GO) **analysis revealed** ...

### Candidate genes in gastric cancer identified by constructing a ... - PeerJ

<https://peerj.com/articles/4692/> ▾ 翻译此页

作者: J Chen - 2018 - 被引用次数: 1 - 相关文章

2018年5月4日 - Background **Gastric cancer** (GC) is one of the most common ... However, the molecular mechanisms of GC are unclear, and the **prognosis** of GC is **poor**. ... was **analyzed** by **weighted gene co-expression network analysis** ... Furthermore, the **significance** of these genes was confirmed via survival **analysis**.[Most recent papers with the keyword Gastric Cancer TCGA | Read by ...](#)

[全部](#)[图片](#)[新闻](#)[视频](#)[购物](#)[更多](#)[设置](#)[工具](#)

找到约 4,730 条结果 (用时 0.58 秒)

### Weighted gene co-expression network analysis reveals potential ...

<https://www.ncbi.nlm.nih.gov/pubmed/29180012> - 翻译此页

作者: Y Li - 2018 - 被引用次数: 1 - 相关文章

2017年11月24日 - **Weighted gene co-expression network analysis** reveals potential genes involved in early metamorphosis process in sea cucumber ...

### Weighted gene co-expression network analysis in identification of ...

<https://www.ncbi.nlm.nih.gov/pubmed/28203405> ▾ 翻译此页

作者: F Tian - 2017 - 被引用次数: 6 - 相关文章

2017年1月4日 - **Weighted gene co-expression network analysis** in identification of metastasis-related genes of lung squamous cell **carcinoma** based on the ...

### Weighted gene co-expression network analysis of colorectal cancer ...

<https://www.ncbi.nlm.nih.gov/pubmed/27571956> ▾ 翻译此页

作者: B Gao - 2016 - 被引用次数: 6 - 相关文章

2016年6月30日 - **Weighted gene co-expression network analysis** of colorectal **cancer** liver ... metastatic **tumor** in the liver, gene ontology (GO) **analysis revealed** ...

### Candidate genes in gastric cancer identified by constructing a ... - PeerJ

<https://peerj.com/articles/4692/> ▾ 翻译此页

作者: J Chen - 2018 - 被引用次数: 1 - 相关文章

2018年5月4日 - Background **Gastric cancer** (GC) is one of the most common ... However, the molecular mechanisms of GC are unclear, and the **prognosis** of GC is **poor**. ... was **analyzed** by **weighted gene co-expression network analysis** ... Furthermore, the **significance** of these genes was confirmed via survival **analysis**.

### Most recent papers with the keyword Gastric Cancer TCGA | Read by ...

<https://www.readbyqxmd.com/keyword/192164> - 翻译此页

The aim of this study was to determine novel gene **expression** signatures from the .... of hub genes with **prognostic values** in **gastric cancer** by bioinformatics **analysis**. ... **Sorting nexin-1** (SNX1) is an important functional protein in cell endocytosis ... was **analyzed** by **weighted gene co-expression**