

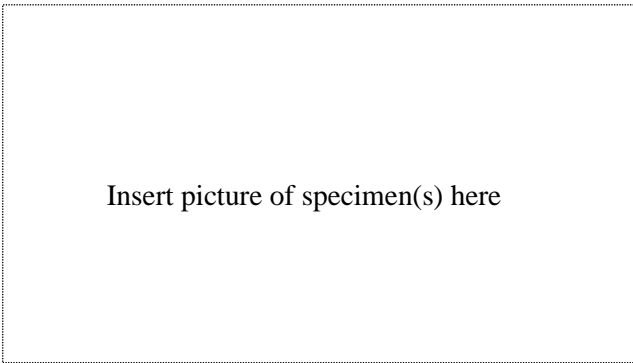
Clinical tissue processing

Date: _____

Processing by: _____

Codename: _____

Time collected: ____ am pm



Tissue descriptions

Tissue type	Weight	Sample description (<i>appearance, abnormalities, size, colour, necrosis</i>)
_____ TURBT: <input type="checkbox"/> Cystectomy: <input type="checkbox"/>		

Gravity wash in 37°C basal media to remove dead cells. Tare empty 50 mL falcon and weigh (g or mg)

Other specimen descriptions

Specimen	# samples	Total volume	# Stored -80°C	Comments on sample
Blood (EDTA)			See PBMC isolation protocol	1mL Plasma taken <input type="checkbox"/>
Serum (SST)				
Urine (whole)	Whole urine	<input type="checkbox"/>		
	Urine sediment pellet	<input type="checkbox"/>		
	PreservCyte (cytology)	<input type="checkbox"/>	Volume: _____ mL urine: _____ : Volume: _____ mL PreservCyte (1: 3 ratio)	
	Urine cytospin	<input type="checkbox"/>		
	Urine agarose pellet	<input type="checkbox"/>		

As per standard specimen processing protocols. Any modifications please outline in general comments

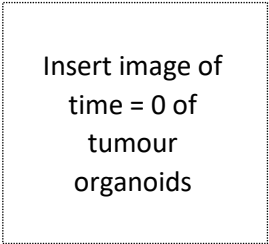
PBMC collection

Sample	Viability	Cell number	# of cryovials @ 1x10 ⁷	Comments
PBMC				
TILs				

Sample storage location: _____

Time: ____ am pm

Cell clusters visible following extraction? Yes No



General comments:

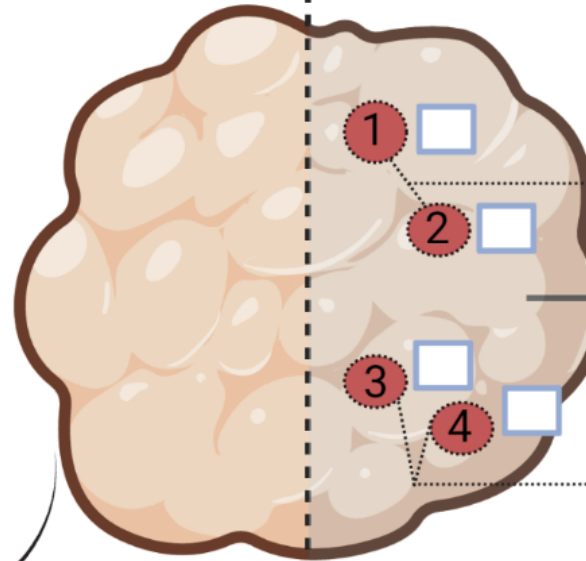
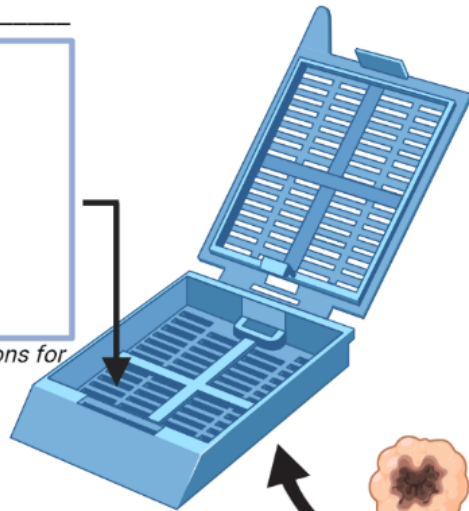
Specimen _____

Specimen _____

Histology



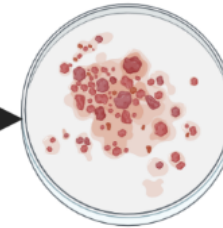
draw layout of sections for histology



fresh frozen tissue



Organoid extraction



Weight/ 10 mL:

Comments:

Weight _____ (g/ mg)