Comparison of tissue and molecular yield between fine-needle biopsy (FNB) and fine-needle aspiration (FNA): a randomized study

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ABSTRACT

Background and study aims Recently, a new Franseen design endoscopic ultrasound-guided fine-needle biopsy (EUS-FNB) needle was developed with the goal of providing more tissue for histology. We compared the tissue adequacy rate and nucleic acid yield of 22G EUS-FNB vs. 22G

endoscopic ultrasound-guided fine-needle aspiration (EUS-FNA), in solid gastrointestinal and extra-intestinal lesions.

Patients and methods We conducted a randomized crossover study and recruited 36 patients. We performed three passes for pancreatic lesions and two passes for other lesions, using each needle. We blinded the pathologist to needle assignment. We assessed the diagnostic tissue adequacy rate and compared the total tissue area, diagnostic tissue area, and desmoplastic stroma (DS) area in cases of carcinoma. We also examined the nucleic acid yield of the two needles in pancreatic lesions.

Results The lesions included 20 pancreatic masses (55%), six gastric subepithelial lesions (17%), five lymph nodes (14%) and five other abdominal masses (14%). Mean $\pm$ SD lesion size was 3.8 ± 2.0 cm. The final diagnosis was malignant in 27 lesions (75%) and benign in nine lesions (25%). We found EUS-FNB procured significantly more median total tissue area (5.2 mm^2 vs. 1.9 mm^2 , $P < 0.001$), diagnostic tissue area (2.2 mm^2 vs. 0.9 mm^2 , $P = 0.029$), and DS area (2 mm^2 vs. 0.1 mm^2 , $P = 0.001$) in lesions diagnosed as carcinoma ($n = 23$), as compared to EUS-FNA. In pancreatic lesions, EUS-FNB obtained significantly more nucleic acid than EUS-FNA (median; $4,085 \text{ ng}$ vs. 2912 ng , $P = 0.02$). There was no difference in the cellblock or rapid on-site cytological evaluation (ROSE) diagnostic yield between the needles.

Conclusion The 22G EUS-FNB provides more histological core tissue and adequate nucleic acid yield compared to 22G EUS-FNA. In this study, the diagnostic performance was similar between the needles

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TRIAL REGISTRATION: Single center, prospective, randomized crossover trial NCT03109639 at clinicaltrials.gov

Introduction

Endoscopic ultrasound-guided fine-needle aspiration (EUS-FNA) is a safe and preferred method for tissue acquisition from solid gastrointestinal and extra-intestinal lesions [1, 2]. However, EUS-FNA has certain drawbacks: The yield is predominantly cellular and rarely provide tissue blocks; need to perform multiple passes; and requirement for rapid onsite cytopathology (ROSE) assessment to improve diagnostic yield [3]. In addition, the cellular yield of EUS-FNA limits architecture assessment, performance of immunohistochemistry and molecular analysis [4, 5]. Such assessments are essential to establish a diagnosis in neoplasms such as lymphoma and gastrointestinal stromal tumors and they are pivotal for clinical trials evaluating molecular markers for personalized oncological treatment. Multiple techniques of EUS-FNA-guided tissue acquisition have been described, but they have failed to consistently show improved diagnostic yield [6–8].

Newer needles with different tip designs and side fenestration (endoscopic ultrasound-guided fine-needle biopsy [EUS-FNB]) have been developed with the goal of obtaining a core sample for histology. Studies comparing the diagnostic efficacy of EUS-FNA and EUS-FNB have yielded conflicting results [9–13]. A meta-analysis comparing one type of EUS-FNB (ProCore, Cook Endoscopy) to standard EUS-FNA showed no difference in sample adequacy, diagnostic accuracy, or acquisition of core specimen between the two needles [14].

Recently, an EUS-FNB needle with Franseen geometry (Acquire, Boston Scientific, United States) has been developed to procure tissue specimen for histology [15–18]. Early evidence assessing diagnostic yield of the Franseen needle in solid pancreatic lesions is promising [19]. However, more evidence of its diagnostic utility in pancreatic and non-pancreatic solid lesions is needed. We performed a randomized trial comparing the outcome of 22G Franseen EUS-FNB (Acquire, Boston Scientific, United States) and 22G standard EUS-FNA (Expect, Boston Scientific, United States) in solid gastrointestinal and extra-intestinal lesions.

Patients and methods

Trial design

We conducted a prospective, randomized, single-blinded crossover trial at Singapore General hospital between April 2017 and November 2017. The institutional review board approved the study. All authors had access to the study data and reviewed and approved the final manuscript (Clinicaltrials.gov, NCT03109639). The study was conducted in accordance with the ethical principles detailed in the Declaration of Helsinki and was consistent with Good Clinical Practices recommendation. We reported our outcome according to the CONSORT recommendation of reporting a randomized trial.

Participants

We enrolled 40 patients who were referred for EUS-guided tissue acquisition. We obtained informed consent from all patients. Inclusions criteria were age older than 18 years, pres-

ence of only solid lesions confirmed by endoscopy or radiology, and ability to comply with the procedure and provide informed consent. Patients were excluded if they had active bleeding, coagulopathy (INR > 1.5, platelet count < 50,000), were concurrently taking anticoagulants (e.g., warfarin) and a thienopyridine (e.g., clopidogrel), had an intervening large blood vessel, or had difficulty tolerating the procedure.

Intervention

Procedure

Four experienced endoscopists trained in EUS and EUS-guided tissue acquisition techniques performed the procedures under moderate sedation using midazolam and fentanyl. The endoscopists used the Franseen EUS-FNB in clinical practice and were familiar with the device before participating in the study. They were aware of the type of needle used. We used the curved linear array echoendoscope for the study (GF-UC140P, Olympus, USA; EG-3870UTK, Pentax, Japan; or EG-580UT, Fujifilm, Japan). We identified the lesions using EUS, confirmed the absence of cystic components and then randomized them to the study needles for tissue acquisition.

We advanced the assigned needle into the lesion under EUS guidance, removed the stylet and applied a negative pressure using a 10-cc suction syringe. We accessed the lesions through the same route, whenever possible, using both the needles. We practiced the fanning method and performed three passes for pancreatic lesions and two passes for other lesions using each needle type. All the assigned passes were completed with the initial device before crossing over to the alternate needle (► Fig. 1). If there was a technical failure (defined as needle malfunction before we reached a diagnosis), the patient crossed over to the alternative needle.

Tissue preparation for pathology

We expressed a portion of the specimen onto a slide using the stylet and prepared air-dried and alcohol-dried smears on-site. We flushed the residual material with normal saline and placed it in formalin solution for cell-block analysis. We collected both the smears and cell-block samples after each needle passes.

The cytotechnician and the pathologist reviewing the smears and slides were blinded to the type of needle used. The slides were reviewed on-site by the cytotechnician after each pass to assess for specimen adequacy. The endoscopist was informed of the outcome of on-site examination only after completion of the assigned number of needle passes to minimize operator bias. We centrifuged and concentrated the cell-block sample and created a tissue clot, fixed the tissue clot in formalin and embedded it with paraffin. We then sectioned and stained it with eosin and hematoxylin for histological assessment. We graded the sample as optimal or suboptimal based on presence of core tissue that enabled adequate evaluation of histological architecture.

We calculated the area of total tissue, diagnostic tissue, and desmoplastic stroma (in carcinoma) using the Philips IntelliSite Pathology solutions image management system. We measured the greatest linear dimension of each tissue core fragments and

summed it to estimate the total tissue length. We then identified a representative tissue core fragment and measured the largest diameter. Using these numerical values, we calculated the area of total tissue (► **Fig. 2**). In a similar manner, we estimated the area of diagnostic tissue and desmoplastic stroma. When needed, we performed immunohistochemical or special staining to characterize the lesion better.

Quantification and quality assessment of nucleic acid

We extracted the genomic DNA and RNA from the formalin-fixed paraffin embedded (FFPE) cell blocks from each needle. We estimated the amount of nucleic acid by NanoDrop spectrophotometry (Thermo Scientific, Wilmington, Massachusetts, United States) and further quantified using the Qubit fluorimetry (Life Technologies, Grand Island, New York, United States). We assessed the quality of the DNA and RNA by reading the ratio of absorbance at 260/280 and 260/230 nm. We calibrated the instruments and performed the procedure as per the manufacturer's instructions.

We also explored the possibility of performing molecular studies from EUS-derived tissue samples using Oncomine Comprehensive panel V3 (Life Technologies, Pleasanton, California, United States), an assay that contains 4648 primer pairs designed for hotspots, targeted regions, and gene fusions of 161 known genes relevant to solid tumors. It provides the reagents for library construction and four pools of multiplex polymerase chain reaction primers for preparation of amplicon libraries from formalin-fixed paraffin embedded tumor samples. We randomly selected two solid pancreatic lesions for this analysis and followed optimized protocols.

► **Fig. 2** Scanning power photomicrograph of a cell-block specimen. The greatest linear diameter of the tissue core fragments was measured (green line) and summed to obtain the total histological core tissue length. A representative core fragment was identified, and the diameter of it was measured (red line). The area of the total histological core was estimated from these measurements.

Outcomes

The primary outcome was to evaluate the diagnostic tissue adequacy rate and compare the area of total tissue, diagnostic tissue, and desmoplastic stroma (in cases with a diagnosis of carcinoma) from the histological core procured using both the needles. Our secondary outcome was to compare the quantity and quality of the DNA and RNA in the samples obtained using the two needles. We defined diagnostic tissue adequacy as presence of histological tissue representative of the sampled lesion and diagnostic yield as the percentage in which a definitive diagnosis could be established from the sampled lesion [20].

We established the final diagnosis based on: pathology from the surgical specimen; radiological imaging including computed tomography, magnetic resonance imaging, and positron emission tomography scan; or clinical progress. We considered a lesion to be benign if there was spontaneous resolution or interval stability in radiological imaging.

Sample Size

In a preliminary study, the EUS-FNB needle provided diagnostic material for histology in >95% of patients [15]. Standard EUS-FNA has a diagnostic histology yield of only 40% [21,22]. We estimated that a total of 36 patients would be needed to detect a 30% difference in histological yield between the two needles, with an α of 0.05 and a power of 90%. We factored in a 10% dropout rate and enrolled a total of 40 patients.

► **Table 1** Comparison of outcomes between EUS-FNB and EUS-FNA

Patients (n = 36)			
	FNB	FNA	P value
Technical success (%)	100%	100%	
Presence of histology core, n (%)	35, (97%)	28, (77%)	0.03
Median total tissue area, mm ² (IQR)	5.2 (2.1 – 14.1)	1.9 (0.4 – 6.6)	<0.001
Median diagnostic tissue area, mm ² (IQR)	2.2 (0.5 – 6.1)	0.9 (0.2 – 3.4)	0.029
Desmoplastic Fibrosis Assessment (n = 23)	20 (87%)	17 (74%)	0.45
Median desmoplastic area, mm ² (IQR)	2 (0.4 – 6.9)	0.1 (0 – 0.5)	0.001
Conducive for IHC staining, (n = 34)	33 (97%)	30 (88%)	0.35
Adverse events, n (%)	0	0	

FNB, fine-needle biopsy; FNA, fine-needle aspiration; IQR, interquartile range; IHC, immunohistochemistry

Randomization and blinding

We randomized the patients to one of two groups (A and B) and recorded the results (► **Fig. 1**). In group A, patients received standard EUS-FNA followed by EUS-FNB. In group B, the assigned passes were performed first using EUS-FNB and then crossed over to EUS-FNA. Randomization was performed in blocks to ensure that the groups were balanced periodically. We used web-based randomization (Sealed Envelope Ltd 2016; <https://www.sealedenvelope.com/simple-randomiser/v1/>) for this purpose. We blinded the cytotechnician and the pathologist to the needle used for specimen collection.

Statistical methods

We expressed continuous variables as the mean ± standard deviation (SD) or median (interquartile range [IQR]). We compared the means using a paired *t*-test and medians using Wilcoxon ranked sum test. We presented categorical variables in percentage, and the correlation with the different technique was studied using McNemar's test. Prior to these tests, we performed a linear mixed model with a fixed period and technique effects, and random patient-level intercepts to investigate if there was a significant period effect. We performed statistical analysis using R 3.4.2 software (R Core Team (2017). R: A language and environment for statistical computing. R foundation for statistical computing, Vienna, Austria). A *P* value < 0.05 was considered significant.

Results

We enrolled 40 patients and excluded four from the study; three had cystic lesions, and one withdrew from the study. The remaining 36 patients were randomized to one of the two study groups. Mean age was 63.5 ± 11.4 years, and the majority were men (56%, n = 20). The lesions sampled included 20 pancreatic masses (55%), six gastric sub-epithelial (SEL) lesions (17%), five lymph nodes (14%) and five other abdominal masses (14%) (2 peri-gastric mass, two retroperitoneal mass, 1 liver metastasis). Mean ± SD size of the lesion was 3.8 ± 2.0 cm. We

accessed the lesion through the trans-gastric route in 22 patients (61%), trans-duodenally in 14 patients (39%) and trans-esophageally in two patients (5%). The final diagnosis was malignancy (22 adenocarcinomas, one liposarcoma, one squamous cell carcinoma, one neuroendocrine tumor, two lymphomas) in 27 patients (75%) and benign lesion (6 spindle cell tumor, 1 splenunculus, 2 inflammatory mass) in 9 (25%). We did not have any technical difficulty, and the procedure was successful in all patients.

Diagnostic tissue adequacy

We found EUS-FNB obtained histological core tissue more frequently than EUS-FNA (97% vs. 77%, *P* = 0.03). Diagnostic adequacy and yield of the histological tissue was similar between EUS-FNB and EUS-FNA (81% vs. 64%, *P* = 0.19). We did not observe any difference in the cell-block diagnostic adequacy and yield between the two needles for pancreatic and non-pancreatic lesions.

When assessing the histology, we found that EUS-FNB provided significantly more median total tissue area than EUS-FNA (5.2 mm² vs. 1.9 mm², *P* < 0.001) in both pancreatic and non-pancreatic lesions (► **Table 1**). Median area of the diagnostic tissue within the histological sample was significantly more with EUS-FNB than with EUS-FNA (2.2 mm² vs. 0.9 mm², *P* = 0.01) in all lesions (► **Fig. 3**). We assessed presence of desmoplastic stroma (DS) in patients with a final diagnosis of carcinoma (64%, n = 23). We found that EUS-FNB provides significantly more DS tissue area than EUS-FNA (2 mm² vs. 0.1 mm², *P* < 0.001) (► **Fig. 4**). When subcategorized, we found EUS-FNB provided significantly more total tissue, diagnostic tissue and desmoplastic stroma in solid pancreatic lesions compared to EUS-FNA. However, in non-pancreatic lesions, EUS-FNB yielded a similar amount of total tissue and diagnostic tissue as EUS-FNA.

We did not observe any period effect in this crossover trial design (► **Table 2**), and EUS-FNB consistently obtained more histological tissue than EUS-FNA. We evaluated suitability for performing immunohistochemistry (IHC) analysis in 34 lesions

► **Fig. 3** ROSE and cell-block assessment of samples obtained from gastric GIST. **a** Diff-Quick staining of cytology specimen obtained using Franseen EUS-FNB. **b** Diff-Quick staining of similar cytology specimen obtained using EUS FNA. **c** H&E staining of histology obtained using Franseen EUS-FNB. **d** H&E staining of similar histology obtained using EUS-FNA.

► **Fig. 4** ROSE and cell-block assessment of samples obtained from a lymph node in metastatic squamous cell carcinoma. **a** Diff-Quick staining of cytology specimen obtained using Franseen EUS-FNB. **b** Diff-Quick staining of similar cellular yield obtained using EUS FNA. **c** H&E staining of histology obtained using Franseen EUS-FNB shows malignant cells surrounded by dense desmoplastic stroma. **d** H&E staining of histology obtained using EUS-FNA shows malignant cells with scanty desmoplastic stroma.

► **Table 2** Linear mixed model with a fixed period (Group) and needle (EUS-FNB) effects and random patient-level intercepts for different measurements.

	Estimation	P value
Total histological core area		
Group B	2.9 (-0.37, 6.26)	0.08
EUS-FNB	3.9 (1.94, 5.88)	<0.001
Total diagnostic tissue area		
Group B	2.4 (-0.82, 5.53)	0.141
EUS-FNB	1.6 (0.17, 3.01)	0.029
Total desmoplastic stroma area		
Group B	0.7 (-1.6, 2.92)	0.549
EUS-FNB	3.5 (1.6, 5.38)	0.001
EUS-FNB, endoscopic ultrasound-guided fine-needle biopsy		

where the cell-block had diagnostic tissue and found no significant difference between EUS-FNB and FNA (97 % vs. 88 %).

On-site (ROSE) diagnostic yield

We collected smears after each needle pass and assessed for on-site (ROSE) diagnostic yield. We found the overall ROSE diagnostic yield was similar between EUS-FNB (81 %) and EUS-FNA (81 %). In pancreatic lesions, we found the first pass diagnostic yield, as determined by ROSE, was similar between EUS-FNB (80 %) and EUS-FNA (70 %). In non-pancreatic lesions, both EUS-FNB (69 %) and EUS-FNA (69 %) performed suboptimally. We found that additional needle passes using FNB and FNA did not significantly increase the ROSE diagnostic yield in any lesions (► **Table 3**).

When combined with cell-block, the overall diagnostic yield improved to 92 % in EUS-FNB group and 86 % in EUS-FNA group.

Quantification and qualification of nucleic acid

We estimated the volume of DNA and RNA obtained from the pancreatic lesions (n = 20) by the two needles. The lesions included 18 pancreatic adenocarcinomas, one neuroendocrine tumor, and one metastatic renal cell carcinoma. We found that EUS-FNB procured significantly more nucleic acid than EUS-FNA [median; 4,085 (2,804–6711) ng vs. 2912 (2,287–4,658) ng, *P* = 0.02). The increased nucleic acid yield corresponds to the increased total tissue yield with EUS-FNB. We noticed the EUS-FNB yielded significantly more RNA [median; 1,634 (1089–

► **Table 3** On-site and cell-block diagnostic rate.

	FNB	FNA	P value
ROSE diagnostic accuracy, n (%)	29 (81 %)	29 (81 %)	1
Pancreatic lesions (n = 20)			
▪ First pass	16 (80 %)	14 (70 %)	
▪ Second pass	2 (50 %)	4 (67 %)	
▪ Third pass	0 (0 %)	0 (0 %)	
Non-pancreatic lesions (n = 16)			
▪ First pass	11 (69 %)	11 (69 %)	
▪ Second pass	0 (0 %)	0 (0 %)	
Cell block diagnostic accuracy, n (%)	29 (81 %)	23 (64 %)	0.19
Pancreatic lesions (n = 20)			
▪ First pass	12 (60 %)	9 (45 %)	
▪ Second pass	1 (12 %)	1 (9 %)	
▪ Third pass	1 (14 %)	2 (20 %)	
Non-pancreatic lesions (n = 16)			
▪ First pass	13 (81 %)	10 (63 %)	
▪ Second pass	2 (40 %)	1 (14 %)	
Combined ROSE and cell block diagnosis, n (%)	33 (92 %)	31 (86 %)	0.71
Pancreatic lesions (n = 20)			
	18 (90 %)	18 (90 %)	
Non-pancreatic lesions (n = 16)			
	15 (94 %)	13 (81 %)	
ROSE, rapid on-site cytological evaluation

► **Fig. 5** Quantification of nucleic acid obtained by the two needles. EUS-FNB provided significantly more nucleic acid than EUS-FNA. EUS-FNB yielded significantly more RNA and a trend towards an increased amount of DNA.

3,939) ng vs. 1295 (986 – 1,782) ng, $P=0.02$] and increased the amount of DNA [median; 2,185 (1,478 – 3,066) ng vs. 1477 (1,151 – 2,522), $P=0.08$] when compared to EUS-FNA (► **Fig. 5**). We assessed the first-pass nucleic acid yield and found a trend towards increased yield with EUS-FNB compared to EUS-FNA [median; 1142 (618 – 2435) vs. 1090 (750 – 1341) ng, $P=0.07$].

We analyzed the purity of the DNA and RNA extracted by measuring the ratio of absorbance at 260/280 nm (optimal threshold range, 1.6 – 2.2) and 260/230 nm (optimal threshold > 1) using Nanodrop spectrophotometer. We found that both EUS-FNB (95%) and FNA (90%) needles procured high-quality DNA in pancreatic lesions. There was a trend towards increased higher-quality RNA yield with EUS-FNB (30%) than with EUS-FNA (10%). We also evaluated the ability to perform sequencing analysis from the nucleic acid obtained using both needles ($n=2$). We found that both needles exceeded the manufacturer recommendation of minimum DNA or RNA concentration (10 ng/reaction) needed for tumor profiling using the next-generation sequencing assay (NGS, OncoPrint™ V3). The NGS results were identical between the two needles.

Complications

There were no complications after EUS-FNB and EUS-FNA in either group.

Discussion

We report the outcome of an RCT comparing Franseen design EUS-FNB versus standard EUS-FNA in solid pancreatic and non-pancreatic lesions. EUS-FNB provided significantly more total histological tissue, diagnostic tissue, and desmoplastic stroma compared to EUS-FNA. However, the diagnostic yield as measured using ROSE and cell block was similar.

The Franseen EUS-FNB needle design has three symmetrical cutting edges. It was postulated that these edges provide stability at puncture, penetrate quickly, and capture maximum tissue with minimal fragmentation. Our study provided some support for this hypothesis and is in agreement with a recent randomized study showing superior histological core and tumor tissue procurements with the 22G Franseen EUS-FNB compared to a standard 22G EUS-FNA needle, in solid pancreatic lesions [19]. It also appears that Franseen EUS-FNB, with its coring ability and tendency to provide a large volume of tissue for cell-block analysis, may overcome the need for ROSE [23, 24].

In a large retrospective study assessing the utility of Franseen EUS-FNB in solid pancreatic and non-pancreatic lesions, the overall cell-block diagnostic yield (62%) was found to be significantly lower [25]. This observation was in sharp contrast to the superior cell-block yield (>90%) reported in the randomized studies on Franseen EUS-FNB in pancreatic lesions [19, 23]. Our results did not support the results of the randomized study. In fact, we found the cell-block diagnostic yield to be lower (81%), and the overall yield improved only with presence of ROSE. One possible cause could be the proportionately larger yield of desmoplastic stroma (10 times higher) than diagnostic tissue (1.5 times higher) compared with EUS-FNA. Other probable contributing factors may include the heterogeneous study cohort, technical difference between the operators, experience of the on-site cytotechnician, cytopathologist expertise and use of specialized imaging software to assess the histology sample.

There are potential issues with obtaining desmoplastic tissue. In pancreatic cancer (PC), a dense, extensive desmoplastic reaction is a typical finding and concern has been raised about its role in drug resistance [26]. Procuring histological samples abundant in desmoplastic stroma may be useful for clinical trials evaluating therapy that targets desmoplastic stroma [27, 28]. Similarly, molecular profiling of PC and application of next-generation sequencing may provide an opportunity to advance development of targeted therapies and improve PC treatment outcomes [20, 29, 30]. Studies evaluating personalized medicine in PC have relied mainly on surgically acquired tissue for performing genetic analysis. Unfortunately, most PC present at an inoperable stage and real-time genetic analysis of advanced cancers is limited. EUS, a safe and minimally invasive technique, is widely used to acquire tissue and establish a diagnosis in such situations. Until now, there was varying evidence on the suitability of EUS-FNA- and EUS-FNB-acquired tissue samples for genomic analysis [31 – 37]. Some studies suggest EUS-FNA-acquired cytology samples and liquid cytology specimens (FNA rinse material) have a higher concentration of intact and pure tumor cells and are superior for NGS. Others consider that the cellular material obtained using EUS-FNA is often graded to be insufficient, contaminated, composed of poor-quality DNA [29, 38, 39] and suboptimal for genetic analysis.

Gleeson et al showed that a cytology slide >5000 cells, tissue volume >1.2 cm² in FFPE sample and at least 5 ng/μL of recoverable DNA has a higher success rate (>90%) at NGS [31]. Our study showed that the Franseen EUS-FNB needle can yield a large amount of tissue and provide an adequate amount of nu-

cleic acids with relatively preserved high-quality DNA and RNA. In the limited samples ($n=2$) with NGS results, the quality metrics were all met with high scores, thus providing evidence that FNB-acquired specimens can serve as a tool for genomic studies and a surrogate to surgical specimens, especially in patients not amenable to surgical intervention [40]. Rodriguez et al showed that reliable RNA sequencing (Illumina, Inc, United States) can be performed if the EUS-TA can acquire 100 ng of total RNA [41]. Our study demonstrated that the increased high-quality RNA yield with EUS-FNB may make performance of RNA sequencing feasible. It would provide a platform for measuring gene expression and capturing novel diagnostic and prognostic information about pancreatic cancer.

Our study has certain limitations. It was not possible to blind the endoscopists to the study needles used. We believe by applying a crossover trial design, may have eliminated the operator-related bias. Second, our study is possibly underpowered as the diagnostic histology yield with EUS-FNA (64%) was higher than the reference rate (40%) used during power calculation. A type II error cannot be excluded. However, post-hoc analysis demonstrated that the study achieved 90% power with α of 0.05, when estimated to detect a 1 mm² difference in the median total tissue area between the two needles [19]. Third, mean size (3.8 ± 2.0 cm) of the lesions included in the study was larger, which may have contributed to the similar diagnostic efficacy for EUS-FNA and EUS-FNB. Performance of EUS-FNB and EUS-FNA in smaller (<2 cm) lesions needs to be assessed. Fourth, our secondary outcome was to perform a DNA and RNA quantification in all the lesions that we sampled. Funding limitations prevented us from pursuing more wide-scale NGS. Fifth, we estimated DNA and RNA from FFPE specimens. Formalin fixation and sectioning of the cell block may have degraded and fragmented DNA and RNA, resulting in lower yields. Performing a dedicated additional pass and using RNA later (Ambion, Austin, Texas, United States) or Cytolyt (Hologic Co, Marlborough, Massachusetts, United States) preservative may have increased the RNA and DNA yield further [42, 43]. Lastly, we did not use any automated or artificial intelligence software for histological assessment and quantification. Nonetheless, such digital software is not widely implemented, and manual histological assessment is more widespread and applicable at this time point.

Conclusion

In conclusion, our study shows that the Franseen EUS-FNB and EUS-FNA device are equally effective in establishing a diagnosis in all solid gastrointestinal and extra-intestinal lesions. We have demonstrated that the Franseen EUS-FNB device can obtain more histological tissue and has better nucleic acid yield with quality and quantity sufficient for downstream genomics applications like NGS. This is a significant step in EUS-FNB becoming a convenient and safe method for obtaining tumor material for precision genomics.

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Competing interests

None

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