Original Article Overexpression of PAK-1 is an independent predictor of disease recurrence in colorectal carcinoma

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Received October 7, 2015; Accepted November 22, 2015; Epub December 1, 2015; Published December 15, 2015

Abstract: Background: Colorectal carcinoma (CRC) is a significant cause of major morbidity and mortality. PAK-1 is a protein that regulates cytoskeletal dynamics and cell motility. The purpose of the present study is to investigate the relationship between PAK-1 immunoexpression and CRC progression and its validity as an independent prognostic factor. Patients and methods: Paraffin blocks of 103 primary CRCs and 37 nodal metastases were retrieved and tissue microarrays were constructed. Immunohistochemistry was performed using anti-PAK-1 antibody. Immunostaining was scored and results were analysed in relation to clinicopathological parameters. Results: PAK-1 was overexpressed in primary CRC (P<0.001). No difference between low and high expression in nodal metastasis (P=0.139). There was no difference between PAK-1 immunoexpression in primary and nodal metastasis (P=0.275). High PAK-1 immunoexpression was associated with disease recurrence (P=0.03). However, there was no association with most clinicopathological parameters. PAK-1 overexpression was detected as an independent predictor of disease recurrence (P=0.05). No association was found between PAK-1 immunoexpression and disease free survival (log-rank =1.287, P=0.257). Conclusion: PAK-1 overexpression may be involved in CRC progression and could be considered an independent predictor of disease recurrence. Further in vivo and in vitro molecular studies are needed to investigate the role of PAK-1 in colorectal carcinogenesis.

Keywords: PAK-1, immunohistochemistry, colorectal carcinoma, prognosis

Introduction

P21-activated kinase (PAK) is a group of serine/threonine kinases that were first discovered in 1994 in a screen for proteins that interact with the small G-proteins Rac1 and Cdc42 [1]. PAKs contribute in numerous cellular signalling pathways [2]. PAKs family is sub-grouped according to structural and architectural characteristics into two categories; group I; including PAK 1, PAK 2, and PAK 3. and group II including PAK 4. PAK 5. and PAK 6. The distribution of PAKs among normal tissues is variable with PAK 1 particularly expressed in the brain, muscle and spleen, PAK 2 in variable tissues, PAK 4 in prostate, testis, colon, and finally PAK 3 and PAK 5 in brain tissue. Group I PAKs are activated by growth factors and extracellular signals through GTPase-dependent and independent mechanisms [2-4].

PAK-1 controls growth factor signals responsible for cell proliferation through the regulation of cell-cycle progression and mitotic activity [5-8]. It plays a central role in stimulating cell motility through the organization of actin cytoskeleton, cell shape and adhesion dynamics which are mandatory for invasion and metastasis [7]. Moreover, PAK-1 has a significant part in regulating cell death and survival signalling through controlling apoptosis [9]. Overexpression of PAK-1 is reported to increase migration potential and abnormal mitotic activity in variable carcinomas such as breast, ovary, thyroid and colon [10-12]. Overexpression of PAK-1 was associated with the development of mammary cell hyperplasia in animal models through

(n=103)		
Parameter		Number (%)
Age	<60 years	60 (58.3%)
	≥60 years	43 (41.7%)
Sex	Male	57 (55.3%)
	Female	46 (44.7%)
Grade	Well-differentiated	18 (17.5%)
	Moderately-differentiated	71 (68.9%)
	Poorly-differentiated	14 (13.6%)
Tumour location	Right colon	34 (33%)
	Left colon	61 (59.2%)
	Rectum	8 (7.8%)
Tumour size	<5 cm	42 (40.8%)
	≥5 cm	61 (59.2%)
Primary tumour	T1	1 (1%)
	T2	15 (14.6%)
	ТЗ	78 (75.7%)
	Τ4	9 (8.7%)
Nodal metastasis	Negative	43 (41.7%)
	Positive	58 (56.3%)
	Cannot be assessed	2 (1.9%)
Lymphovascular invasion	Positive	85 (82.5%)
	Negative	18 (17.5%)
Margin status	Free	99 (96.1%)
	Involved	4 (3.9%)
Distant metastasis	Negative	73 (72.3%)
	Positiv e	28 (27.7%)
	Not available	2 (1.9%)
Recurrence	No recurrence	77 (74.8%)
	Recurrence	26 (25.2%)
Survival	Alive	61 (59.2%)
	Dead	25 (24.3%)
	Not available	17 (16.5%)

 Table 1. Clinicopathological parameters of CRC patients

 (n=103)

T1: Tumour invades submucosa; T2: Tumour invades muscularis propria; T3: Tumour invades through the muscularis propria into the subserosa or into non-peritonealised pericolic or perirectal tissues; T4: Tumour directly invades other organs or structures, and/or perforates visceral peritoneum.

its correlation to increased cyclin D1 promoter activity, which in turn could be related to increased progression of breast cancer [13]. PAK-1 was included into HER-2 pathway controlling invasiveness of breast cancer cells in cooperation with other oncogenes. They showed that loss of PAK-1 leads to reduced expression of beta-catenin and its target genes with subsequent prolonged survival. These results suggest a new therapeutic strategy for HER-2 positive breast cancer patients [14]. Increased PAK-1 expression in the invasive The purpose of the present study is to investigate the relationship between PAK-1 immunoexpression and CRC progression and its validity as an independent prognostic factor.

Materials and methods

Patients

Hundred and three patients diagnosed as CRC constituted the material of the present study. Patients' information was gathered from the records of the Pathology Department in King Abdulaziz University, Jeddah, Saudi Arabia. Paraffin blocks of patients were sliced and stained routinely with haematoxylin and eosin to evaluate histopathological characteristics of the tumours as well as for histological grading and staging (following the AJCC staging system) [15]. For evaluation of tumour invasion, tumour stage was used as an indicator. Clinicopathological parameters of all patients as age, sex, tumour site and size, histological type, grade and stage as well as lymph node and safety margin status, and presence of distant metastasis along with follow up results were collected from the patient's medical records after obtaining the formal ethical approval. Cliniopathological data is shown in

Table 1. The study was approved bythe Research Committee of the Biomedi-cal Ethics Unit, Faculty of Medicine, KingAbdulaziz University, Jeddah, Saudi Arabia.

Tissue microarray construction

A tissue microarray was constructed from 103 primary CRC and 37 lymph nodes showing associated tumour metastasis. TMAs were constructed using an automated tissue arrayer (MASTER 3D HISTECH). Donor Paraffin blocks used in this study was retrieved from surgical Table 2. Categories of PAK-1 immunoexpression in primary tumours and nodal metastases

	Primary	Nodal	
	tumour	Metastasis	P value
	(<i>n</i> =103)	(n=37)	
Low expression	29 (28.2%)	14 (37.8%)	0.275 °
High expression	74 (71.8%)	23 (62.2%)	
P value	<0.001*	0.139*	

*One sample non-parametric chi-square test; *Mann-Whitney test; Low expression: 0+1; High Expression: 2+3.

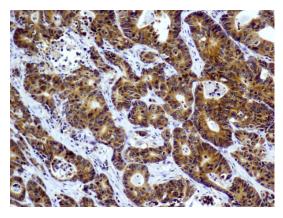


Figure 1. PAK-1 cytoplasmic immunostaining in a well differentiated primary CRC. The expression pattern of PAK-1 protein is strong (+3) and evident in both basal and apical portions of malignant cells (100×).

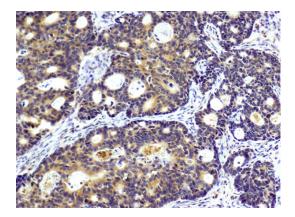


Figure 2. Moderately differentiated colorectal carcinoma showing borderline/moderate (+2) diffuse cytoplasmic expression pattern for PAK-1 (100×).

removed specimens. From selected paraffin blocks, a cylinder of tissue 1 mm in diameter was cut with a TMA instrument and inserted into a new recipient paraffin block. Serial 4-µm sections were then cut from the TMA paraffin blocks and placed on positively charged slides for immunohistochemical staining.

Immunohistochemistry

Immunohistochemical staining was done using a primary anti-p21-activated kinase-1 rabbit polyclonal antibody at a dilution of 1:100 (DakoCytomation Norden A/S, Glostrup, Denmark). Immunostaining was completed by using an automatic immunostainer (Ventana Bench Mark XT, Ventana Inc., Tucson, AZ). Positive controls were used consisting of CRC specimens previously demonstrated to stain with this antibody. Tris-buffered saline in place of the primary antibody was utilized as a negative control.

Interpretation of immunohistochemical staining

PAK-1 positivity was considered when a yellow to brown staining was found in tumour cell cytoplasm and/or nuclei. Immunostaining was evaluated semiquantitatively according to staining intensity on a scale from 0 to 3, with grade 0: negative, 1: weak, 2: moderate and 3: intense. When dichotomized for statistical analysis; negative (-) and weak (+) staining were identified as low expression, while moderate (++) and intense (+++) staining were included in high expression category [13]. Two pathologists (EE and DQ) scored immunostaining independent of clinicopathological data.

Statistical analysis

Difference between two groups of patients was tested by using Mann Whitney test. To test association procedure in three groups of patients on one independent variable the Kruskal Wallis test was used. Non-parametric chi-square was used to test variance along one variable. Multivariate logistic regression analysis was used to predict lymph node metastasis, disease recurrence, lymphovascular invasion, and distant metastasis in relation immunoexpression of PAK-1. Estimated odds ratio (exponential [16]), 95% confidence interval (CI) for exp (B), and significance denoted for each analysis. Tumour stages were dichotomised in two categories; category 1 (Stage 1+2)=superficial invasion and category 2 (Stage 3+4)=deep invasion. The Kaplan-Meier strategy was uti-

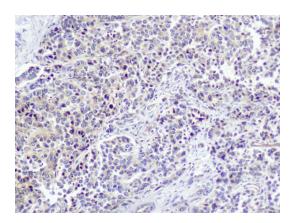


Figure 3. Poorly differentiated colorectal carcinoma showing weak (+1) diffuse cytoplasmic expression pattern for PAK-1 (100×).

 Table 3. Relation of PAK-1 immunoexpression

 to clinicopathological parameters

Parameter	P value
Age	0.625°
Sex	0.196°
Grade	0.826*
Tumour location	0.686*
Tumour size	0.714°
Depth of invasion (pT)	0.371*
Nodal metastasis	0.878*
Distant metastasis	0.611°
Lymphovascular invasion	0.235°
Margin status	0.462°
Recurrence	0.03°

*Kruskal-Wallis Test; [©]Mann-Whitney test.

lised to ascertain the survival probabilities where the Log Rank test was used to look at the contrast between survivals. The end-point for patients was death from tumour (diseasefree). Disease-free survival (DFS) was calculated as the time from diagnosis to the appearance of recurrent disease (or date last seen disease-free). Statistical procedures were performed using SPSS Release 16.0. Statistical significance was determined at *P* value of \leq 0.05 and was 2-sided.

Results

PAK-1 immunoexpression profile

Homogenous yellow-brown PAK-1 immunostaining was shown exclusively in the cytoplasm of malignant cells and in the occasional inflammatory cells. Interestingly, no nuclear staining was noted. Seventy four CRCs (71.8%) showed high PAK-1 expression while 29 CRCs (28.2%) showed low PAK-1 expression; PAK-1 was overexpressed in primary tumours (P<0.001). In nodal metastasis, there was no distinction between low and high expression (P=0.139). There was no statistically significant difference between PAK-1 immunoexpression in primary and nodal metastasis (P=0.275) (**Table 2**). **Figures 1-3** demonstrate the immunostaining expression patterns of PAK-1 protein in different primary tumours.

Relationship between PAK-1 expression and clinicopathological features of CRCs

There was no significant association between PAK-1 immuno expression and age, sex, degree of differentiation, depth of tumour invasion, stage, nodal metastasis, lymphovascular invasion, status of surgical resection margins and distant metastasis. However, overexpression of PAK-1 in CRCs is associated with disease recurrence (P=0.03) (Table 3).

Survival analysis

In Kaplan-Meier survival analysis, at 60 month (5 years) follow up time for example, **Figure 4** showed that PAK-1 expression patterns revealed an association with disease recurrence in that 50% of patients who developed tumor recurrence had high PAK-1 expression pattern of original tumor, as contrasted to only 20% patients who experienced disease recurrence had low PAK-1 expression pattern. In other words, there was a clear trend of difference in DFS between patients with tumors with low PAK-1 expression patterns (longer DFS) and those tumors with high PAK-1 expression patterns (shorter DFS), even though it was not statistically significant (P<0.2, log rank).

Relationship between PAK-1 expression and prognosis

Regression analysis for cytoplasmic immunoexpression revealed that PAK-1 overexpression was an independent predictor of disease recurrence (P=0.05) (**Table 4**). PAK-1 immunoexpression was not related to survival (log-rank =1.287, P=0.257) (**Figure 4**).

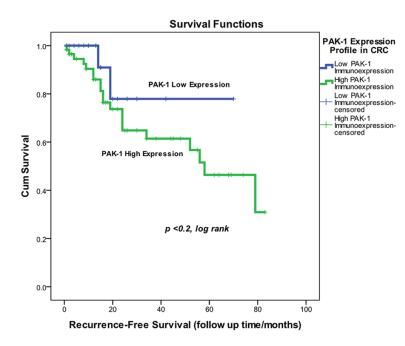


Figure 4. Kaplan-Meier curve showing the recurrence free survival outcomes of patients with low and high PAK-1 expression patterns (log-rank=1.16, *P*=0.28).

Table 4. Regression analysis for cytoplasmic PAK-1 immunoexpres-	
sion	

Variable	Exp (B)	95% CI for exp (B)	P value
Nodal Metastasis	1.180	0.448-3.106	0.738
Distant metastasis	1.223	0.448-3.338	0.694
Tumour invasion	0.614	0.182-2.079	0.434
Lymphovascular invasion	0.606	0.147-2.491	0.487
Margin status	2.567	0.299-22.035	0.390
Recurrence	0.269	0.072-1.006	0.05

Discussion

CRC arises and progresses as a result of cumulative genetic and epigenetic changes in tumour cells [17]. Detection of novel prognostic molecules involved in CRC is required to develop new targeted therapy for CRC. PAK-1 is claimed to be implicated as central player in transformation of normal cell into a malignant one through diverse mechanisms including alteration of cell motility, induction of abnormal cell proliferation and escape from apoptotic signals [3-6, 18]. PAK-1 was found to be overexpressed in a wide variety of cancers including; brain [19], breast [20], liver [21], kidney [22], bladder [23], lung [9], ovary [24], and T-cell lymphoma [25]. Previous reports adopted PAK-1 as a promising potential therapeutic agent in thyroid carcinoma, melanoma and gastric carcinoma [26-28].

In CRC, expression of PAK1 increases with progression through the adenoma to carcinoma sequence, with the most dramatic increases in invasive and metastatic CRC [29]. In the present cohort, the aim was to explore the relationship between PAK-1 overexpression and CRC progression and its validity as independent prognostic factor. PAK-1 was found to be overexpressed in CRCs. This finding is comparable to previous studies [12, 29-31]. In the current study, there was no statistically significant difference between PAK-1 expression in primary tumours and lymph node deposits. This means that PAK-1 may be involved in the progression of nodal metastasis in the same way like in primary tumour. This notion may be supported when knockdown of PAK1 controls growth and metastasis of CRC cell lines and mice models with liver metastasis [32]. In the current study, PAK-1 overexpression is significantly associated with disease recurrence. This further

supports the suggested role of PAK-1 in tumour cells proliferation, survival, and migration. In vitro evidence came from PAK-1 gene transfection which increases PAK-1 expression in SW480 cell line and augmented the invasiveness and metastasis of CRC cells [33].

On the other hand, there is no significant association with other clinicopathological factors. This finding is contradictory to those of Li et al. who reported that PAK-1 activity is associated with depth of invasion, lymph node metastasis, distant metastasis, tumour grade and tumour stage [34].

In the present study, overexpression of PAK-1 is found to be an independent predictor of disease recurrence. Comparably, Carter [29] and

others [35, 36] concluded that PAK-1 expression is significantly increasing with tumour progression and metastasis supporting its functional role in CRC invasiveness and motility. In the current study, there was a clear trend of association, even though it was not statistically significant, between PAK-1 overexpression in primary CRCs and patients' survival i.e., in other words, there was a clear drift of difference in DFS between patients with tumors with low PAK-1 expression patterns (longer DFS) and those patients with tumors of high PAK-1 expression patterns (shorter DFS). This is in conformity with study by Li, L.H., et al. [34] who showed the association between PAK-1 expression and phosphorylation with patient's survival.

Previous reports supported some molecular mechanisms involving PAK-1 role in CRC pathogenesis. PAK-1 is implicated in proliferation, survival, migration, and VEGF secretion of CRC cells which had Ras, PI3K, and Apc mutations. Also, PAK-1 knockdown suppressed growth, survival and migration of CRC cell lines through inactivation of ERK and AKT, the downstream targets of Ras [35, 37-39]. This observation is consistent with the report by Li and co-workers that PAK-1 regulates CRC metastasis through ERK-dependent phosphorylation of FAK [34]. PAK-1 was found to have a stimulatory effect on HIF-1a expression and VEGF secretion by CRC cells in response to hypoxia provide a molecular basis for the stimulation by PAK1 of CRC survival and metastasis [40]. Others found that of PAK-1 downregulation in CRC cell lines reduced beta-catenin levels and cell proliferation via directly phosphorylated beta-catenin [12]. According to the current findings, PAK-1 is an expected as a target for therapy of CRC. Some PAK-1 inhibitors are being tried. Pitts et al., established that this novel agent demonstrates activity against preclinical CRC models cell lines with a range of molecular aberrations [41]. This PAK inhibited CRC cell lines by inhibiting proliferation and migration.

Limitations of the study include the lack of tissues from normal and dysplastic colonic mucosa and short survival period of latest patients included in the study.

Conclusion

In Conclusion, our observations along with previous data supported that PAK-1 overexpression is involved in CRC invasiveness and disease recurrence. PAK-1 overexpression was also an independent prognostic predictor of disease recurrence. Subsequently, targeting PAK-1 as therapy in CRC is promising. Further studies are needed to investigate molecular mechanistic role of PAK-1 in human tumorigenesis. Also testing PAK-1 inhibitors as potential therapeutic target for interruption of tumour cell proliferation and invasion is required to be investigated intensively.

Acknowledgements

This project was funded by the National Plan for Science, Technology and Innovation (MAARIFAH) - King Abdulaziz City for Science and Technology - the Kingdom of Saudi Arabia - award number (11-BIO1524-O3). The authors also, acknowledge with thanks Science and Technology Unit, King Abdulaziz University for technical support.

Disclosure of conflict of interest

None.

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