



A novel nanoprobe-based assay for detecting K-ras mutations in plasma and stool samples from patients with pancreatic cancer: value in diagnosis and prognosis evaluation

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Background: To develop a novel nanoprobe-based assay for detecting K-ras mutations in plasma and stool samples from patients with pancreatic cancer, and assess its value in the diagnosis and prognosis evaluation of this malignancy.

Methods: Fifty-eight pancreatic cancer, 18 chronic pancreatitis, 7 pancreatic cystadenoma, 5 pancreatic cyst, 2 solid pseudopapillary tumor of the pancreas patients, who were treated at the Second Affiliated Hospital of Jiaxing University from June 2013 to October 2015. Thirty-one healthy volunteers were as normal controls. Blood and stool samples were collected from these subjects to detect the K-ras mutation status using a novel nanoprobe-based assay, assess its diagnostic accuracy and analysis the relation between the clinicopathological characteristics and survival in pancreatic cancer.

Results: The detection rates of K-ras mutations in stool and plasma samples from patients with pancreatic cancer were 79.3% and 43.1%, respectively, both of which were significantly higher than those for patients with benign pancreatic diseases (15.6% and 6.3%, respectively; $P < 0.05$) and normal controls (0% for both; $P < 0.05$). The sensitivity and specificity of stool K-ras mutation status alone, plasma K-ras mutation status alone, and the combination of both in the diagnosis of pancreatic cancer were 79.3% and 84.4%, 43.1% and 93.8%, and 86.2% and 96.9%, respectively. Plasma K-ras mutation status was significantly associated with 1-year survival and TNM stage in patients with pancreatic cancer ($P < 0.05$).

Conclusions: The developed novel nanoprobe-based assay allows for the detection of K-ras mutations in trace amounts of plasma and stool DNA.

Keywords: Pancreatic cancer; K-ras; gene mutation; magnetic nanoparticle probe; diagnosis and prognosis

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Introduction

Pancreatic cancer is a kind of highly malignant tumor of the digestive system that has a very poor prognosis, with a 5-year survival rate $< 5\%$ (1). In recent years, the incidence of pancreatic cancer has exhibited a rising trend. According

to the statistics reported by the International Agency for Research on Cancer (2), 3.38 million new cases of pancreatic cancer were diagnosed worldwide in 2012, with a mortality-to-incidence ratio of 0.98. The lack of specific clinical manifestations and reliable tumor biomarkers makes the early diagnosis of pancreatic cancer difficult. As

a consequence, more than half of cases are diagnosed at an advanced stage, and only 15–20% have the opportunity for radical surgery (3), which is presently the only cure for pancreatic cancer. Therefore, early diagnosis is extremely important for improving the prognosis of patients with pancreatic cancer, and at present, there is an urgent need to find a reliable screening tool to improve the early diagnosis of this aggressive malignancy.

The development, invasion and metastasis of pancreatic cancer involve the dysregulation of many oncogenes and tumor suppressor genes (4,5), of which *K-ras* has been the most thoroughly studied due to the nearly ubiquitous presence of *K-ras* mutations in this malignancy (6,7). Since *K-ras* mutations are of paramount importance for the initiation of pancreatic cancer (8), the detection of *K-ras* mutations has been regarded as an important tool for the auxiliary diagnosis and prognostic evaluation of pancreatic cancer (9,10). *K-ras* mutations can be detected in different types of clinical samples, including pancreatic tissue, pancreatic duct brushings, duodenal aspirates, pancreatic juice, bile, plasma, and stool (9). Invasive techniques for obtaining samples such as pancreatic juice or pancreatic tissue are certainly inappropriate for cancer screening, whereas the non-invasive testing of plasma and stool may be applicable for both the diagnosis of symptomatic patients and early screening, especially in populations (11).

Many methods for detecting *K-ras* mutations in plasma and stool samples have been developed, such as polymerase chain reaction (PCR)-restriction fragment length polymorphism (12) and peptide nucleic acid clamp PCR (13). However, these techniques are associated with drawbacks of being poorly reproducible, labor-intensive, time-consuming, or expensive, which make these difficult to be used for screening and the early diagnosis of pancreatic cancer (14,15). The advent of nanotechnology has allowed for the development of nanoprobe appropriate for biomolecular measurements (16), and nanoparticles tagged with short DNA segments can be used to detect the genetic sequence in a sample, improving the diagnostic accuracy in biological samples that contain trace amounts of target DNA (17,18). Thus, the combination of nanoprobe for the specific recognition of *K-ras* mutations with conventional PCR may improve the detection of *K-ras* mutations in plasma and stool samples obtained from patients with pancreatic cancer.

In the present study, we designed novel fluorescent nanoprobe that target *K-ras* G12V and G13D mutations,

which are the most abundant mutation types in pancreatic cancer (6,7), and determined the diagnostic accuracy of this novel nanoprobe-based assay in the detection of *K-ras* mutations in plasma and stool samples obtained from patients with pancreatic cancer, subjects with benign pancreatic diseases, and normal controls. Furthermore, we also assessed the correlation of nanoprobe-based plasma and stool *K-ras* mutation testing with the clinicopathological characteristics and survival of pancreatic cancer patients.

Methods

Patients

A total of 121 patients (64 men and 57 women) treated at the Second Affiliated Hospital of Jiaxing University from June 2013 to October 2015 were included in this study. Among these patients, 58 had pancreatic ductal adenocarcinoma, 18 had chronic pancreatitis, 7 had pancreatic cystadenoma, 5 had pancreatic cyst, and 2 had solid pseudopapillary tumor of the pancreas. All diseases were diagnosed pathologically or by endoscopic ultrasonography-guided fine needle aspiration. In addition, 31 volunteers who underwent a physical examination and exhibited no abnormality were included as normal controls. Informed consent was obtained from all participants, and the study protocol was approved by the Ethics Committee of the Second Affiliated Hospital of Jiaxing University (No. L2015-28).

Sample collection and DNA extraction

Stool specimens were collected, frozen, and stored at -80°C . After stool (200 mg) was processed with TEN and starch, fecal DNA was prepared using the phenol-chloroform method and purified using a DNA purification kit (QIAGEN, Frankfurt, Germany). Then, peripheral blood samples (5 mL) were taken and centrifuged to collect the supernatant as plasma samples, which were preserved at -80°C for further use. Plasma DNA was prepared using a magnetic bead DNA extraction kit (Omega, Norcross, USA). The concentration of the prepared DNA was determined using a NanoDrop 1000 spectrophotometer (NanoDrop, Wilmington, DE, USA), and the $\text{OD}_{260}/\text{OD}_{280}$ ratio was calculated to ensure that the purity of the DNA was good ($\text{OD}_{260}/\text{OD}_{280}$ ratio = 1.6–1.9). The prepared DNA was stored at -80°C .

Nanoparticle probe preparation

Magnetic nanoparticles were prepared through the hydrothermal synthesis method. Briefly, 0.01 M of salicylic acid solution was added to a triangle flask in an argon environment. After adjusting the pH value to 11.0 with NaOH solution, Fe₃O₄ and Fe₂O₃ were added at a Fe ion concentration ratio of 2:1. After fully mixing and heating to 90 °C for 4 hours, the mixture was centrifuged three times at 4,000 rpm to purify the magnetic nanoparticles. After that, the magnetic nanoparticles were observed using a transmission electron microscope, in order to ensure that these were uniform in diameter (~30 nm). Next, the prepared magnetic nanoparticles were adjusted to a concentration of 1 µM with DEPC treated water, mixed with 10 µM of K-*ras* G12V or G13D mutation-specific capture probe in the presence of 1-ethyl-3-(3-dimethylaminopropyl) carbodiimide hydrochloride (Sigma-Aldrich; Merck KGaA, Darmstadt, Germany), and purified in a magnetic field. This was followed by redispersion in Tris-EDTA buffer and the addition of a cross-linker (50 µM of EDC) to react for 2 hours. Then, the magnetic nanoparticles were collected by magnetic separation and re-suspended in TE.

K-ras mutation detection

K-*ras* mutation detection was performed by fluorescent quantitative real-time PCR in a 20-µL reaction system containing 20 ng of sample DNA, 0.9 mmol/L of each primer, 0.9 mmol/L of inhibitor, and 0.20 mmol/L of nanoparticle probe on an ABI 7500 real-time PCR cycler. The PCR cycling parameters were pre-denaturation at 95 °C for 10 minutes and 40 cycles of 95 °C for 20 seconds and 60 °C for 45 seconds. The probe, primers, and blockers used were as follows: K-*ras* capture probe, 5'-CTCTATTGTTGGATCATATTCGTCCACAA AATGATTCTGAATTA-3'; G12V forward primer, 5'-ACTTGTGGTAGTTGGACCT-3' and reverse primer, 5'-TAACTTGAAACCCAAGGTAC-3'; blocker, CCTACGCCACCAGCT (with 4 pentabases); G13D forward primer, 5'-GTTCTAATATAGTCACATT TTCATTATTTTATTATAAAGC-3' and reverse primer, 5'-GTCAAGGCACTCTTGCCTAGG-3'; blocker, CTTGCCTACGCCACCA (with 4 pentabases); β-actin forward primer, 5'-CTCCAT CCTGGCCTCGCTGT-3' and reverse primer, 5'-GCTGTCACCTTCACCGTTCC-3'. Samples with a

Ct value of 21–37 were regarded as having a mutant allele, while samples with a Ct value of 15–20 were regarded as having a wild-type allele.

Statistical analysis

Statistical analyses were performed using SPSS 19.0 statistical software. The sensitivity, specificity, positive predictive value, and negative predictive value of the nanoprobe-based plasma and stool K-*ras* gene detection, alone or in combination, in the diagnosis of pancreatic cancer were calculated. Numerical data, which are expressed as mean ± standard deviation, were compared using the *t*-test. Categorical data, which are expressed as percentages, were compared using the χ^2 test. Survival data were computed using the Kaplan-Meier method and compared using the Log-rank test. Paired dichotomous data were compared using the McNemar test. P values <0.05 were considered statistically significant.

Results

Nanoprobe-based K-ras G12V and G13D mutation detection in plasma and stool samples

Table 1 shows the K-*ras* G12V and G13D mutation status detected in stool and plasma samples from different groups of subjects. The detection rates of K-*ras* mutations in stool and plasma samples from patients with pancreatic cancer were 79.3% and 43.1%, respectively, and G12V was the major mutation type (40/46 and 21/25, respectively). The detection rates of K-*ras* mutations in stool and plasma samples from patients with pancreatic cancer were significantly higher than those from patients with benign pancreatic diseases (15.6% and 6.3%, respectively; P<0.05) and normal controls (0% for both; P<0.05). K-*ras* mutations were detected in both plasma and stool samples from 21 patients with pancreatic cancer and one patient with benign pancreatic disease.

Diagnostic value of nanoprobe-based plasma and stool K-ras mutation testing in patients with pancreatic cancer

As shown in Table 2, stool K-*ras* mutation status detected by nanoprobe-based assay had sensitivity and specificity of 79.3% and 84.4%, respectively, in the diagnosis of pancreatic cancer, and the corresponding values for plasma K-*ras* mutation status were 43.1% and 93.8%, respectively.

Table 1 K-ras G12V and G13D mutations in plasma and stool samples from different groups of subjects

Group	Cases (n)	Stool				Plasma			
		(G12V + G13D)/ G13D (n)	Mutation rate (%)	χ^2	P value	(G12V + G13D)/ G13D (n)	Mutation rate (%)	χ^2	P value
Pancreatic cancer	58	46/40	79.3	–	–	25/21	43.1	–	–
Benign pancreatic diseases	32	5/3	15.6	34.06	0.00*	2/2	6.3	13.34	0.00*
Normal controls	31	0/0	0	50.89	0.00*	0/0	0	18.58	0.00*

*, P<0.05, vs. pancreatic cancer group.

Table 2 Diagnostic value of K-ras mutation status in plasma and stool in pancreatic cancer

Group	Sensitivity (%)	Specificity (%)	PPV (%)	NPV (%)	P value
Stool	79.3	84.4	90.2	69.2	0.000*
Plasma	43.1	93.8	92.6	47.6	0.000*
Stool + plasma	86.2	96.9	98.0	79.5	–

*, P<0.05, vs. stool + plasma. PPV, positive predictive value; NPV, negative predictive value.

When combining the K-ras mutation status in stool and plasma together, the sensitivity and specificity rose to 86.2% and 96.9%, respectively.

Relationship of plasma and stool K-ras mutation status detected by nanoprobe-based assay with the clinicopathological characteristics of pancreatic cancer

We next analyzed the relationship between nanoprobe-based plasma and stool K-ras mutation testing and the clinicopathological characteristics of pancreatic cancer, including gender, age, the presence of clinical symptoms, tumor location, tumor size, tumor differentiation, carbohydrate antigen 19-9 (CA19-9) level, carcinoembryonic antigen (CEA) level, and TNM stage. As shown in Table 3, the K-ras mutation status in stool had no significant association with these clinicopathological characteristics. However, K-ras mutation status in plasma was significantly associated with the TNM stage, although this had no significant association with other clinicopathological characteristics (Table 4).

Relationship of plasma and stool K-ras mutation status detected by nanoprobe-based assay with the survival of patients with pancreatic cancer

All patients were followed by telephone or at the outpatient

clinic. The duration of the follow-up period ranged from 2 to 26 months, with a mean value of 7.3 months. None of the patients were lost to follow-up. There was no significant difference in either the 6-month or 1-year survival rate between patients with and without K-ras mutations in stool (78.8% vs. 81.5%, P=0.884; 41.2% vs. 45.3%, P=0.698) (Figure 1A,B). However, the 1-year survival rate was significantly lower in patients with K-ras mutations in plasma than in those without (24.5% vs. 58.1%, P=0.030), although there was no significant difference in the 6-month survival rate (73.6% vs. 83.7%, P=0.388) (Figure 1C,D).

Discussion

In the present study, we assessed the diagnostic value of the plasma and stool K-ras mutation status detected by a nanoprobe-based assay, and analyzed the association of K-ras mutations in plasma and stool samples with clinicopathological characteristics and survival in patients with pancreatic cancer. It was found that K-ras mutations could be detected in a significant number of patients. However, these mutations could only be detected in a minority of patients with benign pancreatic diseases. The combination of K-ras mutation status in stool and plasma resulted in satisfactory sensitivity and specificity for the diagnosis of pancreatic cancer (86.2% and 96.9%, respectively). In addition, it was noted that the K-ras

Table 3 Relationship between *K-ras* mutation status in stool and clinicopathological characteristics of pancreatic cancer

Variable	K-ras mutation status in stool		<i>t</i> or χ^2	P value
	Yes	No		
Gender			1.353	0.245
Male	22	8		
Female	24	4		
Age, mean \pm SD (years)	66.15 \pm 9.78	64.58 \pm 9.10	0.493	0.624
Clinical symptoms			0.013	0.910
Yes	26	7		
No	20	5		
Tumor location			0.293	0.588
Head and neck	27	6		
Body and tail	19	6		
Tumor size (cm)	4.16 \pm 0.96	3.98 \pm 1.10	0.554	0.582
Differentiation			0.589	0.443
Well/moderate	25	8		
Low	21	4		
CA19-9 (U/L)			0.000	1.000
≥ 37	29	7		
<37	17	5		
CEA (U/L)			0.001	0.972
≥ 5	18	4		
<5	28	8		
TNM stage			0.013	0.910
I + II	20	5		
III + IV	26	7		

mutation status in plasma was significantly associated with the TNM stage and 1-year survival, although this is not the case for *K-ras* mutation status in stool.

Since plasma DNA, in which *K-ras* mutations were detected, is present at extremely low levels (19), their detection requires a highly sensitive system. The nanoprobe-based assay system used in the present study combines the use of fluorescent nanoprobe and the technique of real-time PCR. On one hand, nanoparticles have a high surface area-to-volume ratio, which allows for more target molecules to be attached to a nanoparticle, greatly improving the diagnostic sensitivity (17,18). On the other hand, real-time PCR can produce quantitative

data that spans 7–8 log orders of magnitude, and thereby has an extremely high sensitivity (20). Therefore, the investigators consider that the combination of both can meet the requirement of high sensitivity in detecting a target mutation from trace amounts of DNA.

The reported sensitivities of stool *K-ras* mutation detection for the diagnosis of pancreatic cancer vary greatly from 0% to 80%, and the specificities are lower than 70% (21). In the present study, stool *K-ras* mutation testing had a sensitivity of 79.3% and a specificity of 84.4% in the diagnosis of pancreatic cancer, indicating a moderate diagnostic accuracy. Thus, nanoprobe-based stool *K-ras* mutation testing alone can be used for the screening of

Table 4 Relationship between K-ras mutation status in plasma and clinicopathological characteristics of pancreatic cancer

Variable	K-ras mutation status in plasma		<i>t</i> or χ^2	P value
	Yes	No		
Gender			0.322	0.571
Male	14	16		
Female	11	17		
Age, mean \pm SD (years)	66.80 \pm 9.38	65.09 \pm 9.81	0.658	0.513
Clinical symptoms			0.904	0.342
Yes	16	17		
No	9	16		
Tumor location			0.173	0.678
Head and neck	15	18		
Body and tail	10	15		
Tumor size, mean \pm SD (cm)	3.97 \pm 1.00	4.23 \pm 0.97	-0.969	0.337
Differentiation			0.173	0.678
Well/moderate	15	18		
Low	10	15		
CA19-9 (U/L)			0.070	0.792
≥ 37	16	20		
<37	9	13		
CEA (U/L)			0.080	0.777
≥ 5	10	12		
<5	15	21		
TNM stage			6.539	0.011*
I + II	6	19		
III + IV	19	14		

*, P<0.05, vs. stool + plasma.

pancreatic cancer. In contrast, plasma K-ras mutation detection had a very low sensitivity of 43.1% and a high specificity of 93.8%. This result is similar to the finding of a previous meta-analysis, which reported that plasma K-ras mutation testing had a pooled sensitivity and specificity of 37% and 91%, respectively (22), suggesting that plasma K-ras mutation testing alone, due to its low sensitivity, is only suitable for auxiliary diagnosis to improve diagnostic accuracy. However, when plasma and stool K-ras mutations were simultaneously detected, the sensitivity and specificity rose to 86.2% and 96.9%, respectively.

K-ras mutations could be detected in stool and plasma

samples from a small portion of patients with benign pancreatic diseases (15.6% and 6.3%, respectively). This result suggests that the sensitivity and specificity of nanoprobe-based K-ras mutation testing will inevitably decrease in clinical settings, where the differential diagnosis between pancreatic cancer and benign pancreatic diseases is common. However, considering that benign pancreatic diseases (especially pancreatic epithelial cell hyperplasia) that carry K-ras alterations may be a premalignant condition (11), the detection of K-ras mutations in stool and plasma may identify patients at risk of developing pancreatic cancer.

Many studies have assessed the significance of K-ras

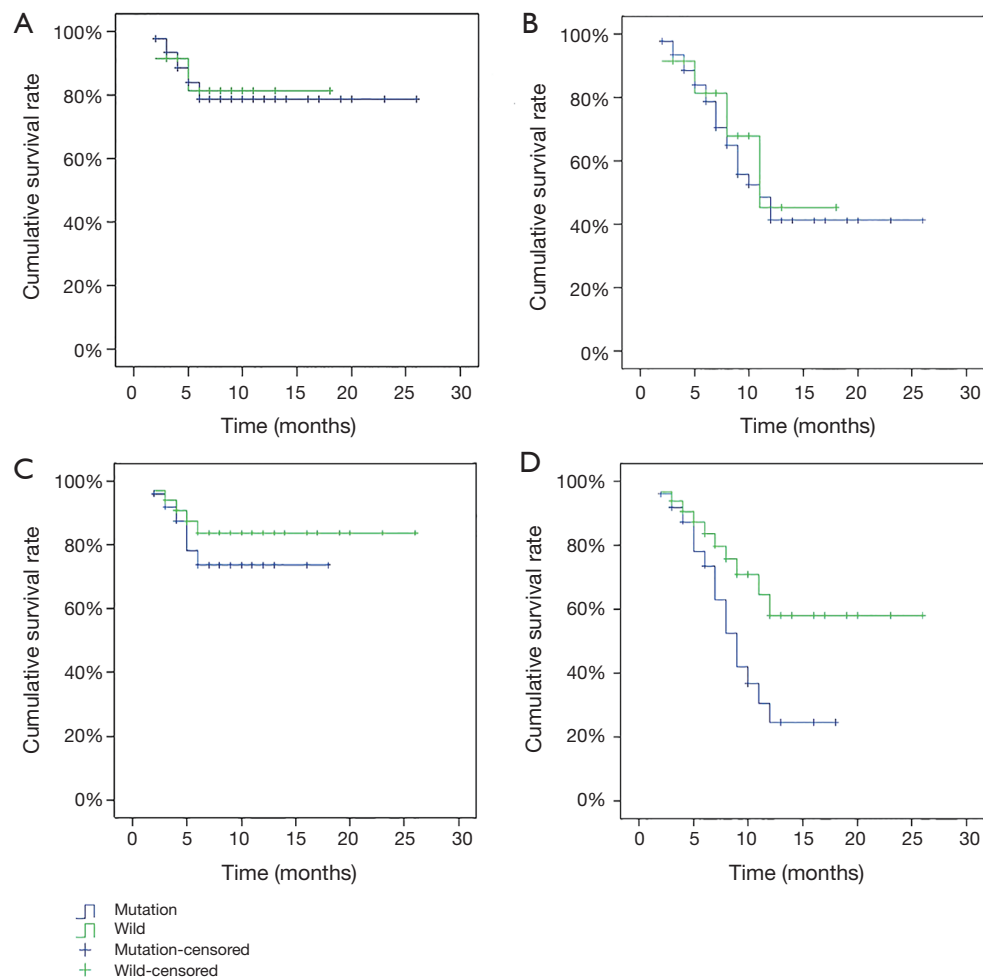


Figure 1 Survival of patients with pancreatic cancer based on plasma and stool K-*ras* mutation status. (A) The 6-month survival rates were compared between patients with and without K-*ras* mutations in stool; (B) the 1-year survival rates were compared between patients with and without K-*ras* mutations in stool; (C) the 6-month survival rates were compared between patients with and without K-*ras* mutations in plasma; (D) the 1-year survival rates were compared between patients with and without K-*ras* mutations in plasma.

mutation status in the prediction of prognosis in patients with pancreatic cancer. However, these results are in conflict with the use of K-*ras* mutation status as a prognostic factor for pancreatic cancer (10). Nonetheless, there appears to be a consensus that K-*ras* mutation status in plasma predicts a worse survival outcome in patients with pancreatic cancer (10,23,24), although survival might not differ by K-*ras* mutations in tissue DNA (25,26). In agreement with these findings, the present study revealed that plasma, not stool, K-*ras* mutation status was significantly associated with the 1-year survival rate of patients with pancreatic cancer (24.5% *vs.* 58.1%). Although the 6-month survival rate did not significantly differ by plasma K-*ras* mutation status, this

exhibited a trend towards this (73.6% *vs.* 83.7%), which may be due to the small sample size of the study. The discrepancy in the prognostic significance of plasma and stool K-*ras* mutation status may be explained by the origin of plasma DNA. It has been hypothesized that plasma DNA might originate from tumor cells detached from metastases (24), and the detection of K-*ras* mutations in plasma may suggest that metastases have occurred. Thus, it is not surprising that plasma K-*ras* mutation status is associated with poorer prognosis.

Since the clinicopathological characteristics of pancreatic cancer often correlate with its prognosis, the relationship of plasma and stool K-*ras* mutation status with the

clinicopathological characteristics of pancreatic cancer was also analyzed. Consistent with a previous finding that K-ras mutation status did not correlate with any of the clinicopathological parameters (24), no significant associations were detected, except that K-ras mutation status in plasma was significantly associated with the TNM stage. Furthermore, a previous study has noted that the presence of K-ras mutations in peripheral blood might reflect different tumor stages (14). This result can also be explained by the hypothesis that the detection of K-ras mutations in plasma suggests the presence of metastases (24), which may upgrade the tumor stage.

The present study has several major limitations. First, as mentioned above, the sample size was small, which may result in some insignificant associations. Second, the retrospective nature of the study may have caused errors and bias inherent to such kind of study. Third, the follow-up period was relatively short. Finally, false positives and false negatives associated with nanoprobe-based assay may lead to the possibility of failing to fully detect the true relationship of K-ras mutation status with the prognosis and clinicopathological characteristics of pancreatic cancer. Future studies are warranted to carefully address these issues.

In conclusion, we have developed a novel fluorescent nanoprobe-based assay for detecting K-ras mutations in plasma and stool samples from patients with pancreatic cancer. This assay exhibits satisfactory sensitivity and specificity when plasma and stool samples are simultaneously detected. The nanoprobe-based detection of plasma K-ras mutation status is associated with survival and tumor stage in patients with pancreatic cancer, and can therefore serve as an auxiliary tool for prognostic evaluation.

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Footnote

Conflicts of Interest: All authors have completed the ICMJE uniform disclosure form (available at <http://dx.doi.org/10.21037/tcr.2018.07.07>). The authors have no conflicts

of interest to declare.

Ethical Statement: The authors are accountable for all aspects of the work in ensuring that questions related to the accuracy or integrity of any part of the work are appropriately investigated and resolved. The study was conducted in accordance with the Declaration of Helsinki (as revised in 2013). This study protocol was approved by the Ethics Committee of the Second Affiliated Hospital of Jiaxing University (No. L2015-28). Informed consent was obtained from all participants.

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