

Current instrument: **Histopathological details**[Preview instrument](#)

Add Field		Add Matrix of Fields		Import from Field Bank	
				Variable: tnm_t	Branching logic: [histo_diagn] = '1' or [histo_diagn] = '2' or [histo_diagn] = '...
T	<input type="radio"/> In situ <input type="radio"/> 1 <input type="radio"/> 2 <input type="radio"/> 3 <input type="radio"/> 4 <input type="radio"/> unknown				reset
TNM classification (from histopathology if available)					
Add Field		Add Matrix of Fields		Import from Field Bank	
				Variable: tnm_n	Branching logic: [histo_diagn] = '1' or [histo_diagn] = '2' or [histo_diagn] = '...
N	<input type="radio"/> 0 <input type="radio"/> 1 <input type="radio"/> 2 <input type="radio"/> unknown				reset
TNM classification (from histopathology if available)					
Add Field		Add Matrix of Fields		Import from Field Bank	
				Variable: tnm_m	Branching logic: [histo_diagn] = '1' or [histo_diagn] = '2' or [histo_diagn] = '...
M	<input type="radio"/> 0 <input type="radio"/> 1 <input type="radio"/> 2 <input type="radio"/> unknown				reset
TNM classification (from histopathology if available)					
Add Field		Add Matrix of Fields		Import from Field Bank	
				Variable: tnm_v	Branching logic: [histo_diagn] = '1' or [histo_diagn] = '2' or [histo_diagn] = '...
V	<input type="radio"/> 0 <input type="radio"/> 1 <input type="radio"/> unknown				reset
TNM classification (from histopathology if available)					
Add Field		Add Matrix of Fields		Import from Field Bank	
				Variable: tnm_l	Branching logic: [histo_diagn] = '1' or [histo_diagn] = '2' or [histo_diagn] = '...
L	<input type="radio"/> 0 <input type="radio"/> 1 <input type="radio"/> unknown				reset
TNM classification (from histopathology if available)					
Add Field		Add Matrix of Fields		Import from Field Bank	
				Variable: tnm_pn	Branching logic: [histo_diagn] = '1' or [histo_diagn] = '2' or [histo_diagn] = '...
Pn	<input type="radio"/> 0 <input type="radio"/> 1 <input type="radio"/> unknown				reset
TNM classification (from histopathology if available)					
Add Field		Add Matrix of Fields		Import from Field Bank	
				Variable: tnm_g	Branching logic: [histo_diagn] = '1' or [histo_diagn] = '2' or [histo_diagn] = '...
Grading	<input type="radio"/> 1 <input type="radio"/> 2 <input type="radio"/> 3 <input type="radio"/> unknown				reset
TNM classification (from histopathology if available)					
Add Field		Add Matrix of Fields		Import from Field Bank	
				Variable: tnm_r	
Resection margin	<input type="radio"/> R0 <input type="radio"/> R1 <input type="radio"/> R2 <input type="radio"/> R unknown/ Rx				reset
Add Field		Add Matrix of Fields		Import from Field Bank	
				Variable: histo_genetic	
Genetic analysis performed	<input type="radio"/> Yes <input type="radio"/> No				reset
Add Field		Add Matrix of Fields		Import from Field Bank	
				Variable: histo_gene_mutation	Branching logic: [histo_genetic] = '1'
Genetic testing positive for	<input type="checkbox"/> BRAF <input type="checkbox"/> MSI <input type="checkbox"/> RAS <input type="checkbox"/> other <input type="checkbox"/> none				
Add Field		Add Matrix of Fields		Import from Field Bank	
				Variable: historische_gene_other	Branching logic: [histo_gene_mutation(4)] = '1'
Specify other positive finding	<input type="text"/>				
Add Field		Add Matrix of Fields		Import from Field Bank	