

STATISTICAL ANALYSIS PLAN

STUDY TITLE: A Randomized, Open-Label, Phase 1/2 Trial of Gemcitabine plus Nab-paclitaxel with or without FG-3019 as Neoadjuvant Chemotherapy in Locally Advanced, Unresectable Pancreatic Cancer

PROTOCOL NUMBER: FGCL-3019-069

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STATISTICAL ANALYSIS PLAN SIGNATURE PAGE**Approvals**

I have reviewed and accepted the information in this document to be a true and accurate representation of the Statistical Analysis Plan, Version 1.0 for Study FGCL-3019-069.

Initiator:**Signature:****Date:** 3/21/2018**Reviewed by****Signature:****Date:** 3/22/2018**Signature:****Date:** 3/21/2018**Signature:****Date:** 3/21/2018**Signature:****Date:** 3/22/2018**Signature:****Date:** 3/22/2018**Signature Significance**

The following significance is lent to the signatures on the Approvals page of this document.

Signatory	Significance
Initiator	By signing, the author is attesting that the content of the document is complete and accurate.

Reviewer	By signing, the reviewer is attesting that the document's approach and contents are compliant with the study protocol, all appropriate, regulatory requirements, and other significant guidelines. This individual(s) has reviewed the document for accuracy and completeness.
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REVISION HISTORY

Version	Date	Description

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ABBREVIATIONS

AE	Adverse Event
AJCC	American Joint Committee on Cancer
BMI	Body Mass Index
BSA	Body Surface Area
BP	Blood Pressure
CA	Carbohydrate Antigen
CBC	Complete Blood Count
CI	Confidence Interval
CMH	Cochran-Mantel-Haenszel
CR	Complete Response
CRF	Case Report Form
CS	Clinically Significant
CTCAE	Common Terminology Criteria for Adverse Events
CTGF	Connective Tissue Growth Factor
DBP	Diastolic Blood Pressure
EBL	Estimated Blood Loss
ECG	Electrocardiogram
ECOG	Eastern Cooperative Oncology Group
eDISH	evaluation of Drug Induce Severe Hepatotoxicity
EFS	Event Free Survival
EOS	End of Study
EOT	End of Treatment
EUS	Endoscopic Ultrasound
FDG-PET	[18F]-fluorodeoxyglucose-positron emission tomography
HAHA	Human Anti-FG-3019 Antibody
HR	Heart Ratio
ICF	Informed Consent Form
ITT	Intent-to-Treat
IV	Intravenous
LLN	Lower Limit of Normal for a laboratory parameter
MedDRA	Medical Dictionary for Regulatory Activities

NCCN	National Comprehensive Cancer Network
NCI	National Cancer Institute
NCS	Not Clinically Significant
ORR	Objective Response Rate
OS	Overall Survival
PD	Progressive Disease
PET	Positron Emission Tomography
PFS	Progression-Free Survival
PK	Pharmacokinetic(s)
PR	Partial Response
PT	Preferred term
RECIST	Response Evaluation Criteria in Solid Tumors
RMST	Restricted mean survival time
SAE	Serious adverse event
SAP	Statistical Analysis Plan
SBP	Systolic Blood Pressure
SD	Stable Disease, Standard deviation
SE	Standard Error
SMQ	Standardized MedDRA queries
SOC	System Organ Class
SSI	Surgical Site Infection
SUVmax	Maximum standardized uptake value
TEAE	Treatment-emergent Adverse Event
TESAE	treatment-emergent serious adverse event
TNM	tumor/node/metastasis
TTP	Time to Progression
ULN	Upper Limit of Normal for a laboratory parameter
WHO-DD	World Health Organization Drug Dictionary

1 INTRODUCTION

This Statistical Analysis Plan (SAP) documents planned analyses for Study FGCL-3019-069 Amendment 6 (September 29, 2016): A Randomized, Open-Label, Phase 1/2 Trial of Gemcitabine plus Nab-paclitaxel with or without FG-3019 as Neoadjuvant Chemotherapy in Locally Advanced, Unresectable Pancreatic Cancer.

This SAP includes statistical analysis methods, statistical models, definitions, and data handling rules. It supersedes the statistical section in the protocol in case of difference.

2 STUDY OBJECTIVES

The objectives of this trial are:

- To investigate the safety, tolerability, and efficacy of FG-3019 administered with gemcitabine and nab-paclitaxel in the treatment of locally advanced, unresectable pancreatic cancer.
- To evaluate the pharmacokinetics of FG-3019 (concentration maximum [C_{\max}] and concentration minimum [C_{\min}]) and correlation with other study endpoints in this treatment setting.

The exploratory objectives of this trial are:

- To assess the effects of neoadjuvant treatments on a panel of tumor and plasma biomarkers.
- To evaluate the effects of neoadjuvant treatment on gene and protein expression profiling using freshly obtained tumor tissues or blood samples.

3 STUDY DESIGN

3.1 Overview

This is a Phase 1/2, randomized, open-label trial to evaluate safety, tolerability and efficacy of gemcitabine plus nab-paclitaxel with FG-3019 (Arm A) and gemcitabine plus nab-paclitaxel (Arm B) in subjects with locally advanced, unresectable pancreatic cancer. Up to 42 evaluable subjects will be included in this trial. Each treatment cycle is 28 days long and subjects may receive up to six cycles of treatment. Subjects will be evaluated for their surgical eligibility at end of treatment. Subjects will be followed for at least 28 weeks post End of Treatment for disease progression and survival.

Blood samples will be collected periodically for the assessment of PK and pharmacodynamics (PD). Study centers participating in this study have the option of participating in the collection of tumor core biopsies that will be used in an exploratory portion of this study evaluating tumor biomarkers including protein expression. At centers not participating, tumor samples obtained from resected specimens may be used in biomarker studies.

This trial has four periods:

- Screening and Disease Staging
- Treatment
- Surgery or Short-Term Follow Up
- Long-Term Follow Up

A schematic overview of the study, including study period timeframes, is provided in Figure 1 of the protocol.

3.2 Study Population

Patients with locally advanced unresectable pancreatic cancer.

3.3 Sample Size Determination

A total sample size of 42 evaluable subjects, 28 in Arm A and 14 in Arm B, is considered appropriate, based on clinical judgment, for initial exploration of safety, tolerability, and treatment effect.

3.4 Randomization and Treatment Assignment

Prior to Amendment 3, eligible subjects were randomized 1:1 to:

- **Arm A:** gemcitabine, nab-paclitaxel, and FG-3019
- or
- **Arm B:** gemcitabine and nab-paclitaxel

In Protocol Amendment 4, the randomization schedule was changed to 3:1 in order to achieve an overall 2:1 randomization (approximately 28 in Arm A and 14 in Arm B at study end).

3.5 Dosing Schedule

Each treatment cycle is 28 days long and subjects may receive up to six cycles of treatment.

- **Arm A treatment consists of:**
 - Gemcitabine, 1000 mg/m² by IV infusion on Days 1, 8 and 15 of each 28-day treatment cycle.
 - Nab-paclitaxel, 125 mg/m² by IV infusion on Days 1, 8 and 15 of each 28-day treatment cycle.
 - FG-3019, 35 mg/kg by IV infusion on Days 1 and 15 of each 28-day treatment cycle. An additional dose will be given on Day 8 of the first cycle.
- **Arm B treatment consists of:**
 - Gemcitabine, 1000 mg/m² by IV infusion on Days 1, 8 and 15 of each 28-day treatment cycle.
 - Nab-paclitaxel, 125 mg/m² by IV infusion on Days 1, 8 and 15 of each 28-day treatment cycle

3.6 Study Assessments

The schedule of assessment is provided in the Appendix 1 of the protocol.

Subject's baseline characteristics, which include demographics, cancer TNM status and staging, medical history, smoking history, prior therapies for cancer including surgery, radiation, and systemic treatment, are collected at Screening.

Eligibility for surgical exploration is evaluated at EOT. For subjects who undergo surgery, surgical outcomes are collected. Post-surgical evaluation is performed 30 days after discharge.

CT scan and RECIST assessment are performed at Screening (within 2 weeks of Cycle 1 Day 1) and approximately every 8 weeks thereafter (prior to Cycle 3, 5 and at EOT) until disease progression or discontinuation of study treatment. Unscheduled scans may be performed at the discretion of the investigators.

PET scan is performed at Screening and at EOT.

CA 19.9 is assessed by local lab at each site at Screening, Day 1 of each cycle (Day 1, Weeks 5, 9, 13, 17, 21), and at EOT.

ECOG performance status is assessed at Screening, Day 1 of Cycle 1, 3, 5, and at EOT. Karnofsky score is additionally evaluated at EOT for the purpose of determining surgical eligibility.

Long-term follow-up for progression and survival is conducted approximately every month for the first 28 weeks following EOT and quarterly thereafter.

Adverse events, concomitant medications, procedures and non-drug therapies are collected from signing ICF through 28 days after the last dose.

Safety lab tests are evaluated at the local lab of each study site at every clinic visit.

Vital signs are measured at every clinic visit.

Physical examination is performed at every clinic visit.

ECG is performed at Screening and at EOT.

Plasma samples for evaluation of FG-3019 concentration level are collected from subjects in Arm A pre- and post-dose on Days 1, 8, 15 of Cycle 1, pre-dose on Day 15 of Cycle 2, pre-and post-dose on Day 1 of Cycles 3 and 6, pre-dose on Day 1 of Cycle 5 and at EOT, and 28 days after last dose.

CTGF and HAHA samples are collected on Day 1 prior to first dose and 28 days after last dose.

Plasma samples for protein expression profile and whole blood sample for PAX gene are collected on Day 1 of Cycles 1, 3, 5 prior to study drug dosing and at EOT.

Up to 4 core biopsies are collected while subjects are undergoing endoscopic ultrasound EUS procedure at Screening. Post-treatment tumor tissues are collected during surgery or during post-treatment EUS. Subjects who are not eligible for surgical exploration are to undergo EUS guided core tumor biopsies to obtain post-treatment tissue samples.

4 STUDY ENDPOINTS AND DEFINITIONS

4.1 Study Endpoints

4.1.1 Safety Endpoints

- Treatment-emergent adverse events (TEAEs), serious treatment-emergent adverse events (TESAEs), death, clinical laboratory tests, and discontinuation of treatment for treatment-related TEAEs.
- Surgical safety with respect to complication rates post resection.

4.1.2 Efficacy Endpoints

- The proportion of subjects who become eligible for surgery.
- The proportion of subjects in whom R0 resection is achieved.
- The proportion of subjects in R0 or R1 resection is achieved.
- Tumor response rates measured by:
 - Complete response (CR) or partial response (PR) per Response Evaluation Criteria in Solid Tumors (RECIST 1.1),
 - At least 50% reduction from baseline in serum CA19-9, or
 - At least 30% reduction from baseline in SUV[max] assessed by FDG-PET
- Median overall survival and 1-year survival rate.
- Median progression free survival and 1-year progression free rate.

4.1.3 Exploratory Outcome Measures

The exploratory outcome measures are:

- FG-3019 plasma concentrations, including trough level (C_{\min}) and maximum concentration (C_{\max}).
- Tissue expression of biomarkers including CTGF, alpha-smooth muscle actin (alpha-SMA), Ki67, Cleaved Caspase-3, and SPARC assessed by immunohistochemistry or ribonucleic acid (RNA).
- Gene and protein expression profiling using freshly obtained tumor tissues, plasma samples or whole blood sample.
- Event free survival (EFS)

4.2 Detail Definitions of the Study Endpoints

4.2.1 Eligibility for Surgical Exploration

Subjects are considered eligible for surgical exploration if one or more of the following criteria are met and have no contraindications:

- Reduction in CA19-9 level by more than 50% at EOT when compared to baseline.
- FDG-PET SUV_{max} decrease by $\geq 30\%$ from baseline.

- Radiological tumor response (PR or CR) per RECIST 1.1.
- Meet the definition of resectable or borderline resectable per NCCN Guidelines

Contraindications include

- Development of distant metastases confirmed by CT scan
- Local progression confirmed by CT scan or progression precluding vascular reconstruction
- Local complication preventing surgery (e.g., PV/SVT thrombosis, pancreatitis)
- Performance status decline to Karnofsky score $\leq 50\%$ or absolute contraindication to surgery
- Other contraindications as specified

4.2.2 Resection Outcome

Resection outcome (R0 or R1) is determined by pathological examination of the surgical specimen after resection and is recorded on Surgical Outcome CRF.

4.2.3 Overall Survival, Time to Progression, Progression-free Survival, and Event-free Survival

Overall survival (OS) is defined as the time from randomization until death from any cause. Date of death is recorded on the Death CRF. For subjects who are alive at data cut or at study closure, the date of last known alive recorded on the Long-Term Follow-up CRF or the date of last clinic visit is defined as the censoring date for OS.

Time to progression (TTP) is defined as the time from randomization until objective tumor progression. Date of disease progression is defined as the date of radiological progression per RECIST 1.1 criteria recorded the Disease Progression CRF. For subjects who have no documented record of radiologic progression at data cut or at study closure, the latest date of the following events is defined as the censoring date for TTP: last post-baseline CT, last post-baseline PET scan, the last record of 'Not Progressed' on the Long-Term Follow-up CRF, and last dose.

Progression free survival (PFS) is defined as the time from randomization until objective tumor progression or death. Specifically, PFS is equal to TTP if the progression event is observed, or equal to OS if the progression event is censored.

Event-free survival (EFS) is defined as the time from randomization to death, or objective tumor progression, or failure to achieve an R0 or R1 resection, or failure to meet surgical eligibility criteria, whichever occurs first.

Event	EFS	Censor or Event
Disease progressed or death before surgical evaluation	EFS =PFS	Event

Discontinued early and had no follow up progression record	EFS = latest date of (last post-baseline CT, last post-baseline PET scan, last dose) – Date of randomization + 1	Censor
Ineligible for surgical exploration	EFS = 24 weeks (168 days)	Event
Eligible but R0/R1 resection not achieved	EFS = Date of surgical exploration or date of surgery cancelation – Date of randomization + 1	Event
Disease progression or death after achieving R0 or R1 resection	EFS = PFS	Censor or event the same as PFS

Alternatively, EFS may be defined as the time from randomization to death, or objective tumor progression, or failure to achieve an R0 or R1 resection, whichever occurs first. For subjects who did not progress at EOT and did not achieve an R0 or R1 resection (regardless of meeting the surgical eligibility criteria or not), EFS is defined as an event on day 168 (week 24). A more specific algorithm is presented in the table below.

Event Timing	EFS	Censor or Event
Death or disease progression before Day 168	EFS = PFS	Event
Discontinued early and had no follow up progression record	EFS = latest date of (last post-baseline CT, last post-baseline PET scan, last dose) – Date of randomization + 1	Censor
Discontinued treatment early and disease progression occurred after Day 168	EFS = 168 days	Event
Completed 6 treatment cycles and R0 or R1 resection not achieved	EFS = 168 days	Event
Death or disease progression after achieving R0 or R1 resection	EFS = PFS	Censor or event the same as PFS

4.2.4 CT Tumor Measurement

Per RECIST 1.1 criteria, up to 10 measurable target lesions are selected at baseline. The longest diameter (LD) for each target lesion is measured. The sum of the LDs of the target lesions, as well as the percent change from the baseline sum LD and the percent change from the smallest sum LD are derived. In this study, only the lesion at the pancreas will be evaluated. The two types of percent changes are used to determine treatment response per RECIST (1.1) criteria.

$\% \text{ change from baseline} = (\text{LD of the lesion at the pancreas} - \text{Baseline}) / \text{Baseline} * 100\%$

Partial response is defined as at least a 30% decrease from baseline.

$\% \text{ change from the smallest sum} = (\text{LD of the lesion at the pancreas} - \text{the smallest measure during the study}) / \text{the smallest measure during the study} * 100\%$

Progressive disease is defined as an absolute increase of at least 5 mm and at least a 20% increase in the sum of diameters of target lesions, taking as reference the smallest sum on study, or the appearance of one or more new lesions.

Non-target lesions are identified and recorded at baseline and are assessed as ‘improved’, ‘stable’, ‘worsened’, or ‘new’ at follow-up visits.

4.2.5 RECIST (1.1) Response and Best Overall Response

Radiological treatment response is determined by radiologists at each site using RECIST 1.1 criteria (Eisenhauer, et al. 2009). Appendix III presents an outline of the RECIST response criteria for target lesions, non-target lesions, and overall response.

Best overall response is the most favorable overall response recorded during the 24-week treatment period, determined by the following order: CR, PR, SD, and PD.

The best objective response is defined as the best response being CR or PR.

4.2.6 CA 19.9: Percent Change from Baseline, Response, and Normalization

CA 19.9 endpoints are defined only in subjects with baseline CA19-9 value higher than the upper limit normal (ULN). ULN is defined as 37 U/mL in this study (Ballehaninna and Chamberlain, 2012). Baseline CA19.9 is defined as the assessment on Day 1 prior to study drug dosing. If Day 1 assessment is missing, then the last assessment prior to dosing is used.

Percent change from baseline is defined as:

$\% \text{ Change from Baseline} = (\text{Assessment at a given time point} - \text{Baseline}) / \text{Baseline} * 100$

The protocol defined CA 19.9 response is reduction from baseline $\geq 50\%$. Additional response criteria, such as reduction from baseline $\geq 70\%$, $\geq 90\%$, and $\geq 95\%$ will be evaluated.

For each subject, best CA19-9 response is defined as the minimum percent change from baseline. (The largest reduction or smallest increase.)

CA 19.9 normalization is defined as CA 19.9 < ULN for those whose baseline value \geq ULN.

4.2.7 PET SUVmax: Percent Change from Baseline, Response, and Normalization

Maximum standardized uptake value (SUVmax) is read by radiologists at each site. Metabolic response is defined as reduction from baseline in $\text{SUV}_{\text{max}} \geq 30\%$ for lesion at the pancreas only. A complete metabolic response is defined as $\text{SUV}_{\text{max}} = 0$.

4.2.8 Exceptional Treatment Response

Exceptional treatment response is defined:

- CA19.9 reduction from baseline $\geq 95\%$ or normalization at any time point.
- PET SUV normalization, i.e., SUV = 0 at EOT

4.2.9 ECOG Performance Status

Assessment on Day 1 or the last assessment prior to first dose is defined as the baseline. The ECOG grading scale is shown below:

GRADE	ECOG PERFORMANCE STATUS
0	Fully active, able to carry on all pre-disease performance without restriction
1	Restricted in physically strenuous activity but ambulatory and able to carry out work of a light or sedentary nature, e.g., light house work, office work
2	Ambulatory and capable of all self-care but unable to carry out any work activities; up and about more than 50% of waking hours
3	Capable of only limited self-care; confined to bed or chair more than 50% of waking hours
4	Completely disabled; cannot carry on any self-care; totally confined to bed or chair
5	Dead

4.2.10 Adverse Events

The definitions of adverse events (AE), serious adverse events (SAE), severity, and relationship to study medication are described in Section 8.4 of the protocol. Adverse events are coded using the Medical Dictionary for Regulatory Activities (MedDRA) for system organ class (SOC) and preferred term for summary.

Treatment Emergent Adverse Event in the Study (TEAE)

TEAEs are defined as new or worsening AEs that occurred in the window of first dose of any study drug (Day 1) and within 28 days of the last dose of study drug or the day before surgery, whichever occurs first.

Definition of TEAE and Study Period When AE Occurred

AE Onset Date Relative to Dosing*	AEPRIOR	TEAE	Study Period (EPOCH)
-----------------------------------	---------	------	----------------------

AE onset date < Day 1	Any	No	screen
AE onset date = Day 1	Yes	No	screen
	No	Yes	treatment
Day 1 < AE onset date ≤ date of last dose in randomized treatment period	Any	Yes	treatment
Date of last dose < AE onset date ≤ earlier date of {last dose + 28, date of surgery - 1}	Any	Yes	Within 28 days from last dose follow-up
AE onset date > date of last dose + 28	Any	No	Additional Follow-up beyond 28 days
*First dose refers to first of any one of the study medications; last dose refers to last of any one of the study medications.			

AEs may be recorded in multiple records due to changes of characteristics such as seriousness, severity, frequency, treatment, and other aspects. These records will be linked together for summary purposes. The most severe grade, seriousness, relatedness to study drug will be used in AE summary tables. Below are the rules:

- (1) Multiple AE records are linked as one AE if they have the same SOC/preferred term and onset date of a later record is the same as stop date of the previous record.
- (2) If the linked AE starts in the TEAE window, it is a TEAE.
- (3) If the linked AE starts in screening, then the post dose record will be compared with the screening record. It is a TEAE if it is worsened comparing to the screening record. 'Worsened' is defined as increased grade in seriousness, severity, and frequency.
- (4) Once an AE record is determined to be a TEAE, all subsequent records of the linked AE, including the records past 28 days after last dose, are considered TEAE. The highest grade of severity and seriousness of the linked TEAE will be included in the AE summary tables.

The hypothetical example below illustrates the algorithm. Of note, AE #1 occurred in Screening; AE #2 and #3 were not worse than AE #1; these 3 AEs are not TEAEs. AE#4 worsened during the TEAE window; the subsequent AE #5 is also a TEAE by the algorithm.

Example of Definition of TEAE for Multiple Records of the Same AE

AE #	Onset	AE	Severity	Seriousness	Related to study drug?	TEAE
1	Screening	Headache	2	N	N	N
2	Day 2	Headache	1	N	N	N

3	Last dose	Headache	2	N	Y	N
4	10 days post last dose	Headache	3	Y	N	Y
5	45 days post last dose	Headache	5	Y	N	Y
6	Day 10	Headache	2	N		
Consolidated for summary		Headache	5	Y	Y	Y

AE Severity Rating

AEs are rated by the investigators as “mild”, “moderate”, “severe”, “life threatening”, or “fatal” based on the CTCAE v4.0 grading system as described in Section 8.4.3 of the protocol. If a subject reports multiple occurrences of an AE within one system organ class or preferred term, the most severe occurrence will be presented in the summary by severity rating. Missing severity rating will not be imputed and it is ranked the lowest in summary by severity rating.

AEs Related to FG-3019, or Gemcitabine, or Nab-paclitaxel

Investigators determine the relationship of AEs with the study medications as described in Section 8.4.4 of the protocol. For summary purposes, ‘Possibly related’ AEs are grouped with the ‘Related’ AEs. If related and unrelated occurrences of an AE within one system organ class or preferred term are reported in a subject, the related occurrence will be presented in the summary by relationship. Missing relationship evaluation will not be imputed and it is ranked the lowest in summary by relationship.

Anaphylactic Reactions and Hypersensitivities

Anaphylactic reactions and hypersensitivities are defined as the TEAEs with matching preferred terms of the Standardized MedDRA Queries (SMQ) (Appendix IX).

The infusion reaction tick box completed by the sites on the AE CRF will not be included in the summary.

4.2.11 Surgical Safety Measures and Post-surgery Follow-up

Surgical safety assessed during surgery

- Duration of the surgery
- Estimated blood loss (EBL) during surgery
- Any blood transfusion
- Amount of blood transfusion

Surgical complication during hospitalization

- Duration of hospitalization post-surgery (date of discharge – date of surgery + 1)

- Intra-abdominal abscess
- Postoperative leaks
- Surgical site wound infection (SSI)
- Other complication

Surgical complication during post-operative follow-up

- Re-admission to hospital within 30 days from discharge
- Return to operating room within 30 days from discharge
- Intra-abdominal abscess
- Postoperative leaks
- Surgical site delayed wound healing and or infection (SSI)
- Other complication

4.2.12 Laboratory Evaluations

The lab tests in the following table are evaluated at every clinic visit to assess treatment tolerability.

Laboratory Tests

Hematology Panel:	Chemistry Panel:
Hemoglobin (Hb)	Sodium
Hematocrit (HCT)	Potassium
Erythrocytes	Blood urea nitrogen (BUN)
Mean corpuscular volume (MCV)	Bicarbonate
Leukocytes (WBCs)	Creatinine
Neutrophils	Glucose
Lymphocytes	Calcium
Monocytes	Phosphorus
Eosinophils	Aspartate aminotransferase (AST)
Platelets	Alanine aminotransferase (ALT)
Absolute neutrophil count (ANC)	Alkaline phosphatase (ALP)
	Total bilirubin

Investigators review and determine whether the lab results are normal or abnormal, and if abnormal, whether they are clinically significant (CS) or not clinically significant (NCS).

In addition to investigators' assessments, CTCAE grading is used to evaluate the lab results. CTCAE grade 3 or higher is considered potentially clinically significant. The CTCAE 4.03 (June 14, 2010) grading for the study-specific tests is presented in Appendix VI.

Instead of using the reference ranges provided by individual local labs, a common set of reference ranges for chemistry and hematology presented in the New England Journal of Medicine 2004; 351: 1548-63, is used in this study. These reference range are presented in Appendix V.

Baseline for lab tests is defined as the last assessment prior to the first dose.

4.2.13 Vital Signs

Vital sign parameters include systolic/diastolic blood pressures, pulse rate, respiratory rate, and body temperature. Baseline for vital signs is defined as the average of the last Screening value and the Day 1 pre-infusion measurement.

Potentially clinically significant vital sign changes are those which meet both criteria in the table below.

Potentially Clinically Significant Changes in Vital Signs

Parameter	Observed Values	Change from Baseline
Systolic BP (mm Hg)	<90 or >140 mm Hg	>20 mm Hg
Diastolic BP (mm Hg)	<50 or >90 mm Hg	>20 mm Hg
Pulse Rate (bpm)	<50 or >100 bpm	>20 bpm
Body temperature (°C)	37	>3

BP=blood pressure; bpm=beats per minute; °C = Celsius

4.2.14 Physical Examination

At each clinic visit, investigators perform physical examination for the following body systems: general appearance, HEENT, lungs, heart, chest and back, abdomen, genitourinary, extremities, neurologic, and skin. Investigators determine whether the body systems are normal or abnormal. For the abnormal cases, descriptions are provided and clinical significance is identified. Baseline is defined as the findings on Day 1 prior to first dose.

4.2.15 ECG

ECG is evaluated by investigator as:

Normal,

Abnormal, CS, or

Abnormal, NCS.

Abnormal findings are described.

4.2.16 PK

Plasma samples are collected from subjects assigned to treatment Arm A for evaluation of FG-3019 concentration level at the following time points. Post-dose samples are collected between 30 minutes to 2 hours from the end of infusion of FG-3019.

CYCLE	DAY	TIME
1	1	Pre-dose
		Post-dose
	8	Pre-dose
		Post-dose
	15	Pre-dose

		Post-dose
2	15	Pre-dose
3	1	Pre-dose
		Post-dose
5	1	Pre-dose
6	1	Pre-dose
		Post-dose
	28/EOT	
Safety FU	28 days post last dose	

4.2.17 CTGF

CTGF samples are collected on Day 1 predose and at safety follow-up visit (70 days after the last dose for subjects enrolled under the original protocol or 28 days after the last dose for subjects enrolled after Amendment 1). CTGF parameters include CTGF N+W and CTGF W.

4.2.18 Protein Expression Profile and PAX Gene

Plasma samples for protein expression profile and whole blood samples for PAX gene evaluation are collected on Day 1 predose of Cycles 1, 3, 5 and at EOT. Analysis of these data will be provided in separate reports.

4.2.19 Tumor Biopsy Evaluation

Analysis of the tumor biopsy samples will be provided in a separate report.

4.2.20 HAHA

Blood samples for human anti-human antibodies (HAHA) measurement are collected from subjects randomized to treatment Arm A in heparinized tubes at pre-dose on Day 1 of Cycle 1 and at the safety follow-up visit approximately 28 days after the last dose with study treatment. HAHA assay results are “Reactive” or “Non-Reactive”. For those with ‘Reactive’ results, further evaluation of “Specific” or “Not Specific”, and % Inhibition by HAHA are provided.

5 GENERAL STATISTICAL CONSIDERATIONS

5.1 Analysis Populations

The Randomized Population includes all subjects who have received a treatment arm assignment, regardless of whether they received study treatment.

In this study, the Intent-to-Treat (ITT) Population and the Safety Population are both defined as all randomized subjects who have taken any amount of study drugs including FG-3019, or gemcitabine, or nab-paclitaxel. The Per Protocol (PP) Population is defined as subjects who have completed the study treatment, as defined on the End-of-Treatment Disposition CRF. If actual treatment received differs from the randomized treatment arm, the actual treatment arm will be used in all data analysis.

The ITT Population will be used in baseline and efficacy summaries; the Safety population will be used for safety summaries. The PP Population will be used in selected efficacy parameters.

5.2 Hypotheses and Decision Rules

Data in this exploratory study will be examined in a hypothesis-generating manner via various statistical methods and models. Adjustment for Multiple Comparisons

No adjustments will be made for multiple tests.

5.3 Handling of Dropouts or Missing Data

Except for the cases described in this section, only observed data without imputation will be used in analyses.

5.3.1 Handling Missing Data in Responder Analyses

When performing a responder analysis, subjects with missing data will be considered non-responders. All subjects in the defined population will be included in the evaluation.

5.3.2 Handling Missing/Incomplete AE Onset Date

If the AE onset date is incomplete or missing, the following rules will be applied to obtain imputed AE onset date.

- If year and month are present, only day is missing,
 - a) If AE onset Year/month = Day 1 Year/month, assign onset date = date of Day 1;
 - b) If AE onset Year/month \neq Day 1 Year/month, assign onset Day = 1;
- If year is present, month and day are missing,
 - a) If onset year = year of Day 1, assign onset date = date of Day 1;
 - b) If onset year \neq year of Day 1, assign January 1st to onset month and day.
- If onset date is completely missing, assign onset date = date of Day 1.

If the stop date is complete and the imputed start date as above is after the stop date, the start date will be imputed by the stop date.

5.3.3 Handling Missing/Incomplete CM Start/Stop Dates

For the purpose of grouping medications in different study period, the following rules will be used to impute incomplete CM start and end date:

- Incomplete CM start date: assign 1 to missing Day, January to missing Month.
- Incomplete CM end date: assign 30 to missing Day, December to missing Month. Impute CM end date only if 'ONGOING' is not checked.

If the imputed stop date is before the start date (imputed or non-imputed start date), then the imputed stop date will be replaced with the start date.

No imputation will be performed for the following cases:

- CM end date will not be imputed if 'ONGOING' is checked.
- Year of CM start or end is missing. If CM end year is missing, the CM will be grouped in the 'Concomitant Medication' category.

5.4 Adjustment for Covariates

Comparisons between treatment arms will be adjusted for the following factors.

- Baseline TNM Stage (T3 vs.T4)
- Baseline TNM Stage (N0 vs. N1/NX)
- Baseline ECOG (0 vs. 1)
- Baseline CA 19.9 (below vs. above median)
- Baseline PET SUV (below vs. above median)
- Baseline tumor size (below vs. above median)
- Baseline SMA Involvement (> 180 degree vs. <180 degree)
- Baseline celiac Abutment Status (Yes vs. No)
- Baseline unreconstructible SMV/Portal Occlusion Status (Yes vs. No)
- Tumor anatomy (head vs. body/tail)

5.5 Definition of Baseline

Unless specified otherwise, baseline is defined as the last evaluable observation collected prior to first dose of study drug.

5.6 Study Day Calculation

The day when a subject receives the first dose of any study drugs (FG-3019, gemcitabine, or nab-paclitaxel) after randomization is designated as Day 1.

Study day of an assessment/procedure is calculated as follows.

- For assessments or procedures on Day 1 or later,
Study day = assessment/procedure date – Day 1 date + 1.
- For assessments or procedures earlier than Day 1,

Study day = assessment/procedure date – Day 1 date.

5.7 Efficacy Analysis Visit Window

Efficacy assessments will be summarized by analysis visit based on actual date of assessment. Visit windows will have the widths of the corresponding assessments centered at the scheduled time, as shown in the table below, except for the upper bound of the Week 24 window, which is set on Week 30 (Day 210)

Assessment Interval	Assessment	Target Assessment	Window
4 weeks	CA19-9	k-th Cycle Day 1	Target day $[4*(k-1)*7 + 1] + 13/-14$ days For EOT $k = 7$.
8 weeks	CT, ECOG	k-th Cycle Day 1	Target day $[4*(k-1)*7 + 1] + 27/-28$ days For EOT $k = 7$.
24 weeks	PET	EOT	Day 85 - 210

All scheduled and unscheduled assessments are included. If two assessments are available in the same window, the one closer to the target day is used. If two assessments with equal distance to the target day, the later assessment will be used in analysis. Assessments that do not fall in any windows will not be included in analyses.

Safety data (lab tests, vital signs, ECG, physical exams) are summarized by nominal clinic visit. Data collected during the unscheduled visits are not included in the summary tables, but are included in data listings and in evaluations for clinically significant abnormal changes.

5.8 Interim Analyses and Data Monitoring

In this open label study efficacy and safety data are monitored periodically throughout the study.

5.9 Pooling Data of Study Sites

All study sites are pooled in all analyses due to the small number of subjects enrolled at each site.

5.10 General Layout

All study parameters, including baseline characteristics, efficacy, safety, PK, and biomarkers, will be summarized descriptively. Descriptive statistics including the number of subjects (n), mean, standard deviation (SD) for baseline variables or standard error (SE) for efficacy parameters, median, minimum and maximum will be presented for continuous variables. For continuous PK parameters, coefficient (CV) and geometric mean may also be presented. Number (n) and percentage (%) of subjects in each category will be summarized for categorical variables.

Efficacy parameters will be summarized analytically. Analytical statistics include LS mean and standard error, 95% CI for the mean or median, p-value. Depending on the nature of the parameter, treatment difference will be expressed in absolute or percent difference, odds ratio, or hazard ratio.

All summaries except PK will be presented by treatment group as well as a combined group (Arm A, Arm B, Overall), unless specified otherwise.

Raw data and derived parameters will be presented in data listings, which will be organized in the order by treatment arm followed by subject ID.

Figures, such as line-chart, bar-chart, box-plot, scatter plot, forest plot, or waterfall plot, are in general included to facilitate comparison between treatment arms and evaluation of trend.

5.11 Data Errata and Hard-Coding

Data errors identified after database lock are documented in an errata log. Under special circumstances, hard-coding in SAS programs is necessary to correct data errors in order to avoid obscure study results. Only limited cases deemed necessary and are approved by study team will be hard-coded. The changed values as well as approval from the study team will be documented.

6 STATISTICAL ANALYSES

6.1 Subject Enrollment and Disposition

The number of subjects enrolled in each study site will be summarized.

The number of subjects in each study population (Randomized, ITT/Safety, and PP) will be summarized by treatment group. The number of subjects who completed or discontinued the study as well as the reasons for early discontinuation will be summarized.

The number of subjects remaining in the study at each visit will be summarized.

Subject who discontinued the study early and the reasons will be listed.

6.2 Protocol Deviations

Protocol deviations will be categorized as follows:

- Entry Deviation: Subject entered study, but did not satisfy eligibility criteria.
- Withdrawal Deviation: Subject met withdrawal criteria during the study but was not withdrawn.
- Dosing Deviation: Subject received the wrong treatment or incorrect dose; including incorrect timing of a dose.
- Prohibited Medication Deviation: Subject received an excluded concomitant treatment.
- Operational Deviation: All other deviations; including, but not limited to: informed consent form-related deviations other than consent not obtained, IRB/IEC approval expired, study drug not stored under protocol-specified conditions, missing laboratory report, out-of window visit, etc.; includes missed visits, and subject refusal of a study procedure or procedures.

Deviations are summarized by subject count and by event count, by cohort and by study site. All recorded protocol deviations are listed. Reported items that are not considered as protocol deviations are not included in summary tables and data listings, but remain in the database for reference.

6.3 Demographics and Baseline Disease Characteristics

Demographics and baseline characteristics will be summarized descriptively for subjects in the ITT population. Each parameter will be presented in data listings.

6.3.1 Subject Demographics and General Characteristics

Demographic parameters and general characteristics include age, age group (< 18, 18-64, 65-74, ≥ 75), sex, ethnicity, race, height, weight, body-mass index (BMI), BSA, smoking status, and number of years that subject smoked.

Age is defined as the age on the day of signing informed consent:

age = INTCK('YEAR', Birth Date, Date of Informed Consent, 'C')

where INTCK is a SAS function.

BSA and BMI are defined below:

$$\text{BSA in m}^2 = \text{Sqrt}(\text{weight in kg} * \text{height in cm} \div 3600)$$

$$\text{BMI} = \text{Weight (kg)} / (\text{Height (m)})^2$$

Smoking pack-years is defined as average packs per day * duration (years).

6.3.2 Baseline Disease Stage

Baseline disease and disease stage include the following variables:

- Tumor name (“PDAC” or “Other”)
- Time from diagnosis (days) defined as date of first dose - date of diagnosis + 1
- Disease stage (1, 2a, 2b, 3, or 4)
- TNM stage
- ECOG performance status

6.3.3 Baseline Tumor Unresectability per NCCN Criteria

The following unresectability characteristics are summarized:

- Does the tumor have greater than 180 degrees SMA encasement
- Does the tumor have any celiac abutment
- Does the tumor have inferior vena cava
- Does the tumor have the unreconstructible SMV/portal occlusion
- Does the tumor have aortic invasion or encasement

6.3.4 Baseline Tumor Burden

Baseline tumor burden will be characterized by the following variables:

- Size of target lesions
- Presence of non-target lesions
- FDG-PET SUV_{max}
- CA19-9

6.3.5 Medical History

Medical conditions, including allergies and surgeries, are coded in system organ class (SOC) and preferred term (PT) using MedDRA.

The medical conditions will be tabulated by SOC and PT. A subject with multiple medical conditions within an SOC is only counted once in this SOC. Similarly, a subject with multiple medical conditions within a PT is only counted once in this PT. The tabulation will be sorted alphabetically by SOC and by decreasing order of frequency of PT within each SOC based on the ITT Population.

6.4 Prior and Concomitant Medications

The World Health Organization Drug Dictionary will be used to classify concomitant medications by therapeutic class and generic name.

Prior and concomitant medications, defined below, are summarized by ATC class and preferred term.

1. Prior medications are those that are stopped prior to the first infusion.
2. Concomitant medications are those that are used concomitantly with the study drug, which are defined as medications that are not stopped before the first infusion.

Subjects reporting more than one use of the same medication will be counted only once in the summary tables.

Prior cancer therapies including systemic, surgical, and radiation will be summarized descriptively and presented in data listing.

All medications and non-drug therapies are presented in data listings.

6.5 Study Drug Exposure and Treatment Compliance

6.5.1 Study Drug Exposure

Study drug exposure will be characterized by the following measures:

- Duration in days from first dose of any study drug to last dose of any study drug
- Number of infusions
- Average dose amount in mg over the entire study period for individual subjects

During the course of 22 weeks of treatment duration (6 treatment cycles, last dose on Day 15 of Cycle 6), subjects in Arm A will receive up to 13 FG-3019 infusions, with 3 infusions administered during the first cycle and 2 during each of the remaining 5 cycles. All subjects will receive up to 18 infusions of gemcitabine and up to 18 infusions of nab-paclitaxel.

FG-3019 dose amount in mg = 35 * body weight in kg.

Gemcitabine dose amount in mg = 1000 * BSA in m².

Nab-paclitaxel dose amount in mg = 125 * BSA in m².

These measures will be summarized descriptively for the Safety population. Study drug administration log will be listed.

6.5.2 Treatment Compliance

Compliance will be calculated as the number of doses the subject received divided by the number of doses the subject is scheduled to receive during the participation in the study.

Compliance = Actual doses received / Scheduled doses while actively in study * 100%

Scheduled doses of the study drugs are listed in the table below.

Day within a Cycle	Number of FG-3019 Doses / Number of Gemcitabine/Nab-paclitaxel Doses					
	Cycle 1	Cycle 2	Cycle 3	Cycle 4	Cycle 5	Cycle 6
1	1/1	4/4	6/7	8/10	10/13	12/16
8	2/2	4/5	6/8	8/11	10/14	12/17
15	3/3	5/6	7/9	9/12	11/15	13/18

Compliance will be summarized in the following categories; 100%, 90-99%, 80-89%, and less than 80 percent.

6.6 Efficacy Analyses

All efficacy analyses described in this section will be based on the ITT Population.

6.6.1 Analysis of Eligibility for Surgical Exploration

Surgical eligibility is defined in Sections 4.2.1. Proportion of subjects in the following categories will be summarized by treatment group:

- met the protocol defined criteria with no contraindication,
- met the protocol defined criteria with contraindication,
- did not meet the protocol defined criteria

Exact 95% CIs for the point estimates as well as the treatment difference will be obtained from SAS PROC FREQ procedure with EXACT option. The two treatment arms will be compared using the Cochran-Mantel-Haenszel (CMH) test controlling for the baseline factors listed in Section 5.5.

6.6.2 Analysis of Resection Outcome

The proportion of subjects achieving different resection outcomes will be summarized descriptively. The CMH test controlling for the baseline factors listed in Section 5.5 is used to compare the two treatment arms.

6.6.3 Analysis of Time-to-Event Parameters

Time-to-event endpoints include overall survival (OS), progression free survival (PFS), and event free survival (EFS). Comparisons between the two treatment arms will be characterized using the following statistics:

- Hazard ratio (HR) and its 95% CI
- Median time-to-event and its 95% CI
- Proportions of subjects with 1-year, {2-year, and 3-year} event-free and the corresponding 95% CIs
- Restricted mean survival time (RMST) over time

The Cox regression model, with adjustment of the baseline factors listed in Section 5.5, will be used to compare the 2 treatment arms and to estimate the hazard ratio and the 95% CI.

proc phreg;

model {time-to-event variable} * censor (1) = ARM baseline-factor;

hazardratio ARM /CL = PL;

run;

Baseline factors are included in the model one at the time. Adjustment of multiple factors may be explored.

Median time-to-event and its 95% CI will be estimated by Kaplan-Meier method using SAS PROC LIFETEST procedure. 95% CI for the event-free rates will be estimated using the Clopper-Pearson method by PROC FREQ with option BINOMIAL (EXACT).

```
proc lifetest;
```

```
    time {time-to-event variable} * censor (1);
```

```
    strata ARM;
```

```
run;
```

Comparison between resected versus non resected subjects, with resection status being used as time dependent covariate, will be performed.

Since the proportional hazards assumption may be violated in EFS, additional analysis using the accelerated failure time model will be performed.

6.6.4 Analysis of RECIST Response and Change in Tumor Measurements

The best RECIST response defined in Section 4.2.5 will be compared between the two arms using the exact procedures described in Section 6.6.1. RECIST response by visit will be summarized descriptively.

Absolute change and percent change from baseline in the sum of the longest diameters (LD) of the target lesions are summarized by visit and treatment arm. Percent change from baseline within each arm is evaluated using one-sample Wilcoxon signed-rank test. Comparison between the two treatment arms is based on two-sample Wilcoxon rank-sum test. Nonparametric methods are used due to the skewed distribution of the percent change data.

Tumor reduction $\geq 30\%$ is also summarized by visit and treatment arm. Comparison between treatment arms will be based on Fisher's exact test.

Status of non-target lesions will be summarized by visit.

6.6.5 Analysis of Change in CA19-9 and CA19-9 Response

CA 19-9 responses defined in Section 4.2.6 will be summarized descriptive by visit. Comparison between the two arms in the dichotomous parameters will be based on the exact procedures described in Section 6.6.1. Ordered categories will be compared using chi-square test with ordinal outcome.

Change from baseline will be evaluated using the one-sample Wilcoxon signed rank test. Comparison between the two arms will be evaluated using the two sample Wilcoxon rank sum test. Nonparametric methods are used due to the skewed distribution of the percent change data.

6.6.6 Summary of Change in FDG-PET SUVmax and PET Response

PET responses defined in Section 4.2.7 will be summarized descriptive. Comparison between the two arms in the dichotomous parameters will be based on the exact procedures described in Section 6.6.1.

Change from baseline will be evaluated using the one-sample Wilcoxon signed rank test. Comparison between the two arms will be evaluated using the two sample Wilcoxon rank sum test. Nonparametric methods are used due to the skewed distribution of the percent change data.

6.6.7 ECOG

Change from baseline in ECOG status will be summarized in shift table of post-baseline versus baseline.

6.6.8 Disease Stage

Disease TNM stage and cancer staging will be summarized in shift table of EOT versus Screening.

6.6.9 NCCN Resectability

NCCN resectability condition will be summarized in shift tables of EOT versus Screening.

6.7 Exploratory Analyses

As an exploration, event free survival will be analyzed using Cox regression model with change from baseline in CA19.9 as time-dependent covariate. Further exploratory analyses will be documented in a separate analysis plan.

6.8 Safety Analyses

Safety analyses will include summary of adverse events, surgical safety parameters, lab test results, vital signs, ECGs, and physical exams. In general, safety data will only be summarized descriptively and no formal inferential statistical test will be applied.

6.8.1 Adverse Events

All reported AEs will be presented in listings. Treatment emergent adverse events (TEAEs), as defined in Section 4.2.10, will be summarized by treatment arm, system organ class (SOC) and preferred term (PT). The summary tables will be sorted alphabetically by SOC and by decreasing order of frequency of PT within each SOC, based on the rate of both treatment arms combined. A subject with multiple adverse events within a SOC is only counted once in this SOC. Similarly, a subject with multiple adverse events within a PT is only counted once in this PT.

In addition to listing of all reported AEs and summary of all TEAEs, the following subgroups of AEs will be summarized and listed separately:

- SAEs (listing) for both arms and relationship to the IP (whether related or unrelated)
- TEAEs and fatal SAEs (tables)
- TEAEs and fatal SAEs
- TEAEs leading to study or treatment discontinuation
- TEAE with incidence $\geq 10\%$ in the overall safety population
- TEAE occurred in 2 or more subjects in either arm
- AEs with severity grade ≥ 3
- AEs with severity grade ≥ 3 that are possibly related to FG-3019 (as assessed by investigators)

6.8.2 Mortality

Deaths as an outcome of TESAEs will be summarized as a safety measure. Long-term all-cause mortality events are captured as an efficacy measure and will be summarized in the efficacy section.

6.8.3 Laboratory Data

Laboratory test results and change from baseline are summarized descriptively by visit.

CTCAE grade 3 or higher lab test results will be considered potentially clinically significant. These results are presented in a data listing.

Shift tables to summarize changes from baseline to each visit in CTCAE categories are tabulated. Shift from baseline to most severe CTCAE category during the study is also summarized. In addition, shift-tables based on investigators' assessments are also presented.

An eDISH (evaluation of drug induce severe hepatotoxicity) analytical graph, which is a scatter plot of maximum observed total bilirubin versus maximum observed ALT or AST, will be generated to identify cases in Hy's law range.

Box-plots for selected lab tests by visit are presented to evaluate trend.

Lab data are summarized by nominal clinic visit. Data collected during the unscheduled visits are not included in the summary tables, but are included in data listings and in evaluations for potentially clinically significant abnormal changes.

6.8.4 Vital Signs

Vital sign observed values and change from baseline are summarized descriptively by visit and by treatment arm.

In addition, potentially clinically significant changes from baseline, which are defined in Section 4.2.13, are tabulated for each visit.

All measurements and change from baseline are presented in data listings. Potential clinically significant changes from baseline are flagged.

Box-plots of vital sign measurements by visit, pre- and post-infusion separately, are presented to evaluate trend.

6.8.5 Physical Examinations

Findings from physical examinations will be summarized in shift tables that include cross-tabulation of 'Normal', 'Abnormal NCS', 'Abnormal CS' at each visit versus baseline. The number and percent of subjects with "Normal", "Abnormal", "Not Done", and "Missing" physical examination results are summarized by body system, cohort, and visit. Shift table of changes from baseline will also be summarized by visit.

The data listing includes abnormal findings.

6.8.6 ECG Data

ECG findings will be summarized in shift tables that include cross-tabulation of investigator assessments of 'Normal', 'Abnormal NCS', 'Abnormal CS' at EOT versus Screening.

Detailed findings are presented in the data listing.

6.8.7 Surgical Safety

Surgical safety measures described in Section 4.2.11 will be summarized descriptively for subjects who have undergone a R0 or R1 resection. Subjects who have undergone surgical exploration but resection was not attempted will be summarized separately.

6.9 Analysis of PK, CTGF, and HAHA

6.9.1 Summary of PK Data

PK concentration level will be summarized descriptively by time point. It will be plotted over time for each subject.

6.9.2 Summary of CTGF Data

CTGF – N+W and CTGF – W are summarized descriptive by scheduled time point.

6.9.3 Summary of HAHA Data

Number of subjects with reactive and specific antibodies to FG-3019 is summarized. HAHA data are also presented in a data listing.

6.9.4 Summary of Biomarker Data

Biomarker data will be presented in a separate report.

7 VALIDATION AND QUALITY ASSURANCE

All datasets (SDTM, ADaM), tables, listings, figures are programmed by two programmers independently. The results must be 100% match.

Both primary and validation programmers will develop programs independently based on the specifications and/or SAP. If the outputs are datasets, the final outputs are compared. If the outputs are TLFs, benchmark results are generated and compared. The validation findings and resolutions are documented on a Validation Worksheet.

The detail process for validation and quality assurance is documented in the Standard Operating Procedures of Q2, Inc.

8 REFERENCES

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New England Journal of Medicine 2004; 351: 1548-63, for lab value reference range

APPENDIX I GENERAL SPECIFICATIONS FOR SUBMISSION DATA

1. Study Data Tabulation Model (SDTM)

Raw datasets will be mapped to SDTM datasets following variable names and attributes specified in the SDTM Implementation Guide (version 3.2 and subsequent update). Data coding is also mapped to the SDTM controlled terminology. OpenCDISC module will be run to confirm compliance to the SDTM guidelines. Deviations from the guidelines will be documented. Variables that exist in different source datasets, such as lab data collection date, will be consolidated and included once in the SDTM datasets. Unscheduled 'Not Done' records are excluded from the SDTM datasets. Detailed mapping specifications are documented in Define.xml and annotation CRFs.

CRF variables that are not included in the SDTM datasets will be marked as 'Not Submitted', since the SDTM datasets will be submitted as the raw datasets and the raw datasets that match the CRFs will not be submitted to regulatory agencies.

The SDTM datasets are the basis to derive the analysis datasets. Summary tables and listings that do not require complex derived variables may be generated directly from SDTM datasets.

Table A contains a list of SDTM datasets to be created for the study. The corresponding supplemental datasets are not included. This set of SDTM datasets includes only data for the Randomized Treatment Period as described in Section 1.

Table A. Study Data Tabulation Model Datasets (SDTM)

SDTM Domain	SDTM Domain Description	SDTM Domain Structure	General Observation Class	Source Data Used	Key Variables
AE	Adverse Events	One record per adverse event per subject	Events	AE	STUDYID, USUBJID, AESTDTC, AEENDTC, AEDECOD, AESPID
CM	Concomitant Medications	One record per recorded medication occurrence per subject	Interventions	CM	STUDYID, USUBJID, CMCAT, CMTRT, CMDECOD, CMSTDTC, CMENDTC, CMSPID
DD	Death Details	One record per subject	Findings	DTH	STUDYID, USUBJID, DDTESTCD, DDDTC
DM	Demographics	One record per subject	Special Purpose Domains	DM, EX, ICF, DTH, SITE_INV	STUDYID, USUBJID
DS	Disposition	One record per disposition status or protocol milestone per subject	Events	DS, ICF, DTH, EX	STUDYID, USUBJID, DSSTDTC, DSDECOD, DSSPID

EC	Exposure as Collected	One record per protocol-specified study treatment per collected-dosing interval per subject	Interventions	EX	STUDYID, USUBJID, VISITNUM, ECTRT, ECSCAT, ECSTDTC, ECENDTC
EG	ECG Test Results	One record per ECG observation per visit per subject	Findings	EG	STUDYID, USUBJID, EGTESTCD, VISITNUM, EGDTC
EX	Exposure	One record per constant dosing interval per subject	Interventions	EX	STUDYID, USUBJID, VISITNUM, EXTRT, EXSCAT, EXSTDTC, EXENDTC
FA	Findings About Events or Interventions	One record per finding, per object, per time point, per visit per subject	Findings	DISE	STUDYID, USUBJID, VISITNUM, FATESTCD, FAOBJ
IE	Inclusion/Exclusion Criteria Not Met	One record per inclusion/exclusion criterion not met per subject	Findings	IE	STUDYID, USUBJID, IETESTCD
LB	Laboratory Test Results	One record per lab test per visit per subject	Findings	LBC, LBH, LBPREG	STUDYID, USUBJID, VISITNUM, LBCAT, LBTESTCD, LBDTC, LBNAM
MH	Medical History	One record per medical history event per subject	Events	MH	STUDYID, USUBJID, MHSPID, MHTERM
PC	Pharmacokinetic Concentrations	One record per sample characteristic or time-point concentration per reference time point or per analyte per subject	Findings	CA, PKCTGF, PKFG, PKPAX, PKPRO	STUDYID, USUBJID, PCTESTCD, VISITNUM, PCTPTNUM
PE	Physical Examination	One record per body system or abnormality per visit per subject	Findings	PE	STUDYID, USUBJID, PETESTCD, VISITNUM

PP	Pharmacokinetic Parameters	One record per PK parameter per time-concentration profile per modeling method per subject	Findings	eDT	STUDYID, USUBJID, PPTESTCD, PPCAT, PPRFTDTC
PR	Procedures	One record per recorded procedure per occurrence per subject	Interventions	NDT, PCASUR, PCARAD, DISE	STUDYID, USUBJID, PRCAT, PRSPID, PRSTDTC, PRENDTC
QS	Questionnaires	One record per questionnaire per question per time point per visit per subject	Findings	ECOG, KARNOF	STUDYID, USUBJID, VISITNUM, QSCAT, QSTESTCD
RS	Disease Response	One record per response assessment per visit per subject per assessor	Findings	RECIST, DISPROG	STUDYID, USUBJID, RSGRPID, VISITNUM, RSTESTCD
SC	Subject Characteristics	One record per characteristic per subject	Findings	DM	STUDYID, USUBJID
SE	Subject Elements	One record per actual Element per subject	Special Purpose Domains	SDTM.DM, SDTM.SV, SDTM.DS, SDTM.EX, SDTM.SS	STUDYID, USUBJID, TAETORD, SESTDTC
SS	Subject Status	One record per finding per visit per subject	Findings	EOS, LTFU	STUDYID, USUBJID, STESTCD, VISITNUM, SSDTC
SU	Substance Use	One record per substance type per reported occurrence per subject	Intervention	SUTOB	STUDYID, USUBJID, SUTRT
SV	Subject Visits	One record per actual visit per subject	Special Purpose Domains	All datasets which include VISIT information	STUDYID, USUBJID, VISITNUM
TA	Trial Arms	One record per planned Element per Arm	Trial Design	N/A	STUDYID, ARMCD, TAETORD
TD	Trial Disease Assessments	One record per planned constant assessment period	Trial Design	N/A	STUDYID, TDORDER
TE	Trial Elements	One record per planned Element	Trial Design	N/A	STUDYID, ETC
TI	Trial Inclusion/Exclusion Criteria	One record per I/E criterion	Trial Design	N/A	STUDYID, TIVERS, IETESTCD

TR	Tumor Results	One record per tumor measurement/assessment per visit per subject per assessor	Findings	TRNTL, TRTL, PETRES, PETRESUV	STUDYID, USUBJID, TRGRPID, VISITNUM, TRLNKGRP, TRTESTCD, TRLNKID
TS	Trial Summary	One record per trial summary parameter per value	Trial Design	N/A	STUDYID, TSPARMCD, TSSEQ
TU	Tumor Identification	One record per identified tumor per subject per assessor	Findings	TRNTL, TRTL, PETRES, PETRESUV	STUDYID, USUBJID, VISITNUM, TUTESTCD, TULNKID
TV	Trial Visits	One record per planned Visit per Arm	Trial Design	N/A	STUDYID, VISITNUM, ARMCD
VS	Vital Signs	One record per vital sign measurement per time point per visit per subject	Findings	VS	STUDYID, USUBJID, VISITNUM, VSTESTCD
YH	Human Anti-human Antibody Assessment	One record per test per visit per subject	Findings	HAHA, eDT HAHA external data	STUDYID, USUBJID, YHTESTCD, VISITNUM, YHDT, YHGRPID
ZA	Surgical Eligibility Assessment	One record per test per subject	Findings	SA	STUDYID, USUBJID, ZATESTCD, VISITNUM, ZAORRES
ZC	Cancer Pathology	One record per test per visit per subject	Findings	CANPATH	STUDYID, USUBJID, ZCTESTCD, VISITNUM
ZF	Surgical (post-operative) Follow-up	One record per test per visit per subject	Findings	SURFU	STUDYID, USUBJID, ZFGRPID, ZFCAT, ZFTTESTCD, VISITNUM
ZO	Surgery/Surgical Outcomes	One record per test per visit per subject	Findings	SURO	STUDYID, USUBJID, VISITNUM, ZOCAT, ZOTESTCD

2 Analysis Data Model (ADaM)

The ADaM datasets are created based on the SDTM datasets following the ADaM Implementation Guide (version 1.0 and subsequent update). They include derived parameters and flags that are needed to generate tables/listings/figures. ADSL and ADAE are created using the specifications provided in the ADaMIG. Study endpoints are included in analysis files in ADaM Basic Data Structure (BDS). Horizontal analysis files are created for analyses on relationship of multiple endpoints. Variable names and labels of the horizontal files are from PARAMCD and PARAM of the corresponding BDS files. Detail derivations of each variable in the ADaM datasets are documented in the Metadata.

Table B. Analysis Data Model (ADaM)

Dataset	Description	Structure	Keys
ADSL	Subject-Level Analysis Dataset	One record per subject	STUDYID, SUBJID
ADAE	Analysis Dataset Adverse Events	One record per subject per adverse event per start date	STUDYID, USUBJID, AESTDTC, AEENDTC, AEDECOD, AESPID
ADCM	Analysis Dataset Concomitant Medications	One record per subject per recorded medication occurrence	STUDYID, USUBJID, CMCAT, CMTRT, CMDECOD, CMSTDTC, CMENDTC, CMSPID
ADDD	Analysis Dataset Death	One record per subject per parameter	STUDYID, USUBJID, PARAMCD
ADDISE	Analysis Dataset Disease Stage	One record per subject per parameter per analysis visit	STUDYID, USUBJID, PARAMCD, AVISITN
ADEFF	Analysis Dataset Efficacy	One record per subject per parameter per analysis visit per Link ID	STUDYID, USUBJID, PARCAT1, PARAMCD, AVISITN, ADT, VISITNUM, TRLNKID
ADEG	Analysis Dataset for ECG Test Results	One record per subject per analysis visit	STUDYID, USUBJID, PARAMCD, AVISITN, ADT
ADEX	Analysis Dataset Exposure	One record per subject per parameter per analysis visit	STUDYID, USUBJID, PARAMCD, AVISITN
ADLB	Analysis Dataset Laboratory Test Results	One record per subject per category per parameter per analysis visit	STUDYID, USUBJID, PARCAT1, PARAMCD, AVISITN
ADMH	Analysis Dataset Medical History	One record per subject per medical history per time interval of medical history	STUDYID, USUBJID, MHSPID, MHTERM
ADPE	Analysis Dataset Physical Examination	One record per subject per parameter per analysis visit	STUDYID, USUBJID, AVISITN, PARAMCD
ADQS	Analysis Dataset for Questionnaire	One record per subject per category per parameter per analysis visit	STUDYID, USUBJID, PARCAT1, PARAMCD, AVISITN

ADRS	Analysis Dataset Disease Response	One record per subject per parameter per analysis visit	STUDYID, USUBJID, PARAMCD, AVISITN
ADSURG	Analysis Dataset Surgical Assessment	One record per subject per category per parameter per analysis visit per analysis value per group ID	STUDYID, USUBJID, PARCAT1, PARAMCD, AVISITN, AVALC, ZFGRPID
ADTR	Analysis Dataset Tumor Results	One record per subject per category per parameter per analysis visit per Link ID	STUDYID, USUBJID, PARCAT1, PARAMCD, AVISITN, TRLNKID
ADTTE	Analysis Dataset Time-to-Event	One record per subject per parameter	STUDYID, USUBJID, ASEQ, PARAMCD
ADVS	Analysis Dataset Vital Signs	One record per subject per parameter per analysis visit	STUDYID, USUBJID, PARAMCD, AVISITN

APPENDIX II GENERAL SPECIFICATIONS FOR TABLES, LISTINGS, FIGURES

1 Software Used

All programming of tables, listings and figures (TLFs) will be performed using the statistical software package SAS® version 9.3 or greater.

2 General

All TLFs are based on SDTM and/or ADaM datasets. By default, data listings reflect the actual values captured in SDTM and ADaM datasets, including date/time variables and missing values. Except for concatenation of some variables for compact display purpose, data are presented directly with minimum manipulation. In general, the character standard result variables, such as –STRESC, are presented in data listings. Date are presented in listings in format yyyy-mm-dd. For incomplete date, CDISC presentation conversion is followed.

For continuous variables that are recorded as “<X” or “>X”, the value of “X” will be used in the calculation of summary statistics. The value “X” is also captured in the numeric variable in the SDTM datasets as well as in the ADaM datasets for consistency, although SDTMIG recommends capturing missing values in the numeric variables.

In general, reported verbatim, such as terms of AE, medical history, medication names, specifications to the ‘Other’ fields, findings, etc., are presented in upper case. However, when reported fields are long, such as comments and protocol deviation descriptions, listing in lower case enhances readability.

3 Table/Listing/Figure Output File Type and Organization

In general, the final set of TLFs will include both PDF and RTF files. Outputs are combined in several large PDF files, eg, all tables, all listings, and all figures, in the order as in the planned TLF in Section 12. A table of content should be included with hyperlinking to individual outputs. True RTF files (in-text format) will be created for tables and listings. SAS outputs for statistical procedures used in analysis of primary, secondary, and exploratory efficacy endpoints will also be included.

4 Page Layout

All column headers (consisting of one or several words) will start with uppercase and thereafter only lowercase characters, except for acronyms and abbreviations. In case values from the database will be displayed in column headers, they may be displayed as in the database. Pages will be numbered as ‘Page x of y’, where ‘y’ is the total number of pages of the corresponding table or listing. The page specifications are presented in Table A.

Table A. Specifications for Page Layout

Paper Size	Letter
Orientation	Landscape
Alignment	Center
Font size	9
Font type	Courier New (default)
Margins	
Top	0.75"
Bottom	0.38"
Left	0.75"
Right	0.38"

The margin sizes and font size for listings may be flexible to provide sufficient information on a single page to facilitate review and comparison.

When created using SAS, tables and listings will be created using ODS, and output files will be produced in RTF. When RTF files are produced, titles and footnotes will appear as document headers/footers.

5 Titles and Footnotes

All tables and listings will have a header showing “FibroGen, Inc.”, the protocol number, database cutoff date or ‘Final Database’, and Page x of y. A footer will show the program file path/name, output file path/name, run date and time.

All titles are written in title format, with uppercase at the beginning of each word; articles, prepositions, and conjunctions, which are of three characters length or less will start with lowercase letters (Mixed Case). Footnotes are in regular text format.

Titles

In total there are up to 10 titles available, defined as following:

- first title “FibroGen, Inc.” (left aligned) and “Database extraction date: ddMMMyyy” or “Final Database” (right aligned)
- second title protocol number + “Clinical Study Report” (left aligned) and “Page x of y” (right aligned)
- third title blank
- fourth title: table/listing/figure number
- fifth title: table/listing/figure title

sixth title: population names if provided in SAP, or brief definition of specific analysis set

Footnotes

Up to 10 footnote lines are available for tables, listings and figures. Footnotes 1, 9 and 10 are standard. Footnotes 2 to 8 (left aligned) might be used as needed. They are to be specified in the Shell.

first footnote	is a separating horizontal line.
second – eighth	are free text which can be used for explanations. Footnotes will be referenced using numbers in square brackets, starting with [1], followed by [2] etc.
ninth footnote	left blank; in case needed may also be used as for explanations.
tenth footnote	the program name (left aligned); the date and time in the format ddMMMyyyy hh:mm when the output was created; the version (e.g. draft or final); and the word “Confidential”.

Footnotes are denoted by [1], [2], and so on.

If footnotes take more than 30% of the space of a long listing, they may be presented only on a standalone first page.

TLF numbers and titles should be inputted from an external file that can be directly copy-and-pasted from the SAP planned TLFs, rather than including in the body of the program. This is to ensure consistency between the SAP and the actual outputs.

Footnotes may be inputted from an external file as well for ease of managing changes.

For summary tables, the corresponding listings with the parameters being summarized should be footnoted as reference. For figures, the corresponding summary table should be footnoted as reference.

6 Table, Listing, Figure Metadata

The table, listing, and figure (TLF) metadata will include the TLF numbers, titles, analysis populations, program names, input dataset names. For tables and figures, PARAMCD, PARAM, and other conditions will be specified. TLF numbers, titles, and footnotes will be imported from this master spreadsheet. In addition, this spreadsheet will record the names of the original programmer and the validator/reviewer and the date of validation approval.

7 Significant Digits of Summary Statistics

- All percentages will be rounded to one decimal place and aligned by the decimal place.
- If the count is zero, the percentage will be suppressed and only ‘0’ will be presented.
- Any p-values will be rounded to four decimal places and will be presented as ‘<.0001’ if they are less than 0.0001 after rounding.
- For variables of direct measurements, summary statistics are displayed with the following specifications of decimal places in Table B.

Table B. Significant Digits of Summary Statistics

Description	Characteristic	Number of decimal places
Count	N	0
Mean	Mean	As in source + 1
Standard deviation	Std	As in source + 1
Standard error of the mean	SEM	As in source + 2
Confidence Interval	CI	As in source + 1
Minimum	Min	As in source
Median	Median	As in source
Maximum	Max	As in source
Q1 / Q3	Q1/Q3	As in source
10% / 90%	10%/90%	As in source
Percentage	%	All percentages will be rounded to one decimal place and lined up by the decimal place. The percentage will be suppressed when the count is zero
Coefficient of variation	CV (%)	1
p-value	p-value	p-values will be rounded to four decimal places and will be presented as '<.0001' if they are less than 0.0001 after rounding

N=number; Std=Standard deviation; CI=Confidence Interval; Min=minimum; Max=maximum; CV=Coefficient of variation

As a general guideline for derived parameters, 3 significant digits may be displayed for a parameter with an overall mean less than 100; otherwise, 1 decimal place may be used. If a derived parameter is in the same scale as some related measured parameters, such as MAP, QTc, the same display format may be used as the measured parameters.

Summary Statistics are to be displayed in the following order: Count, Mean, Standard Deviation, <Coefficient of Variation, Standard Error of the Mean, Confidence Interval>, Minimum, <10%>, <Q1>, Median, <Q3>, <90%>, Maximum.

For categorical variables the categories will be displayed in the TLFs in the same order they appear in the CRF.

8 Figure Specifications

- In general, figures should include annotation of key summary statistics: n, mean, SD or SE, median for continuous variables; n and percent for categorical variables; number of subjects at risk and cumulative number of events as well as median and 95% CI for time-to-event data. Other statistics such as quartiles, ranges may be included depending on need and space.
- P-values should be presented if comparisons are of interest.
- For scatter plots, linear or non-linear trend lines should be included if the association of the two variables is of interest. Correlation coefficient or regression coefficients as well as corresponding p-values should be presented.
- For line-charts, if space allows, 1-sided or 2-sided standard error bars should be presented.
- For box plots, 'BOXSTYLE=SCHEMATIC' should be used. The whiskers are drawn to the most extreme points in the group that lie within the fences. The upper fence is defined as the third quartile (represented by the upper edge of the box) plus 1.5 times the interquartile range. The lower fence is defined as the first quartile (represented by the lower edge of the box) minus 1.5 times the interquartile range. Observations outside the fences are identified with a special symbol.

9 Unit Conversion

Units Presented in TLFs	Units Reported or Derived from CRF	Conversion Formula
Kilogram (kg)	Pound (lb)	$\text{kg} = \text{lb}/2.2$
Centimeter (cm)	Inch (in)	$\text{cm} = 2.54 * \text{in}$
Celsius C°	Fahrenheit (F°)	$\text{C}^\circ = (5/9) * (\text{F}^\circ - 32)$
Year	Day	1 year = 365.25 days
Months	Day	1 month = 30.4375 days

APPENDIX III RECIST (1.1) CRITERIA

Below is an outline of the definition of RECIST response categories, extracted from Eisenhauer EA et al. (2009).

Assessment Category	Response Category	Response Criteria
Target lesions	Complete Response (CR)	Disappearance of all target lesions. Any pathological lymph nodes (whether target or non-target) must have reduction in short axis to <10 mm.
	Partial Response (PR)	At least a 30% decrease in the sum of diameters of target lesions, taking as reference the baseline sum diameters.
	Progressive Disease (PD)	At least a 20% increase in the sum of diameters of target lesions, taking as reference the smallest sum on study (this includes the baseline sum if that is the smallest on study). In addition to the relative increase of 20%, the sum must also demonstrate an absolute increase of at least 5 mm. (Note: the appearance of one or more new lesions is also considered progression).
	Stable Disease (SD)	Neither sufficient shrinkage to qualify for PR nor sufficient increase to qualify for PD, taking as reference the smallest sum diameters while on study.
Non-target Lesions	Complete Response (CR)	Disappearance of all non-target lesions and normalisation of tumour marker level. All lymph nodes must be non-pathological in size (<10mm short axis).
	Non-CR/Non-PD:	Persistence of one or more non-target lesion(s) and/or maintenance of tumour marker level above the normal limits.
	Progressive Disease (PD)	Unequivocal progression of existing non-target lesions. (Note: the appearance of one or more new lesions is also considered progression).

Target Lesions	Non-target Lesions	New Lesions	Overall Response
CR	CR	No	CR
CR	Non-CR/no-PD	No	PR
CR	Not evaluated	No	PR
PR	Non-PD or not all evaluated	No	PR
SD	Non-PD or not all evaluated	No	SD
Not all evaluated	Non-PD	No	inevaluable
PD	Any	Yes or No	PD
Any	PD	Yes or No	PD
Any	Any	Yes	PD

APPENDIX IV TNM AND AJCC STAGING DEFINITIONS

The definitions below are extracted from AJCC Cancer Staging Manual, 7th Edition (2010).

TNM definitions

Primary tumor (T)

TX: Primary tumor cannot be assessed

T0: No evidence of primary tumor

Tis: Carcinoma in situ

T1: Tumor limited to the pancreas, ≤ 2 cm in greatest dimension

T2: Tumor limited to the pancreas, > 2 cm in greatest dimension

T3: Tumor extends beyond the pancreas but without involvement of the celiac axis or the superior mesenteric artery

T4: Tumor involves the celiac axis or the superior mesenteric artery (unresectable primary tumor)

Regional lymph nodes (N)

NX: Regional lymph nodes cannot be assessed

N0: No regional lymph node metastasis

N1: Regional lymph node metastasis

Distant metastasis (M)

MX: Distant metastasis cannot be assessed

M0: No distant metastasis

M1: Distant metastasis

AJCC Pancreatic Cancer Staging

<i>Stage</i>	<i>TNM Category</i>
<i>0</i>	Tis, N0, M0
<i>IA</i>	T1, N0, M0
<i>IB</i>	T2, N0, M0
<i>IIA</i>	T3, N0, M0
<i>IIB</i>	T1, N1, M0 T2, N1, M0 T3, N1, M0
<i>III</i>	T4, any N, M0
<i>IV</i>	Any T, any N, M1

APPENDIX V NEW ENGLAND JOURNAL OF MEDICINE REFERENCE RANGES

The following chemistry and hematology reference ranges are from the New England Journal of Medicine 2004; 351: 1548-63

Chemistry

Lab Test	Units	Lower ^[1]	Upper ^[1]
Alanine Aminotransferase (ALT)	U/L	0	35
Alkaline Phosphatase (ALP)	U/L	30	120
Aspartate Aminotransferase (AST)	U/L	0	35
Bicarbonate	mmol/L	21	28
Total Bilirubin	μmol/L	5.1	17
Blood Urea Nitrogen	mmol/L	3.6	7.1
Calcium	mmol/L	2.2	2.6
Creatinine	μmol/L	0	133
Glucose	mmol/L	4.2	6.4
Phosphate	mmol/L	1	1.4
Potassium	mmol/L	3.5	5
Sodium	mmol/L	136	145

¹ Reference ranges from New England Journal of Medicine 2004; 351: 1548-63

Hematology

Lab Test	Units	Lower ^[1]	Upper ^[1]
Erythrocyte MCV	fL	80	100
Erythrocytes (Male)	10 ¹² /L	4.5	5.9
Erythrocytes (Female)	10 ¹² /L	4.0	5.2
Hematocrit (Male)	proportion of 1	0.41	0.53
Hematocrit (Female)	proportion of 1	0.36	0.46
Hemoglobin (Male)	g/dL	13.5	17.5
Hemoglobin (Female)	g/dL	12	16
Leukocytes	10 ⁹ /L	4.5	11.0
Neutrophils	proportion of 1	0.4	0.7
Neutrophils	10 ⁹ /L	1.8	7.7
Lymphocytes	proportion of 1	0.22	0.44
Lymphocytes	10 ⁹ /L	0.99	4.84
Eosinophils	proportion of 1	0	0.08
Eosinophils	10 ⁹ /L	0	0.88
Platelets	10 ⁹ /L	150	350
ANC	10 ⁹ /L	1.8	7.8

^[1]Reference ranges from New England Journal of Medicine 2004; 351: 1548-63

APPENDIX VI CTCAE TOXICITY GRADING FOR LABORATORY TESTS

The following table is extracted from NCI Common Terminology Criteria for Adverse Events (CTCAE) Version 4.03 (June 14, 2010)

Chemistry

		Grade 1	Grade 2	Grade 3	Grade 4
Albumin	Decreased	3 g/dL – LLN	2 - <3 g/dL	<2 g/dL	
Alkaline phosphatase (ALP)		ULN – 2.5 x ULN	>2.5 – 5.0 x ULN	>5.0 – 20.0 x ULN	> 20.0 x ULN
ALT		ULN – 3.0 x ULN	>3.0 – 5.0 x ULN	>5.0 – 20.0 x ULN	>20.0 x ULN
AST		ULN – 3.0 x ULN	>3.0 – 5.0 x ULN	>5.0 – 20.0 x ULN	>20.0 x ULN
Total bilirubin		ULN – 1.5 x ULN	>1.5 – 3.0 x ULN	>3.0 – 10.0 x ULN	>10.0 x ULN
Calcium (Corrected)	Decreased	8.0 mg/dL – LLN	7.0 - <8.0 mg/dL	6.0 - <7.0 mg/dL	<6.0 mg/dL
		ULN – 11.5 mg/dL	>11.5 – 12.5 mg/dL	>12.5 – 13.5 mg/dL	>13.5 mg/dL
GGT		ULN – 2.5 x ULN	>2.5 – 5.0 x ULN	>5.0 – 20.0 x ULN	> 20.0 x ULN
Glucose (Random)	Decreased	55 mg/dL – LLN	40 - <55 mg/dL	30 - <40 mg/dL	<30 mg/dL
Glucose (Fasting)		ULN – 160 mg/dL	>160 – 250 mg/dL	>250 – 500 mg/dL	> 500 mg/dL
Phosphorous	Decreased	2.5 -<LLN mg/dL	2.0 -<2.5 mg/dL	1.0 - <2.0 mg/dL	<1.0 mg/dL
Potassium	Decreased	3.0 mmol/L – LLN	3.0 mmol/L – LLN ^[1]	2.5 - <3.0 mmol/L	<2.5 mmol/L
		ULN – 5.5 mmol/L	>5.5 – 6.0 mmol/L	>6.0 – 7.0 mmol/L	>7.0 mmol/L
Sodium	Decreased	130 mmol/L – LLN	None	120 - <130 mmol/L	<120 mmol/L
		ULN – 150 mmol/L	>150 – 155 mmol/L	>155 – 160 mmol/L	>160 mmol/L
Magnesium	Decreased	1.2 mg/dL – LLN	0.9 - <1.2 mg/dL	0.7 - <0.9 mg/dL	<0.7 mg/dL
		ULN – 3.0 mg/dL	None	>3.0 – 8.0 mg/dL	>8.0 mg/dL

Serum Hematology

		Grade 1	Grade 2	Grade 3	Grade 4
Uric acid		ULN – 10 mg/dL ^[2]	None	ULN – 10 mg/dL ^[3]	>10 mg/dL
Creatinine Enzymatic		>1 – 1.5 x baseline ^[4] ULN – 1.5 x ULN	>1.5 – 3.0 x baseline ^[4] >1.5 – 3.0 x ULN	>3.0 x baseline ^[4] >3.0 – 6.0 x ULN	>6.0 x ULN
Triglycerides		150 – 300 mg/dL	>300 – 500 mg/dL	>500 – 1,000 mg/dL	>1,000 mg/dL
Hgb	Decreased	10.0 g/dL – LLN	8.0 - <10.0 g/dL	<8.0 g/dL	
		>0 – 2 g/dL (+ ULN/Baseline) ^[5]	>2 – 4 g/dL (+ ULN/Baseline) ^[5]	>4 g/dL (+ ULN/Baseline) ^[5]	
Platelet	Decreased	75,000 /mm ³ – LLN	50,000 – <75,000 /mm ³	25,000 - <50,000 /mm ³	<25,000 /mm ³
WBC	Decreased	3,000 /mm ³ – LLN	2,000 - <3,000 /mm ³	1,000 - <2,000 /mm ³	<1,000 /mm ³
		None	None	>100,000 /mm ³	
aPTT		ULN – 1.5 x ULN	>1.5 – 2.5 x ULN	>2.5 x ULN	
Lymphocytes	Decreased	800 /mm ³ – LLN	500 - <800 /mm ³	200 - <500 /mm ³	<200 /mm ³
		None	>4,000 – 20,000 /mm ³	>20,000 /mm ³	
Neutrophils	Decreased	1,500 /mm ³ – LLN	1,000 - <1,500 /mm ³	500 - <1,000 /mm ³	<500 /mm ³

Decreased: below LLN; Otherwise, above ULN;

[1] Symptomatic, Intervention indicated

[2] without physiologic consequences

[3] with physiologic consequences

[4] Baseline is used if it is above ULN

[5] Increase from ULN/baseline: if baseline is above ULN, the increase should be above the baseline; otherwise, the increase should be above ULN.

APPENDIX VII NCCN VERSION 2. 2014 PANCREATIC CANCER RESECTABILITY CRITERIA

The following is extracted from ‘NCCN Guidelines Version 2.2014, Pancreatic Adenocarcinoma’, page 19.

Tumors considered to be unresectable demonstrate the following:

❖ HEAD

- Distant metastases
- Greater than 180 degrees SMA encasement, any celiac abutment
- Unreconstructible SMV/portal occlusion
- Aortic or inferior vena cava (IVC) invasion or encasement

❖ BODY

- Distant metastases
- SMA or celiac encasement greater than 180 degrees
- Unreconstructible SMV/portal occlusion
- Aortic invasion

❖ TAIL

- Distant metastases
- SMA or celiac encasement greater than 180 degrees

❖ Nodal status

- Metastases to lymph nodes beyond the field of resection should be considered unresectable

APPENDIX VIII TERMINOLOGY USED IN THIS STUDY

Tumor Grade

The grade of the cancer (how abnormal the cells look under the microscope) uses a scale from G1 to G3 (or sometimes G1 to G4), with G1 cancers looking the most like normal cells and having the best outlook.

The details of grading are a little different for pancreatic neuroendocrine tumors (NETs), where measures of how many of the cells are in the process of dividing is an important part of grading. This is determined by counting mitoses (cells that have started to split into two new cells) under a microscope and with a Ki-67 test that recognizes cells that are almost ready to start splitting. Based on these tests, NETs are divided into 2 groups:

- **Well-differentiated NETs** (which includes low-grade [G1] and intermediate-grade [G2] tumors) have 20 or fewer mitoses and a Ki-67 index of 20% or lower.
- **Poorly differentiated tumors** (high-grade [G3] tumors) have more than 20 mitoses or a Ki-67 index of more than 20%. These are also called *neuroendocrine carcinomas*, and they often grow and spread quickly.

Extent of resection

For patients who have surgery, another important factor is the *extent of the resection* — whether or not all of the tumor is removed:

- **R0:** All of the cancer is thought to have been removed. (There are no visible or microscopic signs suggesting that cancer was left behind.)
- **R1:** All visible tumor was removed, but lab tests of the removed specimen show that some small areas of cancer were probably left behind.
- **R2:** Some visible tumor could not be removed.

Resectable versus unresectable pancreatic cancer

The AJCC staging system gives a detailed summary of how far the cancer has spread. But for treatment purposes, doctors use a simpler staging system, which divides cancers into groups based on whether or not they can be removed (resected) with surgery:

Resectable

If the cancer is only in the pancreas (or has spread just beyond it) and the surgeon believes the entire tumor can be removed, it is called *resectable*. (In general, this would include most stage IA, IB, and IIA cancers in the TNM system.)

It's important to note that some cancers might appear to be resectable based on imaging tests such as CT scans, but once surgery is started it might become clear that not all of the cancer can be removed. If

this happens, only a sample of the cancer may be removed to confirm the diagnosis (if a biopsy hasn't been done already), and the rest of the planned operation will be stopped to help avoid the risk of major side effects.

Borderline resectable

This term is used to describe some cancers that might have just reached nearby blood vessels, but which the doctors feel might still be removed completely with surgery. This would include some stage III cancers in the TNM system.

Unresectable

These cancers can't be removed entirely by surgery.

Locally advanced: If the cancer has not yet spread to distant organs but it still can't be removed completely with surgery, it is called *locally advanced*. Often the reason the cancer can't be removed is because it has grown into or surrounded nearby major blood vessels. (In general, this would include stage IIB and most III cancers in the TNM system.)

Surgery to try to remove these tumors would be very unlikely to be helpful and could still have major side effects. Some type of surgery might still be done, but it would be a less extensive operation with the goal of preventing or relieving symptoms or problems like a blocked bile duct or intestinal tract, instead of trying to cure the cancer.

Metastatic: If the cancer has spread to distant organs, it is called *metastatic*. These cancers can't be removed completely. Surgery might still be done, but the goal would be to prevent or relieve symptoms, not to try to cure the cancer.

APPENDIX IX STANDARDIZED MEDDRA QUERIES (SMQ) FOR ANAPHYLACTIC REACTION AND HYPERSENSITIVITY

SMQ for Anaphylactic Reaction

Name	Code	Level
Anaphylactic reaction	10002198	PT
Anaphylactic shock	10002199	PT
Anaphylactic transfusion reaction	10067113	PT
Anaphylactoid reaction	10002216	PT
Anaphylactoid shock	10063119	PT
Circulatory collapse	10009192	PT
Dialysis membrane reaction	10076665	PT
Kounis syndrome	10069167	PT
Shock	10040560	PT
Shock symptom	10040581	PT
Type I hypersensitivity	10045240	PT
Acute respiratory failure	10001053	PT
Asthma	10003553	PT
Bronchial oedema	10056695	PT
Bronchospasm	10006482	PT
Cardio-respiratory distress	10049874	PT
Chest discomfort	10008469	PT
Choking	10008589	PT
Choking sensation	10008590	PT
Circumoral oedema	10052250	PT
Cough	10011224	PT
Cyanosis	10011703	PT
Dyspnoea	10013968	PT
Hyperventilation	10020910	PT
Irregular breathing	10076213	PT
Laryngeal dyspnoea	10052390	PT
Laryngeal oedema	10023845	PT
Laryngospasm	10023891	PT
Laryngotracheal oedema	10023893	PT
Mouth swelling	10075203	PT
Nasal obstruction	10028748	PT
Oedema mouth	10030110	PT

Oropharyngeal oedema	10078783	PT
Oropharyngeal spasm	10031111	PT
Oropharyngeal swelling	10031118	PT
Pharyngeal oedema	10034829	PT
Respiratory arrest	10038669	PT
Respiratory distress	10038687	PT
Respiratory failure	10038695	PT
Reversible airways obstruction	10062109	PT
Sensation of foreign body	10061549	PT
Sneezing	10041232	PT
Stridor	10042241	PT
Swollen tongue	10042727	PT
Tachypnoea	10043089	PT
Throat tightness	10043528	PT
Tongue oedema	10043967	PT
Tracheal obstruction	10044291	PT
Tracheal oedema	10044296	PT
Upper airway obstruction	10067775	PT
Wheezing	10047924	PT
Allergic oedema	10060934	PT
Angioedema	10002424	PT
Erythema	10015150	PT
Eye oedema	10052139	PT
Eye pruritus	10052140	PT
Eye swelling	10015967	PT
Eyelid oedema	10015993	PT
Face oedema	10016029	PT
Flushing	10016825	PT
Generalised erythema	10051576	PT
Injection site urticaria	10022107	PT
Lip oedema	10024558	PT
Lip swelling	10024570	PT
Nodular rash	10075807	PT
Ocular hyperaemia	10030041	PT
Oedema	10030095	PT

Periorbital oedema	10034545	PT
Pruritus	10037087	PT
Pruritus allergic	10063438	PT
Pruritus generalised	10052576	PT
Rash	10037844	PT
Rash erythematous	10037855	PT
Rash generalised	10037858	PT
Rash pruritic	10037884	PT
Skin swelling	10053262	PT
Swelling	10042674	PT
Swelling face	10042682	PT
Urticaria	10046735	PT
Urticaria papular	10046750	PT
Blood pressure decreased	10005734	PT
Blood pressure diastolic decreased	10005737	PT
Blood pressure systolic decreased	10005758	PT
Cardiac arrest	10007515	PT
Cardio-respiratory arrest	10007617	PT
Cardiovascular insufficiency	10065929	PT
Diastolic hypotension	10066077	PT
Hypotension	10021097	PT

SMQ for Hypersensitivity

Name	Code	Level
Acute generalised exanthematous pustulosis	10048799	PT
Administration site dermatitis	10075096	PT
Administration site eczema	10075099	PT
Administration site hypersensitivity	10075102	PT
Administration site rash	10071156	PT
Administration site recall reaction	10075964	PT
Administration site urticaria	10075109	PT
Administration site vasculitis	10075969	PT
Allergic bronchitis	10052613	PT
Allergic colitis	10059447	PT
Allergic cough	10053779	PT

Allergic cystitis	10051394	PT
Allergic eosinophilia	10075185	PT
Allergic gastroenteritis	10075308	PT
Allergic hepatitis	10071198	PT
Allergic keratitis	10057380	PT
Allergic myocarditis	10001715	PT
Allergic oedema	10060934	PT
Allergic otitis externa	10075072	PT
Allergic otitis media	10061557	PT
Allergic pharyngitis	10050639	PT
Allergic reaction to excipient	10078853	PT
Allergic respiratory disease	10063532	PT
Allergic respiratory symptom	10063527	PT
Allergic sinusitis	10049153	PT
Allergic transfusion reaction	10066173	PT
Allergy alert test positive	10075479	PT
Allergy test positive	10056352	PT
Allergy to immunoglobulin therapy	10074079	PT
Allergy to surgical sutures	10077279	PT
Allergy to vaccine	10055048	PT
Alveolitis allergic	10001890	PT
Anaphylactic reaction	10002198	PT
Anaphylactic shock	10002199	PT
Anaphylactic transfusion reaction	10067113	PT
Anaphylactoid reaction	10002216	PT
Anaphylactoid shock	10063119	PT
Anaphylaxis treatment	10002222	PT
Angioedema	10002424	PT
Antiallergic therapy	10064059	PT
Antiendomysial antibody positive	10065514	PT
Anti-neutrophil cytoplasmic antibody positive vasculitis	10050894	PT
Application site dermatitis	10003036	PT
Application site eczema	10050099	PT
Application site hypersensitivity	10063683	PT
Application site rash	10003054	PT

Application site recall reaction	10076024	PT
Application site urticaria	10050104	PT
Application site vasculitis	10076027	PT
Arthritis allergic	10061430	PT
Aspirin-exacerbated respiratory disease	10075084	PT
Atopy	10003645	PT
Blepharitis allergic	10005149	PT
Blood immunoglobulin E abnormal	10005589	PT
Blood immunoglobulin E increased	10005591	PT
Bromoderma	10006404	PT
Bronchospasm	10006482	PT
Catheter site dermatitis	10073992	PT
Catheter site eczema	10073995	PT
Catheter site hypersensitivity	10073998	PT
Catheter site rash	10052271	PT
Catheter site urticaria	10052272	PT
Catheter site vasculitis	10074014	PT
Chronic eosinophilic rhinosinusitis	10071399	PT
Chronic hyperplastic eosinophilic sinusitis	10071380	PT
Circulatory collapse	10009192	PT
Circumoral oedema	10052250	PT
Conjunctival oedema	10010726	PT
Conjunctivitis allergic	10010744	PT
Contact stomatitis	10067510	PT
Contrast media allergy	10066973	PT
Contrast media reaction	10010836	PT
Corneal oedema	10011033	PT
Cutaneous vasculitis	10011686	PT
Dennie-Morgan fold	10062918	PT
Dermatitis	10012431	PT
Dermatitis acneiform	10012432	PT
Dermatitis allergic	10012434	PT
Dermatitis atopic	10012438	PT
Dermatitis bullous	10012441	PT
Dermatitis contact	10012442	PT

Dermatitis exfoliative	10012455	PT
Dermatitis exfoliative generalised	10012456	PT
Dermatitis herpetiformis	10012468	PT
Dermatitis infected	10012470	PT
Dermatitis psoriasiform	10058675	PT
Device allergy	10072867	PT
Dialysis membrane reaction	10076665	PT
Distributive shock	10070559	PT
Documented hypersensitivity to administered product	10076470	PT
Drug cross-reactivity	10076743	PT
Drug eruption	10013687	PT
Drug hypersensitivity	10013700	PT
Drug provocation test	10074350	PT
Drug reaction with eosinophilia and systemic symptoms	10073508	PT
Eczema	10014184	PT
Eczema infantile	10014198	PT
Eczema nummular	10014201	PT
Eczema vaccinatum	10066042	PT
Eczema vesicular	10058681	PT
Eczema weeping	10055182	PT
Encephalitis allergic	10056387	PT
Encephalopathy allergic	10014627	PT
Eosinophilic granulomatosis with polyangiitis	10078117	PT
Epidermal necrosis	10059284	PT
Epidermolysis	10053177	PT
Epidermolysis bullosa	10014989	PT
Epiglottic oedema	10015029	PT
Erythema multiforme	10015218	PT
Erythema nodosum	10015226	PT
Exfoliative rash	10064579	PT
Eye allergy	10015907	PT
Eye oedema	10052139	PT
Eye swelling	10015967	PT
Eyelid oedema	10015993	PT
Face oedema	10016029	PT

Fixed eruption	10016741	PT
Giant papillary conjunctivitis	10018258	PT
Gingival oedema	10049305	PT
Gingival swelling	10018291	PT
Gleich's syndrome	10066837	PT
Haemorrhagic urticaria	10059499	PT
Hand dermatitis	10058898	PT
Henoch-Schonlein purpura	10019617	PT
Henoch-Schonlein purpura nephritis	10069440	PT
Heparin-induced thrombocytopenia	10062506	PT
Hereditary angioedema	10019860	PT
Hypersensitivity	10020751	PT
Hypersensitivity vasculitis	10020764	PT
Idiopathic urticaria	10021247	PT
Immediate post-injection reaction	10067142	PT
Immune thrombocytopenic purpura	10074667	PT
Immune tolerance induction	10070581	PT
Immune-mediated adverse reaction	10077665	PT
Implant site dermatitis	10063855	PT
Implant site hypersensitivity	10063858	PT
Implant site rash	10063786	PT
Implant site urticaria	10063787	PT
Incision site dermatitis	10073168	PT
Incision site rash	10073411	PT
Infusion site dermatitis	10065458	PT
Infusion site eczema	10074850	PT
Infusion site hypersensitivity	10065471	PT
Infusion site rash	10059830	PT
Infusion site recall reaction	10076085	PT
Infusion site urticaria	10065490	PT
Infusion site vasculitis	10074851	PT
Injection site dermatitis	10022056	PT
Injection site eczema	10066221	PT
Injection site hypersensitivity	10022071	PT
Injection site rash	10022094	PT

Injection site recall reaction	10066797	PT
Injection site urticaria	10022107	PT
Injection site vasculitis	10067995	PT
Instillation site hypersensitivity	10073612	PT
Instillation site rash	10073622	PT
Instillation site urticaria	10073627	PT
Interstitial granulomatous dermatitis	10067972	PT
Intestinal angioedema	10076229	PT
Iodine allergy	10052098	PT
Kaposi's varicelliform eruption	10051891	PT
Kounis syndrome	10069167	PT
Laryngeal oedema	10023845	PT
Laryngitis allergic	10064866	PT
Laryngospasm	10023891	PT
Laryngotracheal oedema	10023893	PT
Limbal swelling	10070492	PT
Lip oedema	10024558	PT
Lip swelling	10024570	PT
Mast cell degranulation present	10076606	PT
Medical device site dermatitis	10075572	PT
Medical device site eczema	10075575	PT
Medical device site hypersensitivity	10075579	PT
Medical device site rash	10075585	PT
Medical device site recall reaction	10076140	PT
Medical device site urticaria	10075588	PT
Mouth swelling	10075203	PT
Mucocutaneous rash	10056671	PT
Multiple allergies	10028164	PT
Nephritis allergic	10029120	PT
Nikolsky's sign	10029415	PT
Nodular rash	10075807	PT
Oculomucocutaneous syndrome	10030081	PT
Oculorespiratory syndrome	10067317	PT
Oedema mouth	10030110	PT
Oral allergy syndrome	10068355	PT

Oropharyngeal blistering	10067950	PT
Oropharyngeal oedema	10078783	PT
Oropharyngeal spasm	10031111	PT
Oropharyngeal swelling	10031118	PT
Palatal oedema	10056998	PT
Palatal swelling	10074403	PT
Palisaded neutrophilic granulomatous dermatitis	10068809	PT
Palpable purpura	10056872	PT
Pathergy reaction	10074332	PT
Periorbital oedema	10034545	PT
Pharyngeal oedema	10034829	PT
Pruritus allergic	10063438	PT
Radioallergosorbent test positive	10037789	PT
Rash	10037844	PT
Rash erythematous	10037855	PT
Rash follicular	10037857	PT
Rash generalised	10037858	PT
Rash macular	10037867	PT
Rash maculo-papular	10037868	PT
Rash maculovesicular	10050004	PT
Rash morbilliform	10037870	PT
Rash neonatal	10037871	PT
Rash papulosquamous	10037879	PT
Rash pruritic	10037884	PT
Rash pustular	10037888	PT
Rash rubelliform	10057984	PT
Rash scarlatiniform	10037890	PT
Rash vesicular	10037898	PT
Reaction to azo-dyes	10037973	PT
Reaction to colouring	10037974	PT
Reaction to drug excipients	10064787	PT
Reaction to preservatives	10064788	PT
Red man syndrome	10038192	PT
Rhinitis allergic	10039085	PT
Scleral oedema	10057431	PT

Scleritis allergic	10051126	PT
Scrotal oedema	10039755	PT
Serum sickness	10040400	PT
Serum sickness-like reaction	10040402	PT
Shock	10040560	PT
Shock symptom	10040581	PT
Skin necrosis	10040893	PT
Skin reaction	10040914	PT
Skin test positive	10040934	PT
Solar urticaria	10041307	PT
Solvent sensitivity	10041316	PT
Stevens-Johnson syndrome	10042033	PT
Stoma site hypersensitivity	10074509	PT
Stoma site rash	10059071	PT
Swelling face	10042682	PT
Swollen tongue	10042727	PT
Symmetrical drug-related intertriginous and flexural exanthema	10078325	PT
Tongue oedema	10043967	PT
Toxic epidermal necrolysis	10044223	PT
Toxic skin eruption	10057970	PT
Tracheal oedema	10044296	PT
Type I hypersensitivity	10045240	PT
Type II hypersensitivity	10054000	PT
Type III immune complex mediated reaction	10053614	PT
Type IV hypersensitivity reaction	10053613	PT
Urticaria	10046735	PT
Urticaria cholinergic	10046740	PT
Urticaria chronic	10052568	PT
Urticaria contact	10046742	PT
Urticaria papular	10046750	PT
Urticaria physical	10046751	PT
Urticaria pigmentosa	10046752	PT
Urticaria vesiculosa	10046755	PT
Urticarial vasculitis	10048820	PT
Vaccination site dermatitis	10069477	PT

Vaccination site eczema	10076161	PT
Vaccination site exfoliation	10069489	PT
Vaccination site hypersensitivity	10068880	PT
Vaccination site rash	10069482	PT
Vaccination site recall reaction	10076188	PT
Vaccination site urticaria	10069622	PT
Vaccination site vasculitis	10076191	PT
Vaccination site vesicles	10069623	PT
Vaginal exfoliation	10064483	PT
Vaginal ulceration	10046943	PT
Vasculitic rash	10047111	PT
Vessel puncture site rash	10077117	PT
Vessel puncture site vesicles	10077813	PT
Vulval ulceration	10047768	PT
Vulvovaginal rash	10071588	PT
Vulvovaginal ulceration	10050181	PT
Acute respiratory failure	10001053	PT
Administration site photosensitivity reaction	10075961	PT
Airway remodelling	10075289	PT
Allergy to chemicals	10061626	PT
Allergy to fermented products	10054929	PT
Alpha tumour necrosis factor increased	10059982	PT
Alveolitis	10001889	PT
Antibody test abnormal	10061425	PT
Antibody test positive	10061427	PT
Anti-insulin antibody increased	10053815	PT
Anti-insulin antibody positive	10053814	PT
Anti-insulin receptor antibody increased	10068226	PT
Anti-insulin receptor antibody positive	10068225	PT
Application site photosensitivity reaction	10058730	PT
Asthma	10003553	PT
Asthma late onset	10003559	PT
Asthma-chronic obstructive pulmonary disease overlap syndrome	10077005	PT
Asthmatic crisis	10064823	PT
Auricular swelling	10003800	PT

Blister	10005191	PT
Blister rupture	10073385	PT
Blood immunoglobulin A abnormal	10005584	PT
Blood immunoglobulin A increased	10005586	PT
Blood immunoglobulin D increased	10063244	PT
Blood immunoglobulin G abnormal	10005594	PT
Blood immunoglobulin G increased	10005596	PT
Blood immunoglobulin M abnormal	10005599	PT
Blood immunoglobulin M increased	10005601	PT
Bronchial hyperreactivity	10066091	PT
Bronchial oedema	10056695	PT
Bullous impetigo	10006563	PT
Caffeine allergy	10074895	PT
Capillaritis	10068406	PT
Charcot-Leyden crystals	10008413	PT
Choking	10008589	PT
Choking sensation	10008590	PT
Conjunctivitis	10010741	PT
Corneal exfoliation	10064489	PT
Cytokine release syndrome	10052015	PT
Cytokine storm	10050685	PT
Ear swelling	10014025	PT
Eosinophil count abnormal	10061125	PT
Eosinophil count increased	10014945	PT
Eosinophil percentage abnormal	10058133	PT
Eosinophil percentage increased	10052222	PT
Eosinophilia	10014950	PT
Eosinophilia myalgia syndrome	10014952	PT
Eosinophilic bronchitis	10065563	PT
Eosinophilic oesophagitis	10064212	PT
Eosinophilic pneumonia	10014962	PT
Eosinophilic pneumonia acute	10052832	PT
Eosinophilic pneumonia chronic	10052833	PT
Erythema	10015150	PT
Flushing	10016825	PT

Gastrointestinal oedema	10058061	PT
Generalised erythema	10051576	PT
Generalised oedema	10018092	PT
Genital rash	10018175	PT
Genital swelling	10067639	PT
Haemolytic transfusion reaction	10067122	PT
HLA marker study positive	10067937	PT
Immune complex level increased	10064650	PT
Immunoglobulins abnormal	10021497	PT
Immunoglobulins increased	10021500	PT
Immunology test abnormal	10061214	PT
Implant site photosensitivity	10073415	PT
Infantile asthma	10049585	PT
Infusion site photosensitivity reaction	10065486	PT
Injection site photosensitivity reaction	10053396	PT
Interstitial lung disease	10022611	PT
Laryngeal dyspnoea	10052390	PT
Laryngeal obstruction	10059639	PT
Leukotriene increased	10064663	PT
Lip exfoliation	10064482	PT
Localised oedema	10048961	PT
Mechanical urticaria	10068773	PT
Medical device site photosensitivity reaction	10076137	PT
Mesenteric panniculitis	10063031	PT
Mouth ulceration	10028034	PT
Mucocutaneous ulceration	10028084	PT
Mucosa vesicle	10028103	PT
Mucosal erosion	10061297	PT
Mucosal exfoliation	10064486	PT
Mucosal necrosis	10067993	PT
Mucosal ulceration	10028124	PT
Nasal crease	10078581	PT
Necrotising panniculitis	10062579	PT
Neurodermatitis	10029263	PT
Neutralising antibodies positive	10064980	PT

Noninfective conjunctivitis	10074701	PT
Non-neutralising antibodies positive	10064982	PT
Occupational asthma	10070836	PT
Occupational dermatitis	10030012	PT
Oedema mucosal	10030111	PT
Oral mucosal exfoliation	10064487	PT
Orbital oedema	10031051	PT
Panniculitis	10033675	PT
Penile exfoliation	10064485	PT
Penile oedema	10066774	PT
Penile swelling	10034319	PT
Perineal rash	10075364	PT
Perivascular dermatitis	10064986	PT
Photosensitivity reaction	10034972	PT
Pneumonitis	10035742	PT
Prurigo	10037083	PT
Pruritus	10037087	PT
Pruritus generalised	10052576	PT
Pulmonary eosinophilia	10037382	PT
Reactive airways dysfunction syndrome	10070832	PT
Respiratory arrest	10038669	PT
Respiratory distress	10038687	PT
Respiratory failure	10038695	PT
Respiratory tract oedema	10070774	PT
Reversible airways obstruction	10062109	PT
Rhinitis perennial	10039094	PT
Scrotal swelling	10039759	PT
Seasonal allergy	10048908	PT
Septal panniculitis	10056876	PT
Skin erosion	10040840	PT
Skin exfoliation	10040844	PT
Skin oedema	10058679	PT
Skin swelling	10053262	PT
Sneezing	10041232	PT
Status asthmaticus	10041961	PT

Stomatitis	10042128	PT
Streptokinase antibody increased	10053797	PT
Stridor	10042241	PT
Suffocation feeling	10042444	PT
Throat tightness	10043528	PT
Tongue exfoliation	10064488	PT
Tracheal obstruction	10044291	PT
Tracheostomy	10044320	PT
Transplantation associated food allergy	10075008	PT
Upper airway obstruction	10067775	PT
Vaccination site photosensitivity reaction	10076186	PT
Vaginal oedema	10063818	PT
Visceral oedema	10065768	PT
Vulval oedema	10047763	PT
Vulvovaginal swelling	10071211	PT
Wheezing	10047924	PT