

# Schema documentation for NoSite.xsd

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## Namespace: "http://purl.org/obbr/pathology/cancer/"

### Schemas

#### Main schema NoSite.xsd

Namespace	http://purl.org/obbr/pathology/cancer/
Properties	attribute form default: unqualified element form default: qualified

#### Included schema generic.xsd

Namespace	http://purl.org/obbr/pathology/cancer/
Properties	attribute form default: unqualified element form default: unqualified

### Elements

#### Element size / response

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Properties	content: complex minOccurs: 1 maxOccurs: 3
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'
Children	comment, label, specify
Instance	<pre> &lt;response relation="" unit="" value=""&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/response&gt; </pre>

Attributes	QName	Type	Fixed	Default	Use
	<b>relation</b>	relation.valueset			optional
	<b>unit</b>	lengthUnit.valueset			required
	<b>value</b>	positiveDecimal			required
Source	<pre>&lt;xs:element maxOccurs="3" minOccurs="1" name="response"&gt;   &lt;xs:complexType&gt;     &lt;xs:group ref="generic.content"/&gt;     &lt;xs:attributeGroup ref="dimensionalQuantity"/&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt;</pre>				

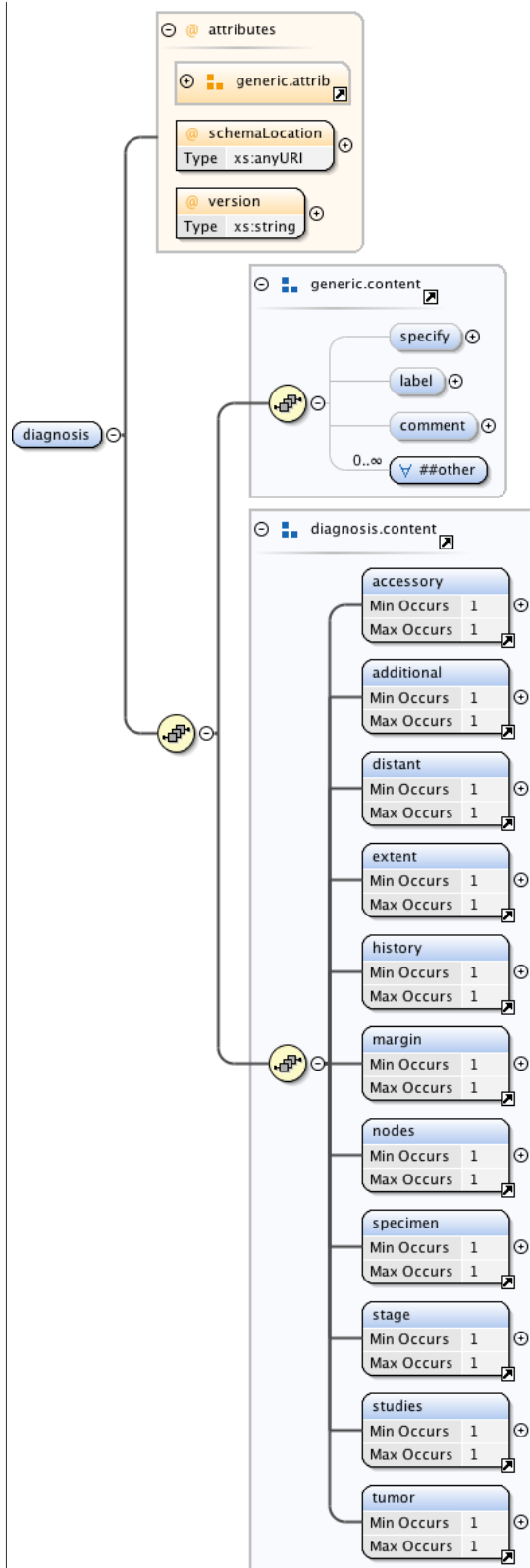
## Element length / response

Namespace	http://purl.org/obbr/pathology/cancer/				
Diagram					
Properties	content:	complex			
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than "http://purl.org/obbr/pathology/cancer/"				
Children	comment, label, specify				
Instance	<pre>&lt;response relation="" unit="" value=""&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/response&gt;</pre>				
Attributes	QName	Type	Fixed	Default	Use
	relation	relation.valueset			optional
	unit	lengthUnit.valueset			required
	value	positiveDecimal			required
Source	<pre>&lt;xs:element name="response"&gt;   &lt;xs:complexType&gt;     &lt;xs:group ref="generic.content"/&gt;     &lt;xs:attributeGroup ref="dimensionalQuantity"/&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt;</pre>				

## Element diagnosis

Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram



Properties	content: complex
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , accessory , additional , distant , extent , history , margin , nodes , specimen , stage , studies , tumor
Children	accessory, additional, comment, distant, extent, history, label, margin, nodes, specify, specimen, stage, studies, tumor
Instance	<pre>&lt;diagnosis id="" schemaLocation="" version=""&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;</pre>

	<code>&lt;label&gt;{0,1}&lt;/label&gt;</code> <code>&lt;comment&gt;{0,1}&lt;/comment&gt;</code> <code>&lt;/diagnosis&gt;</code>				
Attributes	<b>QName</b> <b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b>	<b>Type</b>	<b>Fixed</b>	<b>Default</b>	<b>Use</b>
	<b>id</b>	xs:ID			optional
	<b>schemaLocation</b>	xs:anyURI			required
	<b>version</b>	xs:string			required
Source	<pre> &lt;xs:element name="diagnosis"&gt;   &lt;xs:complexType&gt;     &lt;xs:sequence&gt;       &lt;xs:group ref="generic.content"/&gt;       &lt;xs:group ref="diagnosis.content"/&gt;     &lt;/xs:sequence&gt;     &lt;xs:attributeGroup ref="generic.attrib"/&gt;     &lt;xs:attribute name="schemaLocation" type="xs:anyURI" use="required"/&gt;     &lt;xs:attribute name="version" type="xs:string" use="required"/&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt; </pre>				

## Element accessory

Namespace	http://purl.org/obbr/pathology/cancer/		
Diagram	<pre>graph TD     accessory((accessory)) --- g1(( ))     g1 --- generic_attrib[generic.attrib]     g1 --- g2(( ))     g2 --- generic_content[generic.content]     g2 --- g3(( ))     g3 --- accessory_content[accessory.content]     accessory_content --- g4(( ))     g4 --- treatment_response[treatmentResponse]     g4 --- g5(( ))     g5 --- lymphvascular_invasion[lymphvascularInvasion]     g5 --- g6(( ))     g6 --- perineural_invasion[perineuralInvasion]</pre>		
Properties	content:	complex	
Used by	Element Group	diagnosis.content	
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , treatmentResponse , lymphvascularInvasion , perineuralInvasion		



Children	comment, label, lymphvascularInvasion, perineuralInvasion, specify, treatmentResponse				
Instance	<pre>&lt;accessory id=""&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/accessory&gt;</pre>				
Attributes	<b>QName</b>	<b>Type</b>	<b>Fixed</b>	<b>Default</b>	<b>Use</b>
	ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'				
	id	xs:ID			optional
Source	<pre>&lt;xs:element name="accessory"&gt;   &lt;xs:complexType&gt;     &lt;xs:sequence&gt;       &lt;xs:group ref="generic.content"/&gt;       &lt;xs:group ref="accessory.content"/&gt;     &lt;/xs:sequence&gt;     &lt;xs:attributeGroup ref="generic.attrib"/&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt;</pre>				

### Element accessory.treatment / treatmentResponse

Namespace	http://purl.org/obbr/pathology/cancer/				
Diagram					
Properties	content:	complex			
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , response				
Children	comment, label, response, specify				
Instance	<pre>&lt;treatmentResponse id=""&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/treatmentResponse&gt;</pre>				
Attributes	QName	Type	Fixed	Default	Use
	ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'				
	id	xs:ID			optional
Source	<pre>&lt;xs:element name="treatmentResponse"&gt;   &lt;xs:complexType&gt;</pre>				

```

<xs:sequence>
  <xs:group ref="generic.content"/>
  <xs:group ref="accessory.treatment.content"/>
</xs:sequence>
<xs:attributeGroup ref="generic.attrib"/>
</xs:complexType>
</xs:element>

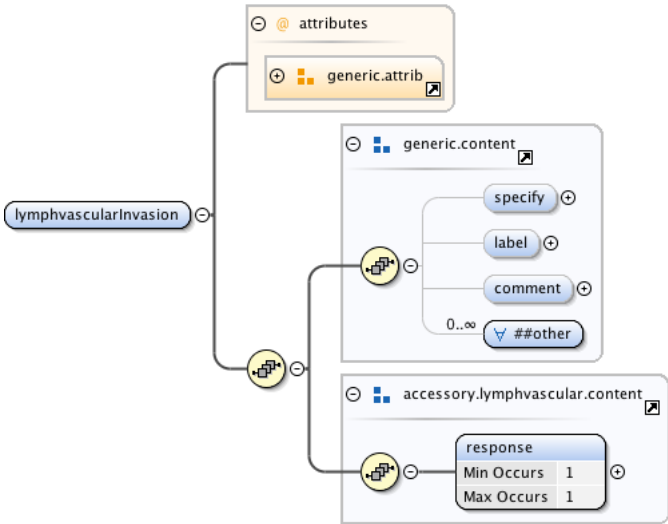
```

## Element `accessory.treatment.content` / `response`

Namespace	http://purl.org/obbr/pathology/cancer/				
Diagram					
Properties	content:	complex			
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'				
Children	comment, label, specify				
Instance	<pre>&lt;response id="" value=""&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/response&gt;</pre>				
Attributes	QName	Type	Fixed	Default	Use
	ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'				
	id	xs:ID			optional
	value	accessory.treatmentResponse.valueset			required
Source	<pre>&lt;xs:element name="response"&gt;   &lt;xs:complexType&gt;     &lt;xs:sequence&gt;       &lt;xs:group ref="generic.content"/&gt;     &lt;/xs:sequence&gt;     &lt;xs:attribute name="value" type="accessory.treatmentResponse.valueset" use="required"/&gt;   &lt;/xs:complexType&gt;   &lt;xs:attributeGroup ref="generic.attrib"/&gt; &lt;/xs:element&gt;</pre>				

## Element `accessory.lymphvascular` / `lymphvascularInvasion`

Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram					
Properties	content:	complex			
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , response				
Children	comment, label, response, specify				
Instance	<pre>&lt;lymphvascularInvasion id=""&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/lymphvascularInvasion&gt;</pre>				
Attributes	QName	Type	Fixed	Default	Use
	ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'				
	id	xs:ID			optional
Source	<pre>&lt;xs:element name="lymphvascularInvasion"&gt;   &lt;xs:complexType&gt;     &lt;xs:sequence&gt;       &lt;xs:group ref="generic.content"/&gt;       &lt;xs:group ref="accessory.lymphvascular.content"/&gt;     &lt;/xs:sequence&gt;     &lt;xs:attributeGroup ref="generic.attrib"/&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt;</pre>				

### Element accessory.lymphvascular.content / response

Namespace	http://purl.org/obbr/pathology/cancer/				
Diagram					

Properties	content: complex				
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than "http://purl.org/obbr/pathology/cancer/"				
Children	comment, label, specify				
Instance	<pre>&lt;response id="" value=""&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/response&gt;</pre>				
Attributes	QName	Type	Fixed	Default	Use
	ANY attribute from ANY namespace OTHER than "http://purl.org/obbr/pathology/cancer/"				
	id	xs:ID			optional
	value	accessory.lymphvascular.valueset			required
Source	<pre>&lt;xs:element name="response"&gt;   &lt;xs:complexType&gt;     &lt;xs:sequence&gt;       &lt;xs:group ref="generic.content"/&gt;     &lt;/xs:sequence&gt;     &lt;xs:attribute name="value" type="accessory.lymphvascular.valueset" use="required"/&gt;     &lt;xs:attributeGroup ref="generic.attrib"/&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt;</pre>				

## Element accessory.perineural / perineuralInvasion

Namespace	http://purl.org/obbr/pathology/cancer/				
Diagram	<p>The diagram illustrates the structure of the <code>perineuralInvasion</code> element. It is a complex element with the following components:</p> <ul style="list-style-type: none"><li><b>Attributes:</b> A group of attributes, including a <code>generic.attrib</code> element.</li><li><b>Content:</b> A sequence of elements:<ul style="list-style-type: none"><li><b>generic.content:</b> A group of elements including <code>specify</code>, <code>label</code>, <code>comment</code>, and <code>##other</code>. The <code>##other</code> element has a cardinality of <code>0..∞</code>.</li><li><b>accessory.perineural.content:</b> A group of elements including a <code>response</code> element. The <code>response</code> element has a <code>Min Occurs</code> of 1 and a <code>Max Occurs</code> of 1.</li></ul></li></ul>				
Properties	content:	complex			
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , response				
Children	comment, label, response, specify				
Instance	<pre>&lt;perineuralInvasion id=""&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/perineuralInvasion&gt;</pre>				
Attributes	QName	Type	Fixed	Default	Use
	ANY attribute from ANY namespace OTHER than 'http://purl.org/				

	QName	Type	Fixed	Default	Use
	<b>obbr/pathology/cancer/</b>				
	<b>id</b>	xs:ID			optional
Source	<pre> &lt;xs:element name="perineuralInvasion"&gt;   &lt;xs:complexType&gt;     &lt;xs:sequence&gt;       &lt;xs:group ref="generic.content"/&gt;       &lt;xs:group ref="accessory.perineural.content"/&gt;     &lt;/xs:sequence&gt;     &lt;xs:attributeGroup ref="generic.attrib"/&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt; </pre>				

## Element accessory.perineural.content / response

Namespace	http://purl.org/obbr/pathology/cancer/				
Diagram					
Properties	content:	complex			
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'				
Children	comment, label, specify				
Instance	<pre>&lt;response id="" value=""&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/response&gt;</pre>				
Attributes	QName	Type	Fixed	Default	Use
	ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'				
	id	xs:ID			optional
	value	accessory.perineural.valueset			required
Source	<pre>&lt;xs:element name="response"&gt;   &lt;xs:complexType&gt;     &lt;xs:sequence&gt;       &lt;xs:group ref="generic.content"/&gt;     &lt;/xs:sequence&gt;     &lt;xs:attribute name="value" type="accessory.perineural.valueset" use="required"/&gt;     &lt;xs:attributeGroup ref="generic.attrib"/&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt;</pre>				

## Element additional

Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram	<p>The diagram illustrates the structure of the 'additional' element. It is a complex type containing two main groups: 'generic.attrib' and 'generic.content'. The 'generic.attrib' group contains a single attribute 'generic.attrib'. The 'generic.content' group contains four elements: 'specify', 'label', 'comment', and '##other'. The 'specify', 'label', and 'comment' elements are optional (indicated by a circle with a minus sign). The '##other' element is optional and has a cardinality of 0..∞. The 'additional' element itself is optional (indicated by a circle with a minus sign). The 'additional.content' group contains a 'finding' element with a cardinality of 0..∞. The 'finding' element has a table with 'Min Occurs' 0 and 'Max Occurs' unbounded.</p>				
Properties	content:	complex			
Used by	Element Group	diagnosis.content			
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , finding*				
Children	comment, finding, label, specify				
Instance	<pre>&lt;additional id=""&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/additional&gt;</pre>				
Attributes	QName	Type	Fixed	Default	Use
	ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'				
	id	xs:ID			optional
Source	<pre>&lt;xs:element name="additional"&gt;   &lt;xs:complexType&gt;     &lt;xs:sequence&gt;       &lt;xs:group ref="generic.content"/&gt;       &lt;xs:group ref="additional.content"/&gt;     &lt;/xs:sequence&gt;     &lt;xs:attributeGroup ref="generic.attrib"/&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt;</pre>				

### Element additional.content / finding

Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram											
Properties	<table><tr><td>content:</td><td>complex</td></tr><tr><td>minOccurs:</td><td>0</td></tr><tr><td>maxOccurs:</td><td>unbounded</td></tr></table>					content:	complex	minOccurs:	0	maxOccurs:	unbounded
content:	complex										
minOccurs:	0										
maxOccurs:	unbounded										
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , response										
Children	comment, label, response, specify										
Instance	<pre>&lt;finding id="" of=""&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/finding&gt;</pre>										
Attributes	QName	Type	Fixed	Default	Use						
	ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'										
	id	xs:ID			optional						
	of	additional.finding.of.valueset			optional						
Source	<pre>&lt;xs:element maxOccurs="unbounded" minOccurs="0" name="finding"&gt;   &lt;xs:complexType&gt;     &lt;xs:sequence&gt;       &lt;xs:group ref="generic.content"/&gt;       &lt;xs:group ref="additional.finding.content"/&gt;     &lt;/xs:sequence&gt;     &lt;xs:attribute name="of" type="additional.finding.of.valueset"/&gt;     &lt;xs:attributeGroup ref="generic.attrib"/&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt;</pre>										

### Element additional.finding.content / response

Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram					
Properties	content:	complex			
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'				
Children	comment, label, specify				
Instance	<pre>&lt;response value=""&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/response&gt;</pre>				
Attributes	QName	Type	Fixed	Default	Use
	value	additional.finding.valueset			required
Source	<pre>&lt;xs:element name="response"&gt;   &lt;xs:complexType&gt;     &lt;xs:sequence&gt;       &lt;xs:group ref="generic.content"/&gt;     &lt;/xs:sequence&gt;     &lt;xs:attribute name="value" type="additional.finding.valueset" use="required"/&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt;</pre>				

## Element distant

Namespace	http://purl.org/obbr/pathology/cancer/		
Diagram			
Properties	content:	complex	
Used by	Element Group	diagnosis.content	
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than "http://purl.org/obbr/pathology/cancer/" , status		
Children	comment, label, specify, status		



Instance	<pre>&lt;distant id=""&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/distant&gt;</pre>				
Attributes	QName	Type	Fixed	Default	Use
	ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'				
	id	xs:ID			optional
Source	<pre>&lt;xs:element name="distant"&gt;   &lt;xs:complexType&gt;     &lt;xs:sequence&gt;       &lt;xs:group ref="generic.content"/&gt;       &lt;xs:group ref="distant.content"/&gt;     &lt;/xs:sequence&gt;     &lt;xs:attributeGroup ref="generic.attrib"/&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt;</pre>				

### Element distant.status / status

Namespace	http://purl.org/obbr/pathology/cancer/				
Diagram					
Properties	content:	complex			
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than "http://purl.org/obbr/pathology/cancer/" , result , response{1,3}				
Children	comment, label, response, result, specify				
Instance	<pre>&lt;status of="" on=""&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/status&gt;</pre>				
Attributes	QName	Type	Fixed	Default	Use
	of	distant.status.of.valueset			required

	QName	Type	Fixed	Default	Use
	<b>on</b>	laterality.valueset			optional
Source	<pre> &lt;xs:element name="status"&gt;   &lt;xs:complexType&gt;     &lt;xs:sequence&gt;       &lt;xs:group ref="generic.content"/&gt;       &lt;xs:group ref="distant.status.result"/&gt;       &lt;xs:group minOccurs="0" ref="size"/&gt;     &lt;/xs:sequence&gt;     &lt;xs:attribute name="of" type="distant.status.of.valueset" use="required"/&gt;     &lt;xs:attribute name="on" type="laterality.valueset"/&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt; </pre>				

## Element distant.status.result / result

Namespace	http://purl.org/obbr/pathology/cancer/				
Diagram					
Properties	content:	complex			
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , response				
Children	comment, label, response, specify				
Instance	<pre>&lt;result date="" duration="" end="" for="" start=""&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/result&gt;</pre>				
Attributes	QName	Type	Fixed	Default	Use
	date	date.valueset			optional
	duration	xs:duration			optional
	end	date.valueset			optional
	for	distant.status.result.for.valueset			optional
	start	date.valueset			optional
Source	<pre>&lt;xs:element name="result"&gt;   &lt;xs:complexType&gt;     &lt;xs:sequence&gt;       &lt;xs:group ref="generic.content"/&gt;       &lt;xs:group ref="distant.status.result.response"/&gt;     &lt;/xs:sequence&gt;     &lt;xs:attribute name="for" type="distant.status.result.for.valueset"/&gt;     &lt;xs:attributeGroup ref="date"/&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt;</pre>				

**Element distant.status.result.response / response**

Namespace	http://purl.org/obbr/pathology/cancer/				
Diagram					
Properties	content:	complex			
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than "http://purl.org/obbr/pathology/cancer/"				
Children	comment, label, specify				
Instance	<pre>&lt;response value=" "&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/response&gt;</pre>				
Attributes	QName	Type	Fixed	Default	Use
	value	union of(val:base.values, val:qualitative.values)			required
Source	<pre>&lt;xs:element name="response"&gt;   &lt;xs:complexType&gt;     &lt;xs:sequence&gt;       &lt;xs:group ref="generic.content" /&gt;     &lt;/xs:sequence&gt;     &lt;xs:attribute name="value" use="required"&gt;       &lt;xs:simpleType&gt;         &lt;xs:union memberTypes="val:base.values val:qualitative.values"/&gt;       &lt;/xs:simpleType&gt;     &lt;/xs:attribute&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt;</pre>				

**Element extent**

Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram	<pre>graph TD     extent((extent)) --&gt; G1(( ))     G1 --&gt; G2(( ))     G2 --&gt; G3(( ))     G3 --&gt; G4(( ))     G4 --&gt; G5(( ))     G5 --&gt; G6(( ))     G6 --&gt; G7(( ))     G7 --&gt; G8(( ))     G8 --&gt; G9(( ))     G9 --&gt; G10(( ))     G10 --&gt; G11(( ))     G11 --&gt; G12(( ))     G12 --&gt; G13(( ))     G13 --&gt; G14(( ))     G14 --&gt; G15(( ))     G15 --&gt; G16(( ))     G16 --&gt; G17(( ))     G17 --&gt; G18(( ))     G18 --&gt; G19(( ))     G19 --&gt; G20(( ))     G20 --&gt; G21(( ))     G21 --&gt; G22(( ))     G22 --&gt; G23(( ))     G23 --&gt; G24(( ))     G24 --&gt; G25(( ))     G25 --&gt; G26(( ))     G26 --&gt; G27(( ))     G27 --&gt; G28(( ))     G28 --&gt; G29(( ))     G29 --&gt; G30(( ))     G30 --&gt; G31(( ))     G31 --&gt; G32(( ))     G32 --&gt; G33(( ))     G33 --&gt; G34(( ))     G34 --&gt; G35(( ))     G35 --&gt; G36(( ))     G36 --&gt; G37(( ))     G37 --&gt; G38(( ))     G38 --&gt; G39(( ))     G39 --&gt; G40(( ))     G40 --&gt; G41(( ))     G41 --&gt; G42(( ))     G42 --&gt; G43(( ))     G43 --&gt; G44(( ))     G44 --&gt; G45(( ))     G45 --&gt; G46(( ))     G46 --&gt; G47(( ))     G47 --&gt; G48(( ))     G48 --&gt; G49(( ))     G49 --&gt; G50(( ))     G50 --&gt; G51(( ))     G51 --&gt; G52(( ))     G52 --&gt; G53(( ))     G53 --&gt; G54(( ))     G54 --&gt; G55(( ))     G55 --&gt; G56(( ))     G56 --&gt; G57(( ))     G57 --&gt; G58(( ))     G58 --&gt; G59(( ))     G59 --&gt; G60(( ))     G60 --&gt; G61(( ))     G61 --&gt; G62(( ))     G62 --&gt; G63(( ))     G63 --&gt; G64(( ))     G64 --&gt; G65(( ))     G65 --&gt; G66(( ))     G66 --&gt; G67(( ))     G67 --&gt; G68(( ))     G68 --&gt; G69(( ))     G69 --&gt; G70(( ))     G70 --&gt; G71(( ))     G71 --&gt; G72(( ))     G72 --&gt; G73(( ))     G73 --&gt; G74(( ))     G74 --&gt; G75(( ))     G75 --&gt; G76(( ))     G76 --&gt; G77(( ))     G77 --&gt; G78(( ))     G78 --&gt; G79(( ))     G79 --&gt; G80(( ))     G80 --&gt; G81(( ))     G81 --&gt; G82(( ))     G82 --&gt; G83(( ))     G83 --&gt; G84(( ))     G84 --&gt; G85(( ))     G85 --&gt; G86(( ))     G86 --&gt; G87(( ))     G87 --&gt; G88(( ))     G88 --&gt; G89(( ))     G89 --&gt; G90(( ))     G90 --&gt; G91(( ))     G91 --&gt; G92(( ))     G92 --&gt; G93(( ))     G93 --&gt; G94(( ))     G94 --&gt; G95(( ))     G95 --&gt; G96(( ))     G96 --&gt; G97(( ))     G97 --&gt; G98(( ))     G98 --&gt; G99(( ))     G99 --&gt; G100(( ))     G100 --&gt; G101(( ))     G101 --&gt; G102(( ))     G102 --&gt; G103(( ))     G103 --&gt; G104(( ))     G104 --&gt; G105(( ))     G105 --&gt; G106(( ))     G106 --&gt; G107(( ))     G107 --&gt; G108(( ))     G108 --&gt; G109(( ))     G109 --&gt; G110(( ))     G110 --&gt; G111(( ))     G111 --&gt; G112(( ))     G112 --&gt; G113(( ))     G113 --&gt; G114(( ))     G114 --&gt; G115(( ))     G115 --&gt; G116(( ))     G116 --&gt; G117(( ))     G117 --&gt; G118(( ))     G118 --&gt; G119(( ))     G119 --&gt; G120(( ))     G120 --&gt; G121(( ))     G121 --&gt; G122(( ))     G122 --&gt; G123(( ))     G123 --&gt; G124(( ))     G124 --&gt; G125(( ))     G125 --&gt; G126(( ))     G126 --&gt; G127(( ))     G127 --&gt; G128(( ))     G128 --&gt; G129(( ))     G129 --&gt; G130(( ))     G130 --&gt; G131(( ))     G131 --&gt; G132(( ))     G132 --&gt; G133(( ))     G133 --&gt; G134(( ))     G134 --&gt; G135(( ))     G135 --&gt; G136(( ))     G136 --&gt; G137(( ))     G137 --&gt; G138(( ))     G138 --&gt; G139(( ))     G139 --&gt; G140(( ))     G140 --&gt; G141(( ))     G141 --&gt; G142(( ))     G142 --&gt; G143(( ))     G143 --&gt; G144(( ))     G144 --&gt; G145(( ))     G145 --&gt; G146(( ))     G146 --&gt; G147(( ))     G147 --&gt; G148(( ))     G148 --&gt; G149(( ))     G149 --&gt; G150(( ))     G150 --&gt; G151(( ))     G151 --&gt; G152(( ))     G152 --&gt; G153(( ))     G153 --&gt; G154(( ))     G154 --&gt; G155(( ))     G155 --&gt; G156(( ))     G156 --&gt; G157(( ))     G157 --&gt; G158(( ))     G158 --&gt; G159(( ))     G159 --&gt; G160(( ))     G160 --&gt; G161(( ))     G161 --&gt; G162(( ))     G162 --&gt; G163(( ))     G163 --&gt; G164(( ))     G164 --&gt; G165(( ))     G165 --&gt; G166(( ))     G166 --&gt; G167(( ))     G167 --&gt; G168(( ))     G168 --&gt; G169(( ))     G169 --&gt; G170(( ))     G170 --&gt; G171(( ))     G171 --&gt; G172(( ))     G172 --&gt; G173(( ))     G173 --&gt; G174(( ))     G174 --&gt; G175(( ))     G175 --&gt; G176(( ))     G176 --&gt; G177(( ))     G177 --&gt; G178(( ))     G178 --&gt; G179(( ))     G179 --&gt; G180(( ))     G180 --&gt; G181(( ))     G181 --&gt; G182(( ))     G182 --&gt; G183(( ))     G183 --&gt; G184(( ))     G184 --&gt; G185(( ))     G185 --&gt; G186(( ))     G186 --&gt; G187(( ))     G187 --&gt; G188(( ))     G188 --&gt; G189(( ))     G189 --&gt; G190(( ))     G190 --&gt; G191(( ))     G191 --&gt; G192(( ))     G192 --&gt; G193(( ))     G193 --&gt; G194(( ))     G194 --&gt; G195(( ))     G195 --&gt; G196(( ))     G196 --&gt; G197(( ))     G197 --&gt; G198(( ))     G198 --&gt; G199(( ))     G199 --&gt; G200(( ))     G200 --&gt; G201(( ))     G201 --&gt; G202(( ))     G202 --&gt; G203(( ))     G203 --&gt; G204(( ))     G204 --&gt; G205(( ))     G205 --&gt; G206(( ))     G206 --&gt; G207(( ))     G207 --&gt; G208(( ))     G208 --&gt; G209(( ))     G209 --&gt; G210(( ))     G210 --&gt; G211(( ))     G211 --&gt; G212(( ))     G212 --&gt; G213(( ))     G213 --&gt; G214(( ))     G214 --&gt; G215(( ))     G215 --&gt; G216(( ))     G216 --&gt; G217(( ))     G217 --&gt; G218(( ))     G218 --&gt; G219(( ))     G219 --&gt; G220(( ))     G220 --&gt; G221(( ))     G221 --&gt; G222(( ))     G222 --&gt; G223(( ))     G223 --&gt; G224(( ))     G224 --&gt; G225(( ))     G225 --&gt; G226(( ))     G226 --&gt; G227(( ))     G227 --&gt; G228(( ))     G228 --&gt; G229(( ))     G229 --&gt; G230(( ))     G230 --&gt; G231(( ))     G231 --&gt; G232(( ))     G232 --&gt; G233(( ))     G233 --&gt; G234(( ))     G234 --&gt; G235(( ))     G235 --&gt; G236(( ))     G236 --&gt; G237(( ))     G237 --&gt; G238(( ))     G238 --&gt; G239(( ))     G239 --&gt; G240(( ))     G240 --&gt; G241(( ))     G241 --&gt; G242(( ))     G242 --&gt; G243(( ))     G243 --&gt; G244(( ))     G244 --&gt; G245(( ))     G245 --&gt; G246(( ))     G246 --&gt; G247(( ))     G247 --&gt; G248(( ))     G248 --&gt; G249(( ))     G249 --&gt; G250(( ))     G250 --&gt; G251(( ))     G251 --&gt; G252(( ))     G252 --&gt; G253(( ))     G253 --&gt; G254(( ))     G254 --&gt; G255(( ))     G255 --&gt; G256(( ))     G256 --&gt; G257(( ))     G257 --&gt; G258(( ))     G258 --&gt; G259(( ))     G259 --&gt; G260(( ))     G260 --&gt; G261(( ))     G261 --&gt; G262(( ))     G262 --&gt; G263(( ))     G263 --&gt; G264(( ))     G264 --&gt; G265(( ))     G265 --&gt; G266(( ))     G266 --&gt; G267(( ))     G267 --&gt; G268(( ))     G268 --&gt; G269(( ))     G269 --&gt; G270(( ))     G270 --&gt; G271(( ))     G271 --&gt; G272(( ))     G272 --&gt; G273(( ))     G273 --&gt; G274(( ))     G274 --&gt; G275(( ))     G275 --&gt; G276(( ))     G276 --&gt; G277(( ))     G277 --&gt; G278(( ))     G278 --&gt; G279(( ))     G279 --&gt; G280(( ))     G280 --&gt; G281(( ))     G281 --&gt; G282(( ))     G282 --&gt; G283(( ))     G283 --&gt; G284(( ))     G284 --&gt; G285(( ))     G285 --&gt; G286(( ))     G286 --&gt; G287(( ))     G287 --&gt; G288(( ))     G288 --&gt; G289(( ))     G289 --&gt; G290(( ))     G290 --&gt; G291(( ))     G291 --&gt; G292(( ))     G292 --&gt; G293(( ))     G293 --&gt; G294(( ))     G294 --&gt; G295(( ))     G295 --&gt; G296(( ))     G296 --&gt; G297(( ))     G297 --&gt; G298(( ))     G298 --&gt; G299(( ))     G299 --&gt; G300(( ))     G300 --&gt; G301(( ))     G301 --&gt; G302(( ))     G302 --&gt; G303(( ))     G303 --&gt; G304(( ))     G304 --&gt; G305(( ))     G305 --&gt; G306(( ))     G306 --&gt; G307(( ))     G307 --&gt; G308(( ))     G308 --&gt; G309(( ))     G309 --&gt; G310(( ))     G310 --&gt; G311(( ))     G311 --&gt; G312(( ))     G312 --&gt; G313(( ))     G313 --&gt; G314(( ))     G314 --&gt; G315(( ))     G315 --&gt; G316(( ))     G316 --&gt; G317(( ))     G317 --&gt; G318(( ))     G318 --&gt; G319(( ))     G319 --&gt; G320(( ))     G320 --&gt; G321(( ))     G321 --&gt; G322(( ))     G322 --&gt; G323(( ))     G323 --&gt; G324(( ))     G324 --&gt; G325(( ))     G325 --&gt; G326(( ))     G326 --&gt; G327(( ))     G327 --&gt; G328(( ))     G328 --&gt; G329(( ))     G329 --&gt; G330(( ))     G330 --&gt; G331(( ))     G331 --&gt; G332(( ))     G332 --&gt; G333(( ))     G333 --&gt; G334(( ))     G334 --&gt; G335(( ))     G335 --&gt; G336(( ))     G336 --&gt; G337(( ))     G337 --&gt; G338(( ))     G338 --&gt; G339(( ))     G339 --&gt; G340(( ))     G340 --&gt; G341(( ))     G341 --&gt; G342(( ))     G342 --&gt; G343(( ))     G343 --&gt; G344(( ))     G344 --&gt; G345(( ))     G345 --&gt; G346(( ))     G346 --&gt; G347(( ))     G347 --&gt; G348(( ))     G348 --&gt; G349(( ))     G349 --&gt; G350(( ))     G350 --&gt; G351(( ))     G351 --&gt; G352(( ))     G352 --&gt; G353(( ))     G353 --&gt; G354(( ))     G354 --&gt; G355(( ))     G355 --&gt; G356(( ))     G356 --&gt; G357(( ))     G357 --&gt; G358(( ))     G358 --&gt; G359(( ))     G359 --&gt; G360(( ))     G360 --&gt; G361(( ))     G361 --&gt; G362(( ))     G362 --&gt; G363(( ))     G363 --&gt; G364(( ))     G364 --&gt; G365(( ))     G365 --&gt; G366(( ))     G366 --&gt; G367(( ))     G367 --&gt; G368(( ))     G368 --&gt; G369(( ))     G369 --&gt; G370(( ))     G370 --&gt; G371(( ))     G371 --&gt; G372(( ))     G372 --&gt; G373(( ))     G373 --&gt; G374(( ))     G374 --&gt; G375(( ))     G375 --&gt; G376(( ))     G376 --&gt; G377(( ))     G377 --&gt; G378(( ))     G378 --&gt; G379(( ))     G379 --&gt; G380(( ))     G380 --&gt; G381(( ))     G381 --&gt; G382(( ))     G382 --&gt; G383(( ))     G383 --&gt; G384(( ))     G384 --&gt; G385(( ))     G385 --&gt; G386(( ))     G386 --&gt; G387(( ))     G387 --&gt; G388(( ))     G388 --&gt; G389(( ))     G389 --&gt; G390(( ))     G390 --&gt; G391(( ))     G391 --&gt; G392(( ))     G392 --&gt; G393(( ))     G393 --&gt; G394(( ))     G394 --&gt; G395(( ))     G395 --&gt; G396(( ))     G396 --&gt; 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G709(( ))     G709 --&gt; G710(( ))     G710 --&gt; G711(( ))     G711 --&gt; G712(( ))     G712 --&gt; G713(( ))     G713 --&gt; G714(( ))     G714 --&gt; G715(( ))     G715 --&gt; G716(( ))     G716 --&gt; G717(( ))     G717 --&gt; G718(( ))     G718 --&gt; G719(( ))     G719 --&gt; G720(( ))     G720 --&gt; G721(( ))     G721 --&gt; G722(( ))     G722 --&gt; G723(( ))     G723 --&gt; G724(( ))     G724 --&gt; G725(( ))     G725 --&gt; G726(( ))     G726 --&gt; G727(( ))     G727 --&gt; G728(( ))     G728 --&gt; G729(( ))     G729 --&gt; G730(( ))     G730 --&gt; G731(( ))     G731 --&gt; G732(( ))     G732 --&gt; G733(( ))     G733 --&gt; G734(( ))     G734 --&gt; G735(( ))     G735 --&gt; G736(( ))     G736 --&gt; G737(( ))     G737 --&gt; G738(( ))     G738 --&gt; G739(( ))     G739 --&gt; G740(( ))     G740 --&gt; G741(( ))     G741 --&gt; G742(( ))     G742 --&gt; G743(( ))     G743 --&gt; G744(( ))     G744 --&gt; G745(( ))     G745 --&gt; G746(( ))     G746 --&gt; G747(( ))     G747 --&gt; 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### Element contiguousExtension / contiguousExtension

Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram	
Properties	content: complex
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , status
Children	comment, label, specify, status
Instance	<pre> &lt;contiguousExtension&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/contiguousExtension&gt; </pre>
Source	<pre> &lt;xs:element name="contiguousExtension"&gt;   &lt;xs:complexType&gt;     &lt;xs:sequence&gt;       &lt;xs:group ref="generic.content" /&gt;       &lt;xs:group maxOccurs="unbounded" minOccurs="0" ref="contiguousExtensionStatus" /&gt;     &lt;/xs:sequence&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt; </pre>

### Element contiguousExtensionStatus / status

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Properties	content: complex
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , result , distance
Children	comment, distance, label, result, specify

Instance	<pre> &lt;status of=""&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/status&gt; </pre>				
Attributes	<b>QName</b>	<b>Type</b>	<b>Fixed</b>	<b>Default</b>	<b>Use</b>
	<b>of</b>	extent.status.of.valueset			optional
Source	<pre> &lt;xs:element name="status"&gt;   &lt;xs:complexType&gt;     &lt;xs:sequence maxOccurs="unbounded"&gt;       &lt;xs:group ref="generic.content"/&gt;       &lt;xs:group ref="contiguousExtension.status.result"/&gt;       &lt;xs:group minOccurs="0" ref="contiguousExtension.status.distance"/&gt;     &lt;/xs:sequence&gt;     &lt;xs:attribute name="of" type="extent.status.of.valueset"/&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt; </pre>				

## Element contiguousExtension.status.result / result

Namespace	http://purl.org/obbr/pathology/cancer/				
Diagram					
Properties	content:	complex			
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , response				
Children	comment, label, response, specify				
Instance	<pre>&lt;result for="" on=""&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/result&gt;</pre>				
Attributes	QName	Type	Fixed	Default	Use
	for	union of(val:base.values, restriction of xs:token)			required
	on	laterality.valueset			optional
Source	<pre>&lt;xs:element name="result"&gt;   &lt;xs:complexType&gt;     &lt;xs:sequence&gt;       &lt;xs:group ref="generic.content"/&gt;       &lt;xs:group maxOccurs="unbounded" minOccurs="0" ref="contiguousExtension.status.result.content"/&gt;     &lt;/xs:sequence&gt;     &lt;xs:attribute name="for" use="required"&gt;</pre>				

```

<xs:simpleType>
  <xs:union memberTypes="val:base.values">
    <xs:simpleType>
      <xs:restriction base="xs:token">
        <xs:enumeration value="invades"/>
        <xs:enumeration value="adheres"/>
        <xs:enumeration value="specify"/>
        <xs:enumeration value="inapplicable"/>
        <xs:enumeration value="unreported"/>
      </xs:restriction>
    </xs:simpleType>
  </xs:union>
</xs:simpleType>
</xs:attribute>
<xs:attribute name="on" type="laterality.valueset"/>
</xs:complexType>
</xs:element>

```

## Element contiguousExtension.status.result.content / response

Namespace	http://purl.org/obbr/pathology/cancer/				
Diagram					
Properties	content:	complex			
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'				
Children	comment, label, specify				
Instance	<pre>&lt;response extent="" value=""&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/response&gt;</pre>				
Attributes	QName	Type	Fixed	Default	Use
	extent	focality.valueset			optional
	value	union of(val:base.values, val:qualitative.values)			required
Source	<pre>&lt;xs:element name="response"&gt;   &lt;xs:complexType&gt;     &lt;xs:group ref="generic.content"/&gt;     &lt;xs:attribute name="value" use="required"&gt;       &lt;xs:simpleType&gt;         &lt;xs:union memberTypes="val:base.values val:qualitative.values"/&gt;       &lt;/xs:simpleType&gt;     &lt;/xs:attribute&gt;     &lt;xs:attribute name="extent" type="focality.valueset"/&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt;</pre>				

## Element contiguousExtension.status.distance / distance

Namespace	http://purl.org/obbr/pathology/cancer/
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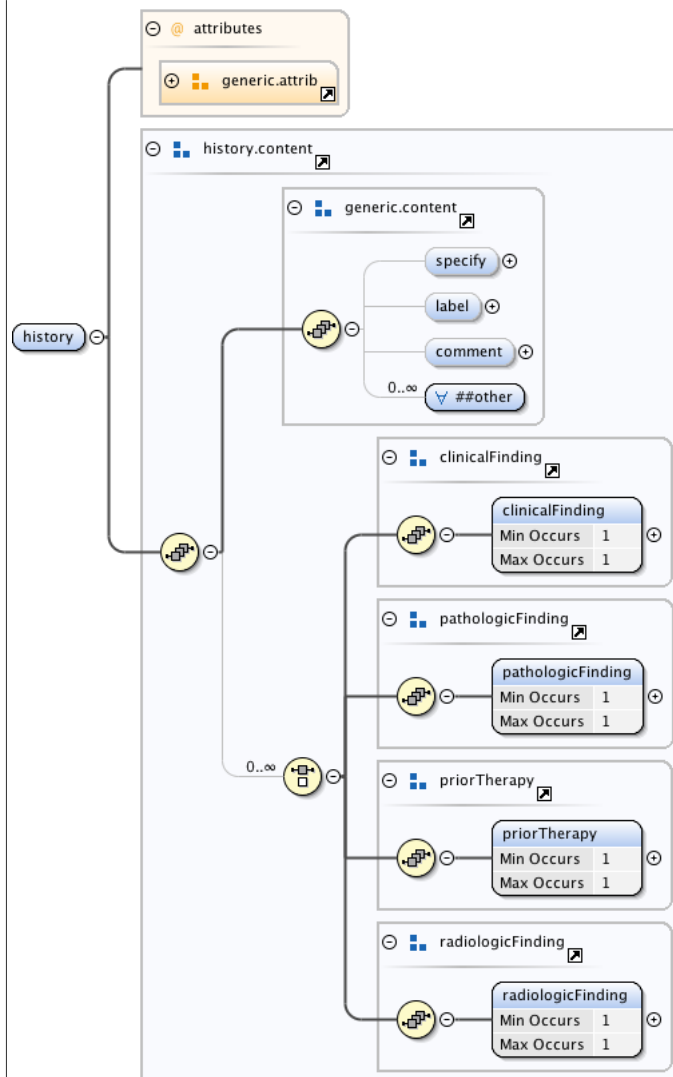
Diagram					
Properties	content:	complex			
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , response				
Children	comment, label, response, specify				
Instance	<pre>&lt;distance for=" " from=" "&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/distance&gt;</pre>				
Attributes	QName	Type	Fixed	Default	Use
	for	restriction of xs:token			optional
	from	val:base.values			optional
Source	<pre>&lt;xs:element name="distance"&gt;   &lt;xs:complexType&gt;     &lt;xs:sequence&gt;       &lt;xs:group ref="generic.content"/&gt;       &lt;xs:group ref="length"/&gt;     &lt;/xs:sequence&gt;     &lt;xs:attribute name="for"&gt;       &lt;xs:simpleType&gt;         &lt;xs:restriction base="xs:token"&gt;           &lt;xs:enumeration value="invades"/&gt;           &lt;xs:enumeration value="adheres"/&gt;         &lt;/xs:restriction&gt;       &lt;/xs:simpleType&gt;     &lt;/xs:attribute&gt;     &lt;xs:attribute name="from" type="val:base.values"/&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt;</pre>				

## Element history

Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram



Properties	content: complex				
Used by	Element Group diagnosis.content				
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , ((clinicalFinding)   (pathologicFinding)   (priorTherapy)   (radiologicFinding))				
Children	clinicalFinding, comment, label, pathologicFinding, priorTherapy, radiologicFinding, specify				
Instance	<pre>&lt;history id=""&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/history&gt;</pre>				
Attributes	QName	Type	Fixed	Default	Use
	ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'				
	id	xs:ID			optional
Source	<pre>&lt;xs:element name="history"&gt;   &lt;xs:complexType&gt;     &lt;xs:group ref="history.content"/&gt;     &lt;xs:attributeGroup ref="generic.attrib"/&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt;</pre>				

**Element clinicalFinding / clinicalFinding**

Namespace	http://purl.org/obbr/pathology/cancer/				
Diagram					
Properties	content:	complex			
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , response				
Children	comment, label, response, specify				
Instance	<pre>&lt;clinicalFinding date="" duration="" end="" start=""&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/clinicalFinding&gt;</pre>				
Attributes	QName	Type	Fixed	Default	Use
	date	date.valueset			optional
	duration	xs:duration			optional
	end	date.valueset			optional
	start	date.valueset			optional
Source	<pre>&lt;xs:element name="clinicalFinding"&gt;   &lt;xs:complexType&gt;     &lt;xs:sequence&gt;       &lt;xs:group ref="generic.content"/&gt;       &lt;xs:group ref="clinicalFinding.content"/&gt;     &lt;/xs:sequence&gt;     &lt;xs:attributeGroup ref="date"/&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt;</pre>				

**Element clinicalFinding.content / response**

Namespace	http://purl.org/obbr/pathology/cancer/				
Diagram					

Properties	content: complex				
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'				
Children	comment, label, specify				
Instance	<pre>&lt;response value=""&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/response&gt;</pre>				
Attributes	QName	Type	Fixed	Default	Use
	value	restriction of xs:token			required
Source	<pre>&lt;xs:element name="response"&gt;   &lt;xs:complexType&gt;     &lt;xs:group ref="generic.content" /&gt;     &lt;xs:attribute name="value" use="required"&gt;       &lt;xs:simpleType&gt;         &lt;xs:restriction base="xs:token"&gt;           &lt;xs:enumeration value="specify" /&gt;           &lt;xs:enumeration value="inapplicable" /&gt;           &lt;xs:enumeration value="unreported" /&gt;         &lt;/xs:restriction&gt;       &lt;/xs:simpleType&gt;     &lt;/xs:attribute&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt;</pre>				

### Element pathologicFinding / pathologicFinding

Namespace	http://purl.org/obbr/pathology/cancer/				
Diagram					
Properties	content: complex				
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , response				
Children	comment, label, response, specify				
Instance	<pre>&lt;pathologicFinding date="" duration="" end="" start=""&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/pathologicFinding&gt;</pre>				
Attributes	QName	Type	Fixed	Default	Use
	date	date.valueset			optional
	duration	xs:duration			optional
	end	date.valueset			optional
	start	date.valueset			optional
Source	<xs:element name="pathologicFinding">				

```

<xs:complexType>
  <xs:sequence>
    <xs:group ref="generic.content" />
    <xs:group ref="pathologicFinding.content" />
  </xs:sequence>
  <xs:attributeGroup ref="date" />
</xs:complexType>
</xs:element>

```

## Element pathologicFinding.content / response

Namespace	http://purl.org/obbr/pathology/cancer/				
Diagram					
Properties	content:	complex			
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'				
Children	comment, label, specify				
Instance	<pre>&lt;response value=""&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/response&gt;</pre>				
Attributes	QName	Type	Fixed	Default	Use
	value	restriction of xs:token			required
Source	<pre>&lt;xs:element name="response"&gt;   &lt;xs:complexType&gt;     &lt;xs:group ref="generic.content"/&gt;     &lt;xs:attribute name="value" use="required"&gt;       &lt;xs:simpleType&gt;         &lt;xs:restriction base="xs:token"&gt;           &lt;xs:enumeration value="specify"/&gt;           &lt;xs:enumeration value="inapplicable"/&gt;           &lt;xs:enumeration value="unreported"/&gt;         &lt;/xs:restriction&gt;       &lt;/xs:simpleType&gt;     &lt;/xs:attribute&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt;</pre>				

## Element priorTherapy / priorTherapy

Namespace	http://purl.org/obbr/pathology/cancer/
-----------	----------------------------------------

Diagram					
Properties	content:	complex			
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , response				
Children	comment, label, response, specify				
Instance	<pre>&lt;priorTherapy date="" duration="" end="" start=""&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/priorTherapy&gt;</pre>				
Attributes	QName	Type	Fixed	Default	Use
	date	date.valueset			optional
	duration	xs:duration			optional
	end	date.valueset			optional
	start	date.valueset			optional
Source	<pre>&lt;xs:element name="priorTherapy"&gt;   &lt;xs:complexType&gt;     &lt;xs:sequence&gt;       &lt;xs:group ref="generic.content"/&gt;       &lt;xs:group ref="priorTherapy.content"/&gt;     &lt;/xs:sequence&gt;     &lt;xs:attributeGroup ref="date"/&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt;</pre>				

### Element priorTherapy.content / response

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Properties	content: complex
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'

Children	comment, label, specify				
Instance	<pre>&lt;response value=" "&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/response&gt;</pre>				
Attributes	QName	Type	Fixed	Default	Use
	value	restriction of xs:token			required
Source	<pre>&lt;xs:element name="response"&gt;   &lt;xs:complexType&gt;     &lt;xs:group ref="generic.content" /&gt;     &lt;xs:attribute name="value" use="required"&gt;       &lt;xs:simpleType&gt;         &lt;xs:restriction base="xs:token"&gt;           &lt;xs:enumeration value="specify" /&gt;           &lt;xs:enumeration value="inapplicable" /&gt;           &lt;xs:enumeration value="unreported" /&gt;           &lt;xs:enumeration value="chemotherapy" /&gt;           &lt;xs:enumeration value="radiation" /&gt;           &lt;xs:enumeration value="surgery" /&gt;         &lt;/xs:restriction&gt;       &lt;/xs:simpleType&gt;     &lt;/xs:attribute&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt;</pre>				

## Element radiologicFinding / radiologicFinding

Namespace	http://purl.org/obbr/pathology/cancer/					
Diagram						
Properties	content: complex					
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , response					
Children	comment, label, response, specify					
Instance	<pre>&lt;radiologicFinding date="" duration="" end="" start=""&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/radiologicFinding&gt;</pre>					
Attributes	QName	Type	Fixed	Default	Use	
	date	date.valueset			optional	
	duration	xs:duration			optional	
	end	date.valueset			optional	
	start	date.valueset			optional	
Source	<pre>&lt;xs:element name="radiologicFinding"&gt;   &lt;xs:complexType&gt;     &lt;xs:sequence&gt;</pre>					

```

<xs:group ref="generic.content" />
<xs:group ref="radiologicFinding.content" />
</xs:sequence>
<xs:attributeGroup ref="date" />
</xs:complexType>
</xs:element>

```

## Element radiologicFinding.content / response

Namespace	http://purl.org/obbr/pathology/cancer/				
Diagram					
Properties	content:	complex			
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'				
Children	comment, label, specify				
Instance	<pre>&lt;response value=" "&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/response&gt;</pre>				
Attributes	QName	Type	Fixed	Default	Use
	value	restriction of xs:token			required
Source	<pre>&lt;xs:element name="response"&gt;   &lt;xs:complexType&gt;     &lt;xs:group ref="generic.content"/&gt;     &lt;xs:attribute name="value" use="required"&gt;       &lt;xs:simpleType&gt;         &lt;xs:restriction base="xs:token"&gt;           &lt;xs:enumeration value="specify"/&gt;           &lt;xs:enumeration value="inapplicable"/&gt;           &lt;xs:enumeration value="unreported"/&gt;         &lt;/xs:restriction&gt;       &lt;/xs:simpleType&gt;     &lt;/xs:attribute&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt;</pre>				

## Element margin

Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram					
Properties	content:	complex			
Used by	Element Group	diagnosis.content			
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , status				
Children	comment, label, specify, status				
Instance	<pre>&lt;margin id=""&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/margin&gt;</pre>				
Attributes	QName	Type	Fixed	Default	Use
	ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'				
	id	xs:ID			optional
Source	<pre>&lt;xs:element name="margin"&gt;   &lt;xs:complexType&gt;     &lt;xs:group ref="margin.content"/&gt;     &lt;xs:attributeGroup ref="generic.attrib"/&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt;</pre>				

## Element margins.margin.status / status

Namespace	http://purl.org/obbr/pathology/cancer/
-----------	----------------------------------------



Diagram					
Properties	content:	complex			
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , result , distance				
Children	comment, distance, label, result, specify				
Instance	<pre>&lt;status of="" on=""&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/status&gt;</pre>				
Attributes	QName	Type	Fixed	Default	Use
	of	margin.status.of.valueset			required
	on	laterality.valueset			optional
Source	<pre>&lt;xs:element name="status"&gt;   &lt;xs:complexType&gt;     &lt;xs:sequence&gt;       &lt;xs:group ref="generic.content"/&gt;       &lt;xs:group ref="margins.margin.status.result"/&gt;       &lt;xs:group ref="distance"/&gt;     &lt;/xs:sequence&gt;     &lt;xs:attribute name="of" type="margin.status.of.valueset" use="required"/&gt;     &lt;xs:attribute name="on" type="laterality.valueset"/&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt;</pre>				

### Element margins.margin.status.result / result

Namespace	http://purl.org/obbr/pathology/cancer/
-----------	----------------------------------------

Diagram					
Properties	content: complex				
Model	response , specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'				
Children	comment, label, response, specify				
Instance	<pre>&lt;result for=""&gt;   &lt;response closest="" extent="" value=""&gt;{1,1}&lt;/response&gt; &lt;/result&gt;</pre>				
Attributes	QName	Type	Fixed	Default	Use
	for	restriction of xs:token			required
Source	<pre>&lt;xs:element name="result"&gt;   &lt;xs:complexType&gt;     &lt;xs:sequence&gt;       &lt;xs:group ref="margins.margin.status.result.content"/&gt;       &lt;xs:group ref="generic.content"/&gt;     &lt;/xs:sequence&gt;     &lt;xs:attribute name="for" use="required"&gt;       &lt;xs:simpleType&gt;         &lt;xs:restriction base="xs:token"&gt;           &lt;xs:enumeration value="in-situ"/&gt;           &lt;xs:enumeration value="invasive"/&gt;           &lt;xs:enumeration value="specify"/&gt;           &lt;xs:enumeration value="inapplicable"/&gt;           &lt;xs:enumeration value="unreported"/&gt;         &lt;/xs:restriction&gt;       &lt;/xs:simpleType&gt;     &lt;/xs:attribute&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt;</pre>				

### Element margins.margin.status.result.content / response

Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram	<pre>graph TD     response(response) --- attributes_group[attributes]     response --- generic_content_group[generic content]     attributes_group --- value[value]     attributes_group --- extent[extent]     attributes_group --- closest[closest]     generic_content_group --- specify(specify)     generic_content_group --- label(label)     generic_content_group --- comment(comment)     generic_content_group --- other[0..∞ ##other]</pre>				
Properties	content:	complex			
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'				
Children	comment, label, specify				
Instance	<pre>&lt;response closest="" extent="" value=""&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/response&gt;</pre>				
Attributes	QName	Type	Fixed	Default	Use
	closest	union of(val:base.values, val:qualitative.values)			optional
	extent	focality.valueset			optional
	value	restriction of xs:token			required
Source	<pre>&lt;xs:element name="response"&gt;   &lt;xs:complexType&gt;     &lt;xs:group ref="generic.content"/&gt;     &lt;xs:attribute name="value" use="required"&gt;       &lt;xs:simpleType&gt;         &lt;xs:restriction base="xs:token"&gt;           &lt;xs:enumeration value="specify"/&gt;           &lt;xs:enumeration value="inapplicable"/&gt;           &lt;xs:enumeration value="unreported"/&gt;           &lt;xs:enumeration value="positive"/&gt;           &lt;xs:enumeration value="negative"/&gt;           &lt;xs:enumeration value="equivocal"/&gt;           &lt;xs:enumeration value="indeterminate"/&gt;         &lt;/xs:restriction&gt;       &lt;/xs:simpleType&gt;     &lt;/xs:attribute&gt;     &lt;xs:attribute name="extent" type="focality.valueset"/&gt;     &lt;xs:attribute name="closest"&gt;       &lt;xs:simpleType&gt;         &lt;xs:union memberTypes="val:base.values val:qualitative.values"/&gt;       &lt;/xs:simpleType&gt;     &lt;/xs:attribute&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt;</pre>				

## Element distance / distance

Namespace	http://purl.org/obbr/pathology/cancer/
-----------	----------------------------------------

Diagram					
Properties	content:	complex			
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , response				
Children	comment, label, response, specify				
Instance	<pre>&lt;distance for=" " from=" "&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/distance&gt;</pre>				
Attributes	QName	Type	Fixed	Default	Use
	for	restriction of xs:token			required
	from	restriction of xs:token			optional
Source	<pre>&lt;xs:element name="distance"&gt;   &lt;xs:complexType&gt;     &lt;xs:sequence&gt;       &lt;xs:group ref="generic.content"/&gt;       &lt;xs:group ref="length"/&gt;     &lt;/xs:sequence&gt;     &lt;xs:attribute name="for" use="required"&gt;       &lt;xs:simpleType&gt;         &lt;xs:restriction base="xs:token"&gt;           &lt;xs:enumeration value="in-situ"/&gt;           &lt;xs:enumeration value="invasive"/&gt;         &lt;/xs:restriction&gt;       &lt;/xs:simpleType&gt;     &lt;/xs:attribute&gt;     &lt;xs:attribute name="from"&gt;       &lt;xs:simpleType&gt;         &lt;xs:restriction base="xs:token"&gt;           &lt;xs:enumeration value="margin"/&gt;         &lt;/xs:restriction&gt;       &lt;/xs:simpleType&gt;     &lt;/xs:attribute&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt;</pre>				

## Element nodes

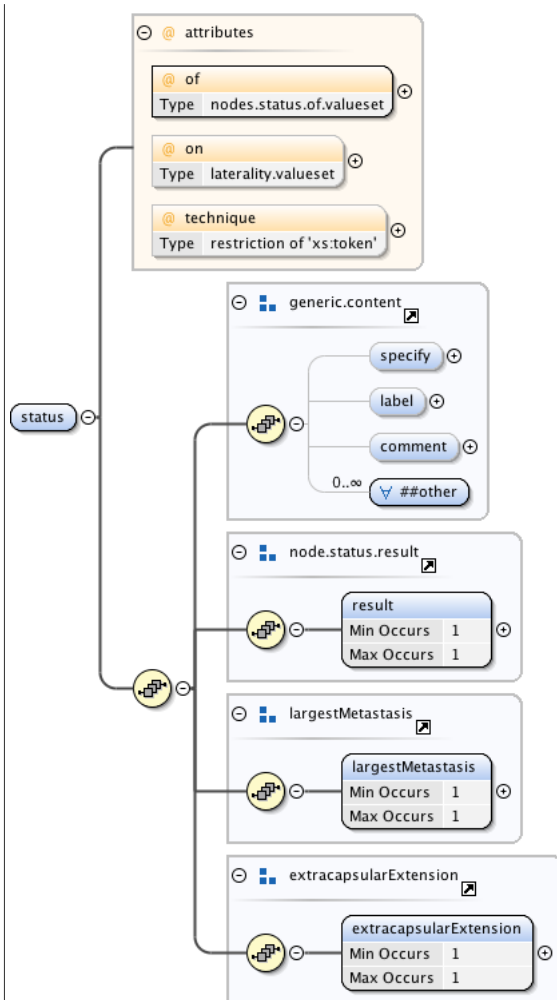
Namespace	http://purl.org/obbr/pathology/cancer/
-----------	----------------------------------------

Diagram					
Properties	content:	complex			
Used by	Element Group	diagnosis.content			
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , status				
Children	comment, label, specify, status				
Instance	<pre>&lt;nodes id=""&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/nodes&gt;</pre>				
Attributes	QName	Type	Fixed	Default	Use
	ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'				
	id	xs:ID			optional
Source	<pre>&lt;xs:element name="nodes"&gt;   &lt;xs:complexType&gt;     &lt;xs:group ref="nodes.content"/&gt;     &lt;xs:attributeGroup ref="generic.attrib"/&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt;</pre>				

## Element nodesStatus / status

Namespace	http://purl.org/obbr/pathology/cancer/
-----------	----------------------------------------

## Diagram



## Properties

content: complex

## Model

specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , result , largestMetastasis , extracapsularExtension

## Children

comment, extracapsularExtension, label, largestMetastasis, result, specify

## Instance

```

<status of="" on="" technique="">
  <specify>{0,1}</specify>
  <label>{0,1}</label>
  <comment>{0,1}</comment>
</status>

```

## Attributes

QName	Type	Fixed	Default	Use
<b>of</b>	nodes.status.of.valueset			required
<b>on</b>	laterality.valueset			optional
<b>technique</b>	restriction of xs:token			optional

## Source

```

<xs:element name="status">
  <xs:complexType>
    <xs:sequence>
      <xs:group ref="generic.content"/>
      <xs:group ref="node.status.result"/>
      <xs:group ref="largestMetastasis"/>
      <xs:group ref="extracapsularExtension"/>
    </xs:sequence>
    <xs:attribute name="of" type="nodes.status.of.valueset" use="required"/>
    <xs:attribute name="on" type="laterality.valueset"/>
    <xs:attribute name="technique">
      <xs:simpleType>
        <xs:restriction base="xs:token">
          <xs:enumeration value="H+E one level"/>
          <xs:enumeration value="H+E multiple level(s)"/>
          <xs:enumeration value="H+E plus immunohistochemistry"/>
          <xs:enumeration value="specify"/>
          <xs:enumeration value="inapplicable"/>
        </xs:restriction>
      </xs:simpleType>
    </xs:attribute>
  </xs:complexType>
</xs:element>

```

```

        <xs:enumeration value="unreported" />
      </xs:restriction>
    </xs:simpleType>
  </xs:attribute>
</xs:complexType>
</xs:element>

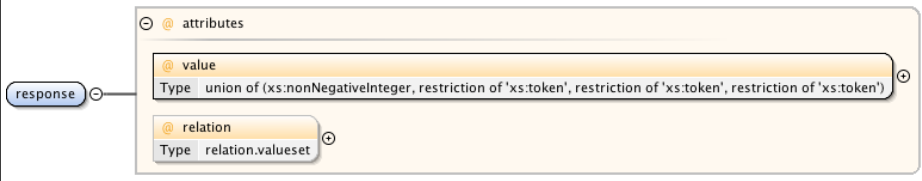
```

## Element node.status.result / result

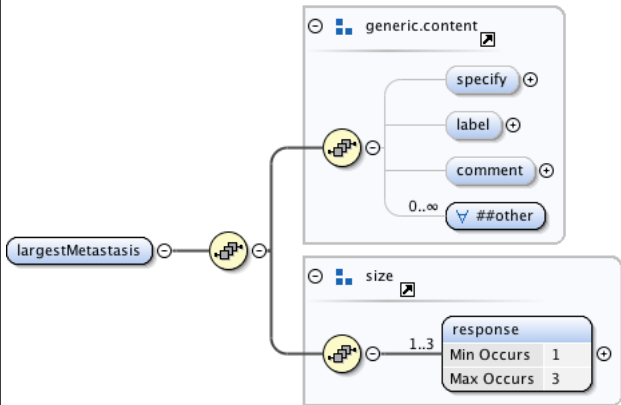
Namespace	http://purl.org/obbr/pathology/cancer/				
Diagram					
Properties	content:	complex			
Model	response , specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'				
Children	comment, label, response, specify				
Instance	<pre>&lt;result for="" type=""&gt;   &lt;response relation="" value=""&gt;{1,1}&lt;/response&gt; &lt;/result&gt;</pre>				
Attributes	QName	Type	Fixed	Default	Use
	for	union of(restriction of xs:token)			required
	type	restriction of xs:token			optional
Source	<pre>&lt;xs:element name="result"&gt;   &lt;xs:complexType&gt;     &lt;xs:sequence&gt;       &lt;xs:group ref="node.status.result.response" /&gt;       &lt;xs:group ref="generic.content" /&gt;     &lt;/xs:sequence&gt;     &lt;xs:attribute name="for" use="required"&gt;       &lt;xs:simpleType&gt;         &lt;xs:union&gt;           &lt;xs:simpleType&gt;             &lt;xs:restriction base="xs:token"&gt;               &lt;xs:enumeration value="total" /&gt;               &lt;xs:enumeration value="negative" /&gt;               &lt;xs:enumeration value="positive" /&gt;             &lt;/xs:restriction&gt;           &lt;/xs:simpleType&gt;         &lt;/xs:union&gt;       &lt;/xs:simpleType&gt;     &lt;/xs:attribute&gt;     &lt;xs:attribute name="type"&gt;       &lt;xs:simpleType&gt;         &lt;xs:restriction base="xs:token"&gt;           &lt;xs:enumeration value="necroinflammatory" /&gt;         &lt;/xs:restriction&gt;       &lt;/xs:simpleType&gt;     &lt;/xs:attribute&gt;   &lt;/xs:complexType&gt;</pre>				

&lt;/xs:element&gt;

**Element node.status.result.response / response**

Namespace	http://purl.org/obbr/pathology/cancer/				
Diagram					
Properties	content:	complex			
Attributes	<b>QName</b>	<b>Type</b>	<b>Fixed</b>	<b>Default</b>	<b>Use</b>
	<b>relation</b>	relation.valueset			optional
	<b>value</b>	union of(xs:nonNegativeInteger, restriction of xs:token, restriction of xs:token, restriction of xs:token)			required
Source	<pre> &lt;xs:element name="response"&gt;   &lt;xs:complexType&gt;     &lt;xs:attribute name="value" use="required"&gt;       &lt;xs:simpleType&gt;         &lt;xs:union memberTypes="xs:nonNegativeInteger"&gt;           &lt;xs:simpleType&gt;             &lt;xs:restriction base="xs:token"&gt;               &lt;xs:enumeration value="specify"/&gt;             &lt;/xs:restriction&gt;           &lt;/xs:simpleType&gt;           &lt;xs:simpleType&gt;             &lt;xs:restriction base="xs:token"&gt;               &lt;xs:enumeration value="inapplicable"/&gt;             &lt;/xs:restriction&gt;           &lt;/xs:simpleType&gt;           &lt;xs:simpleType&gt;             &lt;xs:restriction base="xs:token"&gt;               &lt;xs:enumeration value="unreported"/&gt;             &lt;/xs:restriction&gt;           &lt;/xs:simpleType&gt;         &lt;/xs:union&gt;       &lt;/xs:simpleType&gt;     &lt;/xs:attribute&gt;     &lt;xs:attribute name="relation" type="relation.valueset"/&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt; </pre>				

**Element largestMetastasis / largestMetastasis**

Namespace	http://purl.org/obbr/pathology/cancer/				
Diagram					
Properties	content:	complex			



Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , response{1,3}
Children	comment, label, response, specify
Instance	<pre> &lt;largestMetastasis&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/largestMetastasis&gt; </pre>
Source	<pre> &lt;xs:element name="largestMetastasis"&gt;   &lt;xs:complexType&gt;     &lt;xs:sequence&gt;       &lt;xs:group ref="generic.content" /&gt;       &lt;xs:group ref="size" /&gt;     &lt;/xs:sequence&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt; </pre>

## Element extracapsularExtension / extracapsularExtension

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Properties	content: complex
Model	response , specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'
Children	comment, label, response, specify
Instance	<pre> &lt;extracapsularExtension&gt;   &lt;response value=""&gt;{1,1}&lt;/response&gt; &lt;/extracapsularExtension&gt; </pre>
Source	<pre> &lt;xs:element name="extracapsularExtension"&gt;   &lt;xs:complexType&gt;     &lt;xs:sequence&gt;       &lt;xs:group ref="nodes.extracapsularExtension.content" /&gt;       &lt;xs:group ref="generic.content" /&gt;     &lt;/xs:sequence&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt; </pre>

## Element nodes.extracapsularExtension.content / response

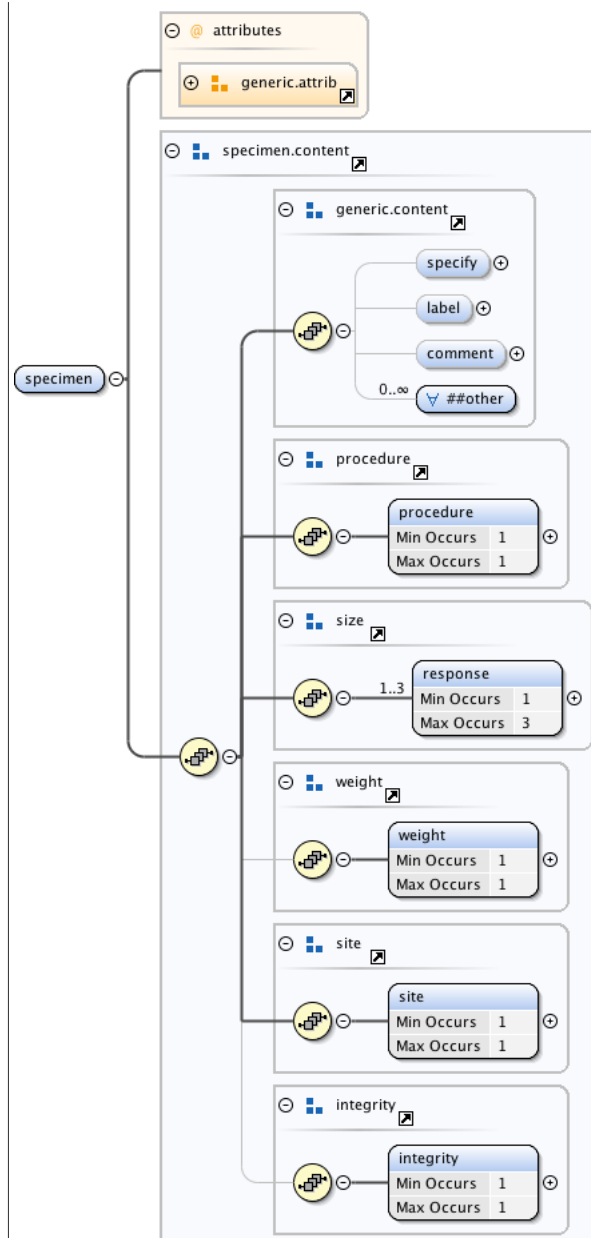
Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	

Properties	content: complex				
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'				
Children	comment, label, specify				
Instance	<pre>&lt;response value=""&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/response&gt;</pre>				
Attributes	QName	Type	Fixed	Default	Use
	value	union of(val:base.values, val:qualitative.values)			required
Source	<pre>&lt;xs:element name="response"&gt;   &lt;xs:complexType&gt;     &lt;xs:group ref="generic.content"/&gt;     &lt;xs:attribute name="value" use="required"&gt;       &lt;xs:simpleType&gt;         &lt;xs:union memberTypes="val:base.values val:qualitative.values"/&gt;       &lt;/xs:simpleType&gt;     &lt;/xs:attribute&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt;</pre>				

## Element specimen

Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram



Properties	content:                      complex				
Used by	Element Group                      diagnosis.content				
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , procedure , response{1,3} , weight , site , integrity				
Children	comment, integrity, label, procedure, response, site, specify, weight				
Instance	<specimen id=""> <specify>{0,1}</specify> <label>{0,1}</label> <comment>{0,1}</comment> </specimen>				
Attributes	QName	Type	Fixed	Default	Use
	ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'				
	id	xs:ID			optional

Source	<pre> &lt;xs:element name="specimen"&gt;   &lt;xs:complexType&gt;     &lt;xs:group ref="specimen.content"/&gt;     &lt;xs:attributeGroup ref="generic.attrib"/&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt; </pre>
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## Element procedure / procedure

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Properties	content: complex
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , response+
Children	comment, label, response, specify
Instance	<pre> &lt;procedure&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/procedure&gt; </pre>
Source	<pre> &lt;xs:element name="procedure"&gt;   &lt;xs:complexType&gt;     &lt;xs:sequence&gt;       &lt;xs:group ref="generic.content"/&gt;       &lt;xs:group ref="specimen.procedure.content"/&gt;     &lt;/xs:sequence&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt; </pre>

## Element specimen.procedure.content / response

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Properties	content: complex maxOccurs: unbounded

Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'				
Children	comment, label, specify				
Instance	<pre>&lt;response on="" value=""&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/response&gt;</pre>				
Attributes	QName	Type	Fixed	Default	Use
	on	laterality.valueset			optional
	value	specimen.procedure.response.valueset			required
Source	<pre>&lt;xs:element maxOccurs="unbounded" name="response"&gt;   &lt;xs:complexType&gt;     &lt;xs:group ref="generic.content"/&gt;     &lt;xs:attribute name="value" type="specimen.procedure.response.valueset" use="required"/&gt;   &gt;   &lt;xs:attribute name="on" type="laterality.valueset"/&gt; &lt;/xs:complexType&gt; &lt;/xs:element&gt;</pre>				

## Element weight / weight

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	<p>The diagram illustrates the structure of the <code>weight</code> element. It is a sequence of two complex content types: <code>generic.content</code> and <code>specimen.weight.content</code>. The <code>generic.content</code> type contains four optional elements: <code>specify</code>, <code>label</code>, <code>comment</code>, and <code>##other</code> (which can occur 0 to infinity times). The <code>specimen.weight.content</code> type contains a <code>response</code> element that must occur exactly once (Min Occurs 1, Max Occurs 1).</p>
Properties	content: complex
Model	<code>specify{0,1}</code> , <code>label{0,1}</code> , <code>comment{0,1}</code> , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , <code>response</code>
Children	<code>comment</code> , <code>label</code> , <code>response</code> , <code>specify</code>
Instance	<pre>&lt;weight&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/weight&gt;</pre>
Source	<pre>&lt;xs:element name="weight"&gt;   &lt;xs:complexType&gt;     &lt;xs:sequence&gt;       &lt;xs:group ref="generic.content"/&gt;       &lt;xs:group ref="specimen.weight.content"/&gt;     &lt;/xs:sequence&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt;</pre>

## Element specimen.weight.content / response

Namespace	http://purl.org/obbr/pathology/cancer/
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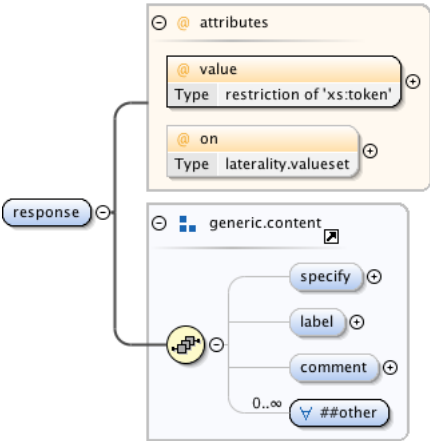
Diagram					
Properties	content:	complex			
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'				
Children	comment, label, specify				
Instance	<pre>&lt;response id="" relation="" unit="" value=""&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/response&gt;</pre>				
Attributes	QName	Type	Fixed	Default	Use
	ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'				
	id	xs:ID			optional
	relation	relation.valueset			optional
	unit	weightUnit.valueset			required
	value	weight.valueset			required
Source	<pre>&lt;xs:element name="response"&gt;   &lt;xs:complexType&gt;     &lt;xs:group ref="generic.content"/&gt;     &lt;xs:attributeGroup ref="weight"/&gt;     &lt;xs:attributeGroup ref="generic.attrib"/&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt;</pre>				

## Element site / site

Namespace	http://purl.org/obbr/pathology/cancer/				
Diagram					
Properties	content:	complex			

Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , response
Children	comment, label, response, specify
Instance	<pre> &lt;site&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/site&gt; </pre>
Source	<pre> &lt;xs:element name="site"&gt;   &lt;xs:complexType&gt;     &lt;xs:sequence&gt;       &lt;xs:group ref="generic.content" /&gt;       &lt;xs:group ref="site.content" /&gt;     &lt;/xs:sequence&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt; </pre>

## Element site.content / response

Namespace	http://purl.org/obbr/pathology/cancer/				
Diagram					
Properties	content:	complex			
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'				
Children	comment, label, specify				
Instance	<pre>&lt;response on="" value=""&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/response&gt;</pre>				
Attributes	QName	Type	Fixed	Default	Use
	on	lateralities.valueset			optional
	value	restriction of xs:token			required
Source	<pre>&lt;xs:element name="response"&gt;   &lt;xs:complexType&gt;     &lt;xs:group ref="generic.content"/&gt;     &lt;xs:attribute name="value" use="required"&gt;       &lt;xs:simpleType&gt;         &lt;xs:restriction base="xs:token"&gt;           &lt;xs:enumeration value="specify"/&gt;           &lt;xs:enumeration value="inapplicable"/&gt;           &lt;xs:enumeration value="unreported"/&gt;         &lt;/xs:restriction&gt;       &lt;/xs:simpleType&gt;     &lt;/xs:attribute&gt;     &lt;xs:attribute name="on" type="lateralities.valueset"/&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt;</pre>				

## Element integrity / integrity

Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram	
Properties	content: complex
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , response
Children	comment, label, response, specify
Instance	<pre>&lt;integrity&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/integrity&gt;</pre>
Source	<pre>&lt;xs:element name="integrity"&gt;   &lt;xs:complexType&gt;     &lt;xs:sequence&gt;       &lt;xs:group ref="generic.content"/&gt;       &lt;xs:group ref="specimen.integrity.content"/&gt;     &lt;/xs:sequence&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt;</pre>

### Element specimen.integrity.content / response

Namespace	http://purl.org/obbr/pathology/cancer/				
Diagram					
Properties	content: complex				
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'				
Children	comment, label, specify				
Instance	<pre>&lt;response value=""&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/response&gt;</pre>				
Attributes	QName	Type	Fixed	Default	Use
	value	restriction of xs:token			required
Source	<pre>&lt;xs:element name="response"&gt;   &lt;xs:complexType&gt;     &lt;xs:group ref="generic.content"/&gt;     &lt;xs:attribute name="value" use="required"&gt;</pre>				



```

<xs:simpleType>
  <xs:restriction base="xs:token">
    <xs:enumeration value="specify"/>
    <xs:enumeration value="inapplicable"/>
    <xs:enumeration value="unreported"/>
    <xs:enumeration value="intact"/>
    <xs:enumeration value="fragmented"/>
  </xs:restriction>
</xs:simpleType>
</xs:attribute>
</xs:complexType>
</xs:element>

```

## Element stage

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Properties	content: complex

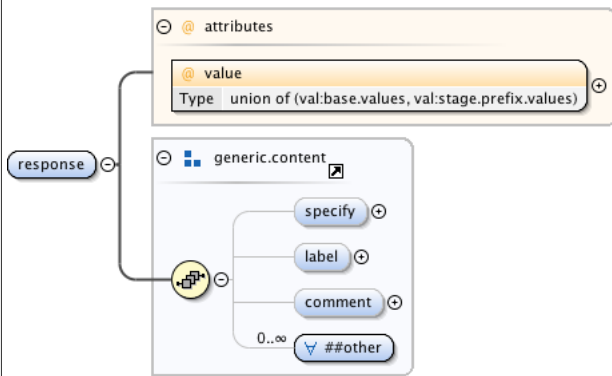
Used by	Element Group diagnosis.content				
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , prefix , multiplicity , pT , pN , pM , group				
Children	comment, group, label, multiplicity, pM, pN, pT, prefix, specify				
Instance	<pre>&lt;stage id=""&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/stage&gt;</pre>				
Attributes	QName	Type	Fixed	Default	Use
	ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'				
	id	xs:ID			optional
Source	<pre>&lt;xs:element name="stage"&gt;   &lt;xs:complexType&gt;     &lt;xs:group ref="stage.content"/&gt;     &lt;xs:attributeGroup ref="generic.attrib"/&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt;</pre>				

### Element prefix / prefix

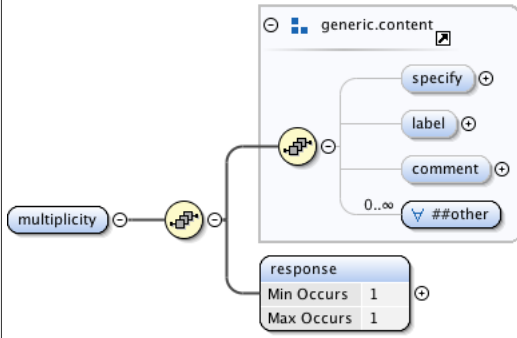
Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Properties	content: complex
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , response
Children	comment, label, response, specify
Instance	<pre>&lt;prefix&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/prefix&gt;</pre>
Source	<pre>&lt;xs:element name="prefix"&gt;   &lt;xs:complexType&gt;     &lt;xs:sequence&gt;       &lt;xs:group ref="generic.content"/&gt;       &lt;xs:group ref="stage.prefix.response"/&gt;     &lt;/xs:sequence&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt;</pre>

### Element stage.prefix.response / response

Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram					
Properties	content:	complex			
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'				
Children	comment, label, specify				
Instance	<pre>&lt;response value=""&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/response&gt;</pre>				
Attributes	QName	Type	Fixed	Default	Use
	value	union of(val:base.values, val:stage.prefix.values)			required
Source	<pre>&lt;xs:element name="response"&gt;   &lt;xs:complexType&gt;     &lt;xs:group ref="generic.content"/&gt;     &lt;xs:attribute name="value" use="required"&gt;       &lt;xs:simpleType&gt;         &lt;xs:union memberTypes="val:base.values val:stage.prefix.values"/&gt;       &lt;/xs:simpleType&gt;     &lt;/xs:attribute&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt;</pre>				

## Element multiplicity / multiplicity

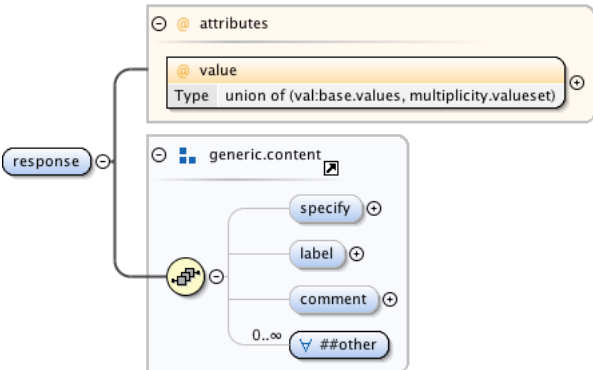
Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Properties	content: complex
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , response
Children	comment, label, response, specify
Instance	<pre>&lt;multiplicity&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt;   &lt;response value=""&gt;{1,1}&lt;/response&gt; &lt;/multiplicity&gt;</pre>
Source	<pre>&lt;xs:element name="multiplicity"&gt;   &lt;xs:complexType&gt;</pre>

```

<xs:sequence>
  <xs:group ref="generic.content"/>
  <xs:element name="response">
    <xs:complexType>
      <xs:group ref="generic.content"/>
      <xs:attribute name="value" use="required">
        <xs:simpleType>
          <xs:union memberTypes="val:base.values multiplicity.valueset"/>
        </xs:simpleType>
      </xs:attribute>
    </xs:complexType>
  </xs:element>
</xs:sequence>
</xs:complexType>
</xs:element>

```

## Element multiplicity / multiplicity / response

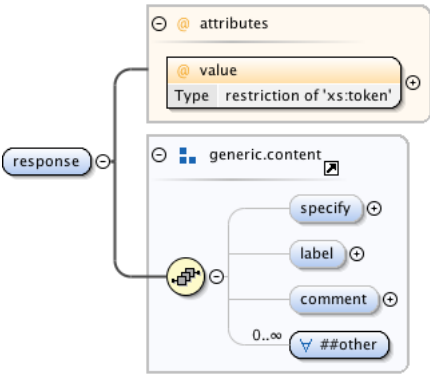
Namespace	http://purl.org/obbr/pathology/cancer/				
Diagram					
Properties	content:	complex			
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'				
Children	comment, label, specify				
Instance	<pre>&lt;response value=""&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/response&gt;</pre>				
Attributes	QName	Type	Fixed	Default	Use
	value	union of(val:base.values, multiplicity.valueset)			required
Source	<pre>&lt;xs:element name="response"&gt;   &lt;xs:complexType&gt;     &lt;xs:group ref="generic.content"/&gt;     &lt;xs:attribute name="value" use="required"&gt;       &lt;xs:simpleType&gt;         &lt;xs:union memberTypes="val:base.values multiplicity.valueset"/&gt;       &lt;/xs:simpleType&gt;     &lt;/xs:attribute&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt;</pre>				

## Element pT / pT

Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram	
Properties	content: complex
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , response
Children	comment, label, response, specify
Instance	<pre>&lt;pT&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/pT&gt;</pre>
Source	<pre>&lt;xs:element name="pT"&gt;   &lt;xs:complexType&gt;     &lt;xs:sequence&gt;       &lt;xs:group ref="generic.content"/&gt;       &lt;xs:group ref="stage.pT.content"/&gt;     &lt;/xs:sequence&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt;</pre>

### Element stage.pT.content / response

Namespace	http://purl.org/obbr/pathology/cancer/				
Diagram					
Properties	content:	complex			
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'				
Children	comment, label, specify				
Instance	<pre>&lt;response value=""&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/response&gt;</pre>				
Attributes	QName	Type	Fixed	Default	Use
	value	restriction of xs:token			required
Source	<pre>&lt;xs:element name="response"&gt;   &lt;xs:complexType&gt;     &lt;xs:group ref="generic.content"/&gt;     &lt;xs:attribute name="value" use="required"&gt;</pre>				

```

<xs:simpleType>
  <xs:restriction base="xs:token">
    <xs:enumeration value="specify"/>
    <xs:enumeration value="inapplicable"/>
    <xs:enumeration value="unreported"/>
  </xs:restriction>
</xs:simpleType>
</xs:attribute>
</xs:complexType>
</xs:element>

```

**Element pN / pN**

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Properties	content: complex
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , response
Children	comment, label, response, specify
Instance	<pre> &lt;pN&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/pN&gt; </pre>
Source	<pre> &lt;xs:element name="pN"&gt;   &lt;xs:complexType&gt;     &lt;xs:sequence&gt;       &lt;xs:group ref="generic.content"/&gt;       &lt;xs:group ref="stage.pN.content"/&gt;     &lt;/xs:sequence&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt; </pre>

**Element stage.pN.content / response**

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Properties	content: complex
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'

Children	comment, label, specify				
Instance	<pre>&lt;response value=" "&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/response&gt;</pre>				
Attributes	QName	Type	Fixed	Default	Use
	value	restriction of xs:token			required
Source	<pre>&lt;xs:element name="response"&gt;   &lt;xs:complexType&gt;     &lt;xs:group ref="generic.content" /&gt;     &lt;xs:attribute name="value" use="required"&gt;       &lt;xs:simpleType&gt;         &lt;xs:restriction base="xs:token"&gt;           &lt;xs:enumeration value="specify" /&gt;           &lt;xs:enumeration value="inapplicable" /&gt;           &lt;xs:enumeration value="unreported" /&gt;         &lt;/xs:restriction&gt;       &lt;/xs:simpleType&gt;     &lt;/xs:attribute&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt;</pre>				

## Element pM / pM

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	<p>The diagram illustrates the structure of the <code>pM</code> element. It is a sequence of two groups: <code>generic.content</code> and <code>stage.pM.content</code>. The <code>generic.content</code> group contains three optional elements: <code>specify</code>, <code>label</code>, and <code>comment</code>, followed by a sequence of zero or more <code>##other</code> elements. The <code>stage.pM.content</code> group contains a <code>response</code> element with a minimum and maximum occurrence of 1.</p>
Properties	content: complex
Model	<code>specify{0,1}</code> , <code>label{0,1}</code> , <code>comment{0,1}</code> , ANY element from ANY namespace OTHER than "http://purl.org/obbr/pathology/cancer/" , <code>response</code>
Children	<code>comment</code> , <code>label</code> , <code>response</code> , <code>specify</code>
Instance	<pre>&lt;pM&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/pM&gt;</pre>
Source	<pre>&lt;xs:element name="pM"&gt;   &lt;xs:complexType&gt;     &lt;xs:sequence&gt;       &lt;xs:group ref="generic.content"/&gt;       &lt;xs:group ref="stage.pM.content"/&gt;     &lt;/xs:sequence&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt;</pre>

## Element stage.pM.content / response

Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram					
Properties	content:	complex			
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'				
Children	comment, label, specify				
Instance	<pre>&lt;response value=""&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/response&gt;</pre>				
Attributes	QName	Type	Fixed	Default	Use
	value	restriction of xs:token			required
Source	<pre>&lt;xs:element name="response"&gt;   &lt;xs:complexType&gt;     &lt;xs:group ref="generic.content"/&gt;     &lt;xs:attribute name="value" use="required"&gt;       &lt;xs:simpleType&gt;         &lt;xs:restriction base="xs:token"&gt;           &lt;xs:enumeration value="specify"/&gt;           &lt;xs:enumeration value="inapplicable"/&gt;           &lt;xs:enumeration value="unreported"/&gt;         &lt;/xs:restriction&gt;       &lt;/xs:simpleType&gt;     &lt;/xs:attribute&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt;</pre>				

### Element stage.group / group

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Properties	content: complex
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , response
Children	comment, label, response, specify
Instance	<pre>&lt;group&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt;</pre>



	</group>
Source	<pre> &lt;xs:element name="group"&gt;   &lt;xs:complexType&gt;     &lt;xs:sequence&gt;       &lt;xs:group ref="generic.content" /&gt;       &lt;xs:group ref="stage.group.response" /&gt;     &lt;/xs:sequence&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt; </pre>

### Element stage.group.response / response

Namespace	http://purl.org/obbr/pathology/cancer/				
Diagram					
Properties	content:	complex			
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'				
Children	comment, label, specify				
Instance	<pre>&lt;response value=""&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/response&gt;</pre>				
Attributes	QName	Type	Fixed	Default	Use
	value	stage.group.valueset			required
Source	<pre>&lt;xs:element name="response"&gt;   &lt;xs:complexType&gt;     &lt;xs:group ref="generic.content"/&gt;     &lt;xs:attribute name="value" type="stage.group.valueset" use="required"/&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt;</pre>				

### Element studies

Namespace	http://purl.org/obbr/pathology/cancer/				
Diagram					

Properties	content: complex				
Used by	Element Group diagnosis.content				
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than "http://purl.org/obbr/pathology/cancer/" , study				
Children	comment, label, specify, study				
Instance	<pre>&lt;studies id=""&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt;   &lt;study&gt;{1,1}&lt;/study&gt; &lt;/studies&gt;</pre>				
Attributes	QName	Type	Fixed	Default	Use
	ANY attribute from ANY namespace OTHER than "http://purl.org/obbr/pathology/cancer/"				
	id	xs:ID			optional
Source	<pre>&lt;xs:element name="studies"&gt;   &lt;xs:complexType&gt;     &lt;xs:group ref="studies.content"/&gt;     &lt;xs:attributeGroup ref="generic.attrib"/&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt;</pre>				

### Element studies.content / study

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Properties	content: complex
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' ,
Children	comment, label, specify
Instance	<pre> &lt;study&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/study&gt; </pre>
Source	<pre> &lt;xs:element name="study"&gt;   &lt;xs:complexType&gt;     &lt;xs:sequence&gt;       &lt;xs:group ref="generic.content"/&gt;       &lt;xs:group ref="studies.study.content"/&gt;     &lt;/xs:sequence&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt; </pre>

### Element tumor

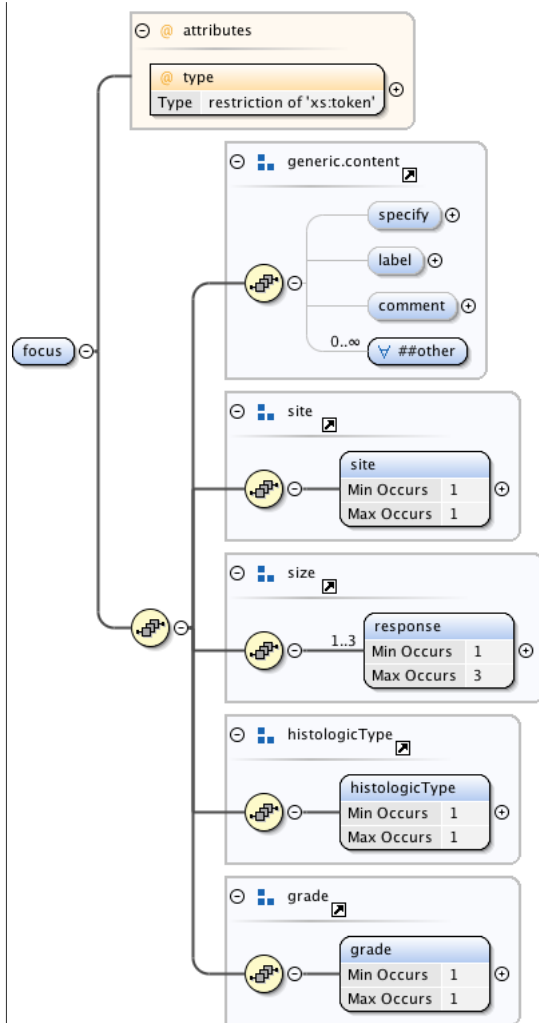
Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram	<pre>graph TD     tumor((tumor)) --- choice1(( ))     choice1 --- generic_attr[generic.attrib]     choice1 --- choice2(( ))     choice2 --- generic_content[generic.content]     choice2 --- focus_element[focus]     generic_content --- choice3(( ))     choice3 --- specify(specify)     choice3 --- label(label)     choice3 --- comment(comment)     choice3 --- other[##other 0..∞]     focus_element --- focus_type[focus Min Occurs 1 Max Occurs 1]</pre>				
Properties	content:	complex			
Used by	Element Group	diagnosis.content			
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , focus				
Children	comment, focus, label, specify				
Instance	<pre>&lt;tumor id=""&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/tumor&gt;</pre>				
Attributes	QName	Type	Fixed	Default	Use
	ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'				
	id	xs:ID			optional
Source	<pre>&lt;xs:element name="tumor"&gt;   &lt;xs:complexType&gt;     &lt;xs:group ref="tumor.content"/&gt;     &lt;xs:attributeGroup ref="generic.attrib"/&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt;</pre>				

## Element focus / focus

Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram



Properties	content: complex				
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , site , response{1,3} , histologicType , grade				
Children	comment, grade, histologicType, label, response, site, specify				
Instance	<pre> &lt;focus type=""&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/focus&gt; </pre>				
Attributes	QName	Type	Fixed	Default	Use
	type	restriction of xs:token			required
Source	<pre> &lt;xs:element name="focus"&gt;   &lt;xs:complexType&gt;     &lt;xs:sequence&gt;       &lt;xs:group ref="generic.content" /&gt;       &lt;xs:group ref="site" /&gt;       &lt;xs:group ref="size" /&gt;       &lt;xs:group ref="histologicType" /&gt;       &lt;xs:group ref="grade" /&gt;     &lt;/xs:sequence&gt;     &lt;xs:attribute name="type" use="required"&gt;       &lt;xs:simpleType&gt;         &lt;xs:restriction base="xs:token"&gt;           &lt;xs:enumeration value="principal"/&gt;           &lt;xs:enumeration value="sole"/&gt;           &lt;xs:enumeration value="additional"/&gt;         &lt;/xs:restriction&gt;       &lt;/xs:simpleType&gt;     &lt;/xs:attribute&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt; </pre>				

**Element histologicType / histologicType**

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Properties	content: complex
Model	response , specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'
Children	comment, label, response, specify
Instance	<pre>&lt;histologicType&gt;   &lt;response value="" {1,1}&lt;/response&gt; &lt;/histologicType&gt;</pre>
Source	<pre>&lt;xs:element name="histologicType"&gt;   &lt;xs:complexType&gt;     &lt;xs:sequence&gt;       &lt;xs:group ref="tumor.histologicType.content"/&gt;       &lt;xs:group ref="generic.content"/&gt;     &lt;/xs:sequence&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt;</pre>

**Element tumor.histologicType.content / response**

Namespace	http://purl.org/obbr/pathology/cancer/				
Diagram					
Properties	content:	complex			
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'				
Children	comment, label, specify				
Instance	<pre>&lt;response value=""&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/response&gt;</pre>				
Attributes	QName	Type	Fixed	Default	Use
	value	union of(val:base.values, tumor.histologicType.valueset)			required

Source	<pre> &lt;xs:element name="response"&gt;   &lt;xs:complexType&gt;     &lt;xs:group ref="generic.content"/&gt;     &lt;xs:attribute name="value" use="required"&gt;       &lt;xs:simpleType&gt;         &lt;xs:union memberTypes="val:base.values tumor.histologicType.valueset"/&gt;       &lt;/xs:simpleType&gt;     &lt;/xs:attribute&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt; </pre>
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## Element grade / grade

Namespace	http://purl.org/obbr/pathology/cancer/				
Diagram					
Properties	content:	complex			
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , response				
Children	comment, label, response, specify				
Instance	<pre>&lt;grade system=" "&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/grade&gt;</pre>				
Attributes	QName	Type	Fixed	Default	Use
	system	tumor.gradeSystem.valueset			required
Source	<pre>&lt;xs:element name="grade"&gt;   &lt;xs:complexType&gt;     &lt;xs:sequence&gt;       &lt;xs:group ref="generic.content"/&gt;       &lt;xs:group ref="tumor.grade.response"/&gt;     &lt;/xs:sequence&gt;     &lt;xs:attribute name="system" type="tumor.gradeSystem.valueset" use="required"/&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt;</pre>				

## Element tumor.grade.response / response

Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram					
Properties	content:	complex			
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'				
Children	comment, label, specify				
Instance	<pre>&lt;response value=""&gt;   &lt;specify&gt;{0,1}&lt;/specify&gt;   &lt;label&gt;{0,1}&lt;/label&gt;   &lt;comment&gt;{0,1}&lt;/comment&gt; &lt;/response&gt;</pre>				
Attributes	QName	Type	Fixed	Default	Use
	value	union of(val:base.values, tumor.grade.valueset)			required
Source	<pre>&lt;xs:element name="response"&gt;   &lt;xs:complexType&gt;     &lt;xs:group ref="generic.content"/&gt;     &lt;xs:attribute name="value" use="required"&gt;       &lt;xs:simpleType&gt;         &lt;xs:union memberTypes="val:base.values tumor.grade.valueset"/&gt;       &lt;/xs:simpleType&gt;     &lt;/xs:attribute&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt;</pre>				

## Simple Types

### Simple Type lengthUnit.valueset


Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	<p>The diagram illustrates a derivation union. On the left, a box labeled 'lengthUnit.valueset' contains the text 'Derivation union'. An arrow points from this box to a box on the right labeled 'val:lengthUnit.values'. Both boxes have a small circle with a minus sign at their top-right corners.</p>
Type	union of(val:lengthUnit.values)
Used by	Attribute                      dimensionalQuantity/@unit
Source	<pre>&lt;xs:simpleType name="lengthUnit.valueset"&gt;   &lt;xs:union memberTypes="val:lengthUnit.values"/&gt; &lt;/xs:simpleType&gt;</pre>

### Simple Type date.valueset

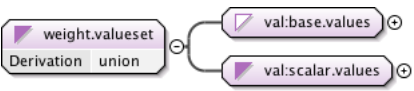
Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	<p>The diagram shows a central box labeled 'date.valueset' with a 'Derivation' tab set to 'union'. To its right, a circle with a minus sign connects to three boxes: 'xs:date', 'xs:gYearMonth', and 'xs:gYear'. Each box has a small icon in the top-left corner.</p>
Type	union of(xs:date, xs:gYearMonth, xs:gYear)
Used by	Attributes date/@date, date/@end, date/@start
Source	<pre>&lt;xs:simpleType name="date.valueset"&gt;   &lt;xs:union memberTypes="xs:date xs:gYearMonth xs:gYear" /&gt; &lt;/xs:simpleType&gt;</pre>

```
</xs:simpleType>
```


### Simple Type weightUnit.valueset

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(val:weightUnit.values)
Used by	Attribute weight/@unit
Source	<pre>&lt;xs:simpleType name="weightUnit.valueset"&gt;   &lt;xs:union memberTypes="val:weightUnit.values" /&gt; &lt;/xs:simpleType&gt;</pre>


### Simple Type weight.valueset

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(val:base.values, val:scalar.values)
Used by	Attribute weight/@value
Source	<pre>&lt;xs:simpleType name="weight.valueset"&gt;   &lt;xs:union memberTypes="val:base.values val:scalar.values" /&gt; &lt;/xs:simpleType&gt;</pre>

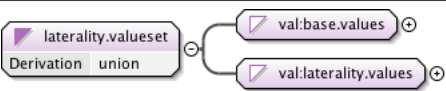
### Simple Type relation.valueset

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(val:relation.values)
Used by	Attributes dimensionalQuantity/@relation, node.status.result.response/response/@relation, weight/@relation
Source	<pre>&lt;xs:simpleType name="relation.valueset"&gt;   &lt;xs:union memberTypes="val:relation.values" /&gt; &lt;/xs:simpleType&gt;</pre>

### Simple Type multiplicity.valueset

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(val:multiplicity.values)
Source	<pre>&lt;xs:simpleType name="multiplicity.valueset"&gt;   &lt;xs:union memberTypes="val:multiplicity.values" /&gt; &lt;/xs:simpleType&gt;</pre>


### Simple Type laterality.valueset

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(val:base.values, val:laterality.values)
Used by	Attributes contiguousExtension.status.result/result/@on, distant.status/status/@on, generic.content/@on, margins.margin.status/status/@on, nodesStatus/status/@on, site.content/response/@on, specimen.procedure.content/response/@on
Source	<pre>&lt;xs:simpleType name="laterality.valueset"&gt;   &lt;xs:union memberTypes="val:base.values val:laterality.values" /&gt; &lt;/xs:simpleType&gt;</pre>

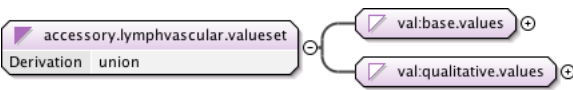


```
</xs:simpleType>
```

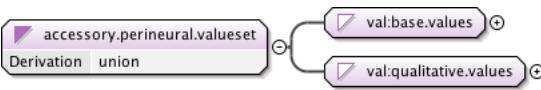
### Simple Type **focality.valueset**

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(val:focality.values)
Used by	Attributes contiguousExtension.status.result.content/response/@extent, generic.content/@extent, margins.margin.status.result.content/response/@extent
Source	<pre>&lt;xs:simpleType name="focality.valueset"&gt;   &lt;xs:union memberTypes="val:focality.values" /&gt; &lt;/xs:simpleType&gt;</pre>


### Simple Type **accessory.lymphvascular.valueset**

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(val:base.values, val:qualitative.values)
Used by	Attribute accessory.lymphvascular.content/response/@value
Source	<pre>&lt;xs:simpleType name="accessory.lymphvascular.valueset"&gt;   &lt;xs:union memberTypes="val:base.values val:qualitative.values" /&gt; &lt;/xs:simpleType&gt;</pre>

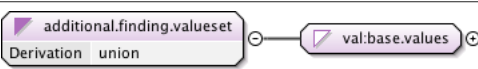
### Simple Type **accessory.perineural.valueset**

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(val:base.values, val:qualitative.values)
Used by	Attribute accessory.perineural.content/response/@value
Source	<pre>&lt;xs:simpleType name="accessory.perineural.valueset"&gt;   &lt;xs:union memberTypes="val:base.values val:qualitative.values" /&gt; &lt;/xs:simpleType&gt;</pre>

### Simple Type **accessory.treatmentResponse.valueset**

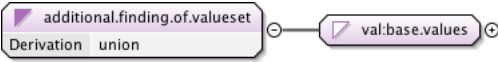
Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(val:accessory.treatmentResponse.values)
Used by	Attribute accessory.treatment.content/response/@value
Source	<pre>&lt;xs:simpleType name="accessory.treatmentResponse.valueset"&gt;   &lt;xs:union memberTypes="val:accessory.treatmentResponse.values" /&gt; &lt;/xs:simpleType&gt;</pre>

### Simple Type **additional.finding.valueset**


Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(val:base.values)
Used by	Attribute additional.finding.content/response/@value
Source	<pre>&lt;xs:simpleType name="additional.finding.valueset"&gt;   &lt;xs:union memberTypes="val:base.values" /&gt; &lt;/xs:simpleType&gt;</pre>

	</xs:simpleType>
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### Simple Type additional.finding.of.valueset

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(val:base.values)
Used by	Attribute additional.content/finding/@of
Source	<pre>&lt;xs:simpleType name="additional.finding.of.valueset"&gt;   &lt;xs:union memberTypes="val:base.values" /&gt; &lt;/xs:simpleType&gt;</pre>


### Simple Type extent.contiguousExtension.status.result.for.valueset

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(val:extent.contiguousExtension.status.result.for.values)
Source	<pre>&lt;xs:simpleType name="extent.contiguousExtension.status.result.for.valueset"&gt;   &lt;xs:union memberTypes="val:extent.contiguousExtension.status.result.for.values" /&gt; &lt;/xs:simpleType&gt;</pre>

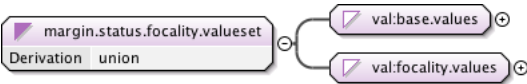
### Simple Type extent.status.of.valueset

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(val:base.values)
Used by	Attribute contiguousExtensionStatus/status/@of
Source	<pre>&lt;xs:simpleType name="extent.status.of.valueset"&gt;   &lt;xs:union memberTypes="val:base.values" /&gt; &lt;/xs:simpleType&gt;</pre>

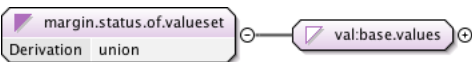
### Simple Type history.priorTherapy.valueset

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(val:history.priorTherapy.values)
Source	<pre>&lt;xs:simpleType name="history.priorTherapy.valueset"&gt;   &lt;xs:union memberTypes="val:history.priorTherapy.values" /&gt; &lt;/xs:simpleType&gt;</pre>

### Simple Type margin.status.focality.valueset

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(val:base.values, val:focality.values)
Source	<pre>&lt;xs:simpleType name="margin.status.focality.valueset"&gt;   &lt;xs:union memberTypes="val:base.values val:focality.values" /&gt; &lt;/xs:simpleType&gt;</pre>

### Simple Type margin.status.of.valueset

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	

Type	union of(val:base.values)
Used by	Attribute margins.margin.status/status/@of
Source	<pre>&lt;xs:simpleType name="margin.status.of.valueset"&gt;   &lt;xs:union memberTypes="val:base.values" /&gt; &lt;/xs:simpleType&gt;</pre>

### Simple Type margin.status.result.for.valueset

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(val:margin.status.result.for.values)
Source	<pre>&lt;xs:simpleType name="margin.status.result.for.valueset"&gt;   &lt;xs:union memberTypes="val:margin.status.result.for.values" /&gt; &lt;/xs:simpleType&gt;</pre>

### Simple Type distant.status.of.valueset

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(val:distant.status.of.values)
Used by	Attribute distant.status/status/@of
Source	<pre>&lt;xs:simpleType name="distant.status.of.valueset"&gt;   &lt;xs:union memberTypes="val:distant.status.of.values" /&gt; &lt;/xs:simpleType&gt;</pre>

### Simple Type distant.status.result.for.valueset

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(val:distant.status.result.for.values)
Used by	Attribute distant.status.result/result/@for
Source	<pre>&lt;xs:simpleType name="distant.status.result.for.valueset"&gt;   &lt;xs:union memberTypes="val:distant.status.result.for.values" /&gt; &lt;/xs:simpleType&gt;</pre>


### Simple Type nodes.status.result.for.valueset

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(val:nodes.status.result.for.values)
Source	<pre>&lt;xs:simpleType name="nodes.status.result.for.valueset"&gt;   &lt;xs:union memberTypes="val:nodes.status.result.for.values" /&gt; &lt;/xs:simpleType&gt;</pre>


### Simple Type nodes.status.result.type.valueset

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(val:base.values, val:nodes.status.result.type.values)
Source	<pre>&lt;xs:simpleType name="nodes.status.result.type.valueset"&gt;   &lt;xs:union memberTypes="val:base.values val:nodes.status.result.type.values" /&gt; &lt;/xs:simpleType&gt;</pre>

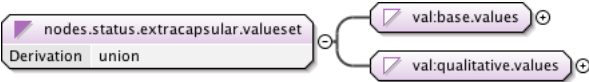
**Simple Type nodes.status.of.valueset**

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(val:base.values)
Used by	Attribute nodesStatus/status/@of
Source	<pre>&lt;xs:simpleType name="nodes.status.of.valueset"&gt;   &lt;xs:union memberTypes="val:base.values" /&gt; &lt;/xs:simpleType&gt;</pre>

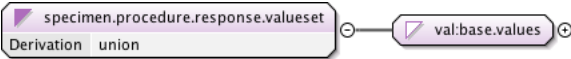
**Simple Type nodes.status.technique.valueset**

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(val:nodes.status.technique.values)
Source	<pre>&lt;xs:simpleType name="nodes.status.technique.valueset"&gt;   &lt;xs:union memberTypes="val:nodes.status.technique.values" /&gt; &lt;/xs:simpleType&gt;</pre>


**Simple Type nodes.status.extracapsular.valueset**

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(val:base.values, val:qualitative.values)
Source	<pre>&lt;xs:simpleType name="nodes.status.extracapsular.valueset"&gt;   &lt;xs:union memberTypes="val:base.values val:qualitative.values" /&gt; &lt;/xs:simpleType&gt;</pre>


**Simple Type specimen.procedure.response.valueset**

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(val:base.values)
Used by	Attribute specimen.procedure.content/response/@value
Source	<pre>&lt;xs:simpleType name="specimen.procedure.response.valueset"&gt;   &lt;xs:union memberTypes="val:base.values" /&gt; &lt;/xs:simpleType&gt;</pre>

**Simple Type specimen.integrity.valueset**

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(val:specimen.integrity.values)
Source	<pre>&lt;xs:simpleType name="specimen.integrity.valueset"&gt;   &lt;xs:union memberTypes="val:specimen.integrity.values" /&gt; &lt;/xs:simpleType&gt;</pre>

**Simple Type stage.prefix.valueset**

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(val:stage.prefix.values)

Source	<pre>&lt;xs:simpleType name="stage.prefix.valueset"&gt;   &lt;xs:union memberTypes="val:stage.prefix.values" /&gt; &lt;/xs:simpleType&gt;</pre>
--------	-------------------------------------------------------------------------------------------------------------------------------------------------

### Simple Type stage.group.valueset

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	<pre> graph LR     A[stage.group.valueset Derivation union] --- B[val:base.values]   </pre>
Type	union of(val:base.values)
Used by	Attribute stage.group.response/response/@value
Source	<pre>&lt;xs:simpleType name="stage.group.valueset"&gt;   &lt;xs:union memberTypes="val:base.values" /&gt; &lt;/xs:simpleType&gt;</pre>

### Simple Type tumor.histologicType.valueset

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	<pre> graph LR     A[tumor.histologicType.valueset Derivation union] --- B[val:base.values]   </pre>
Type	union of(val:base.values)
Source	<pre>&lt;xs:simpleType name="tumor.histologicType.valueset"&gt;   &lt;xs:union memberTypes="val:base.values" /&gt; &lt;/xs:simpleType&gt;</pre>

### Simple Type tumor.grade.valueset

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	<pre> graph LR     A[tumor.grade.valueset Derivation union] --- B[val:tumor.grade.values]   </pre>
Type	union of(val:tumor.grade.values)
Source	<pre>&lt;xs:simpleType name="tumor.grade.valueset"&gt;   &lt;xs:union memberTypes="val:tumor.grade.values" /&gt; &lt;/xs:simpleType&gt;</pre>

### Simple Type tumor.gradeSystem.valueset

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	<pre> graph LR     A[tumor.gradeSystem.valueset Derivation union] --- B[val:tumor.gradeSystem.values]   </pre>
Type	union of(val:tumor.gradeSystem.values)
Used by	Attribute grade/grade/@system
Source	<pre>&lt;xs:simpleType name="tumor.gradeSystem.valueset"&gt;   &lt;xs:union memberTypes="val:tumor.gradeSystem.values" /&gt; &lt;/xs:simpleType&gt;</pre>

### Simple Type positiveDecimal

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	<pre> graph LR     A[positiveDecimal Derivation restriction Base Type xs:decimal] --- B[xs:decimal]   </pre>
Type	restriction of xs:decimal
Facets	minInclusive 0.0
Used by	Attribute dimensionalQuantity/@value
Source	<pre>&lt;xs:simpleType name="positiveDecimal"&gt;   &lt;xs:restriction base="xs:decimal"&gt;     &lt;xs:minInclusive value="0.0" /&gt;   &lt;/xs:restriction&gt; &lt;/xs:simpleType&gt;</pre>

## Element Groups

### Element Group size

Namespace	http://purl.org/obbr/pathology/cancer/	
Diagram		
Used by	Elements	distant.status/status, focus/focus, largestMetastasis/largestMetastasis
	Element Group	specimen.content
Model	response{1,3}	
Children	response	
Source	<pre> &lt;xs:group name="size"&gt;   &lt;xs:sequence&gt;     &lt;xs:element maxOccurs="3" minOccurs="1" name="response"&gt;       &lt;xs:complexType&gt;         &lt;xs:group ref="generic.content" /&gt;         &lt;xs:attributeGroup ref="dimensionalQuantity" /&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>	

### Element Group generic.content

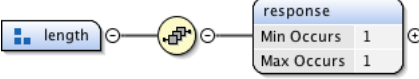
Namespace	http://purl.org/obbr/pathology/cancer/	
Diagram		
Used by	Elements	accessory, accessory.lymphvascular.content/response, accessory.lymphvascular/lymphvascularInvasion, accessory.perineural.content/response, accessory.perineural/perineuralInvasion, accessory.treatment.content/response, accessory.treatment/treatmentResponse, additional, additional.content/finding, additional.finding.content/response, clinicalFinding.content/response, clinicalFinding/clinicalFinding, contiguousExtension.status.distance/distance, contiguousExtension.status.result.content/response, contiguousExtension.status.result/result, contiguousExtension/contiguousExtension, contiguousExtensionStatus/status, diagnosis, distance/distance, distant, distant.status.result.response/response, distant.status.result/result, distant.status/status, extent, extracapsularExtension/extracapsularExtension, focus/focus, grade/grade, histologicType/histologicType, integrity/integrity, largestMetastasis/largestMetastasis, length/response, margins.margin.status.result.content/response, margins.margin.status.result/result, margins.margin.status/status, multiplicity/multiplicity, multiplicity/multiplicity/response, node.status.result/result, nodes.extracapsularExtension.content/response, nodesStatus/status, pM/pM, pN/pN, pT/pT, pathologicFinding.content/response, pathologicFinding/pathologicFinding, prefix/prefix, priorTherapy.content/response, priorTherapy/priorTherapy, procedure/procedure, radiologicFinding.content/response, radiologicFinding/radiologicFinding, site.content/response, site/site, size/response, specimen.integrity.content/response, specimen.procedure.content/response, specimen.weight.content/response, stage.group.response/response, stage.group/group, stage.pM.content/response, stage.pN.content/response, stage.pT.content/response, stage.prefix.response/response, studies.content/study, tumor.grade.response/response, tumor.histologicType.content/response, weight/weight
	Element Groups	history.content, margin.content, nodes.content, specimen.content, stage.content, studies.content, tumor.content
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'	
Children	comment, label, specify	
Source	<pre> &lt;xs:group name="generic.content"&gt;   &lt;xs:sequence&gt;     &lt;xs:element minOccurs="0" name="specify" type="xs:string" /&gt;     &lt;xs:element minOccurs="0" name="label" type="xs:string" /&gt;     &lt;xs:element minOccurs="0" name="comment" type="xs:string" /&gt; </pre>	

```

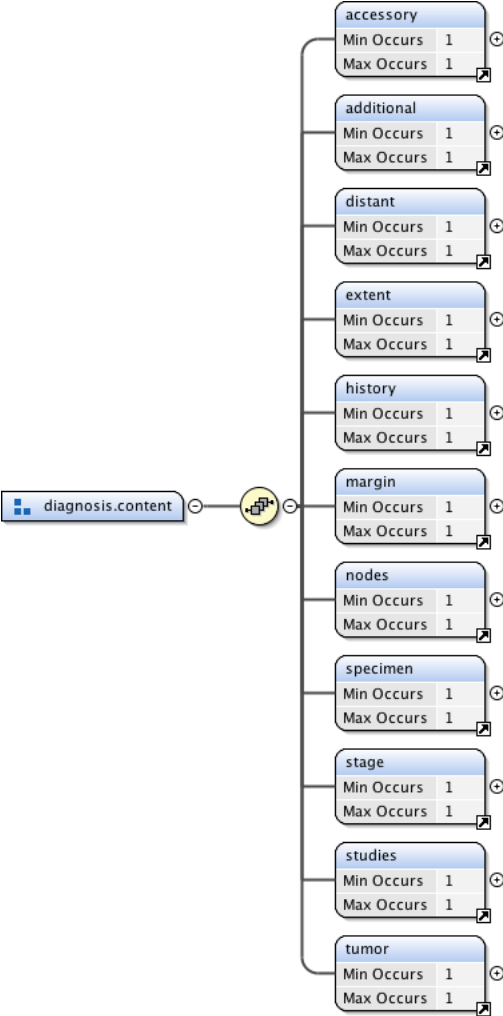
<xs:any maxOccurs="unbounded" minOccurs="0" namespace="##other"
processContents="skip" />
</xs:sequence>
</xs:group>

```

## Element Group length

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Elements contiguousExtension.status.distance/distance, distance/distance
Model	response
Children	response
Source	<pre> &lt;xs:group name="length"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="response"&gt;       &lt;xs:complexType&gt;         &lt;xs:group ref="generic.content" /&gt;         &lt;xs:attributeGroup ref="dimensionalQuantity" /&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>

## Element Group diagnosis.content

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element diagnosis

Model	accessory , additional , distant , extent , history , margin , nodes , specimen , stage , studies , tumor
Children	accessory, additional, distant, extent, history, margin, nodes, specimen, stage, studies, tumor
Source	<pre> &lt;xs:group name="diagnosis.content"&gt;   &lt;xs:sequence&gt;     &lt;xs:element ref="accessory"/&gt;     &lt;xs:element ref="additional"/&gt;     &lt;xs:element ref="distant"/&gt;     &lt;xs:element ref="extent"/&gt;     &lt;xs:element ref="history"/&gt;     &lt;xs:element ref="margin"/&gt;     &lt;xs:element ref="nodes"/&gt;     &lt;xs:element ref="specimen"/&gt;     &lt;xs:element ref="stage"/&gt;     &lt;xs:element ref="studies"/&gt;     &lt;xs:element ref="tumor"/&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>

### Element Group accessory.content

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element      accessory
Model	treatmentResponse , lymphvascularInvasion , perineuralInvasion
Children	lymphvascularInvasion, perineuralInvasion, treatmentResponse
Source	<pre> &lt;xs:group name="accessory.content"&gt;   &lt;xs:sequence&gt;     &lt;xs:group minOccurs="0" ref="accessory.treatment"/&gt;     &lt;xs:group minOccurs="0" ref="accessory.lymphvascular"/&gt;     &lt;xs:group minOccurs="0" ref="accessory.perineural"/&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>

### Element Group accessory.treatment

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element Group      accessory.content
Model	treatmentResponse
Children	treatmentResponse
Source	<pre> &lt;xs:group name="accessory.treatment"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="treatmentResponse"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt;           &lt;xs:group ref="generic.content"/&gt;           &lt;xs:group ref="accessory.treatment.content"/&gt;         &lt;/xs:sequence&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt;   &lt;xs:attributeGroup ref="generic.attrib"/&gt; &lt;/xs:group&gt; </pre>




```

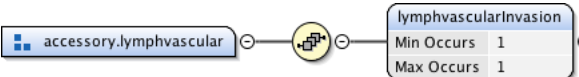
</xs:complexType>
</xs:element>
</xs:sequence>
</xs:group>

```

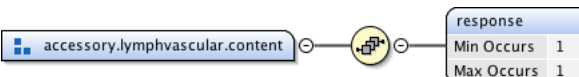
### Element Group accessory.treatment.content

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element accessory.treatment/treatmentResponse
Model	response
Children	response
Source	<pre> &lt;xs:group name="accessory.treatment.content"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="response"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt;           &lt;xs:group ref="generic.content" /&gt;         &lt;/xs:sequence&gt;         &lt;xs:attribute name="value" type="accessory.treatmentResponse.valueset"           use="required" /&gt;         &lt;xs:attributeGroup ref="generic.attrib" /&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>

### Element Group accessory.lymphvascular

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element Group accessory.content
Model	lymphvascularInvasion
Children	lymphvascularInvasion
Source	<pre> &lt;xs:group name="accessory.lymphvascular"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="lymphvascularInvasion"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt;           &lt;xs:group ref="generic.content" /&gt;           &lt;xs:group ref="accessory.lymphvascular.content" /&gt;         &lt;/xs:sequence&gt;         &lt;xs:attributeGroup ref="generic.attrib" /&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>

### Element Group accessory.lymphvascular.content

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element accessory.lymphvascular/lymphvascularInvasion
Model	response
Children	response
Source	<pre> &lt;xs:group name="accessory.lymphvascular.content"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="response"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt;           &lt;xs:group ref="generic.content" /&gt;         &lt;/xs:sequence&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>

```

</xs:sequence>
<xs:attribute name="value" type="accessory.lymphvascular.valueset" use="required"/
>
<xs:attributeGroup ref="generic.attrib"/>
</xs:complexType>
</xs:element>
</xs:sequence>
</xs:group>

```

### Element Group accessory.perineural

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element Group      accessory.content
Model	perineuralInvasion
Children	perineuralInvasion
Source	<pre> &lt;xs:group name="accessory.perineural"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="perineuralInvasion"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt;           &lt;xs:group ref="generic.content"/&gt;           &lt;xs:group ref="accessory.perineural.content"/&gt;         &lt;/xs:sequence&gt;         &lt;xs:attributeGroup ref="generic.attrib"/&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>

### Element Group accessory.perineural.content

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element      accessory.perineural/perineuralInvasion
Model	response
Children	response
Source	<pre> &lt;xs:group name="accessory.perineural.content"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="response"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt;           &lt;xs:group ref="generic.content"/&gt;         &lt;/xs:sequence&gt;         &lt;xs:attribute name="value" type="accessory.perineural.valueset" use="required"/&gt;         &lt;xs:attributeGroup ref="generic.attrib"/&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>

### Element Group additional.content

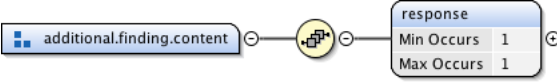
Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element      additional
Model	finding*
Children	finding
Source	<pre> &lt;xs:group name="additional.content"&gt;   &lt;xs:sequence&gt;     &lt;xs:element maxOccurs="unbounded" minOccurs="0" name="finding"&gt; </pre>

```

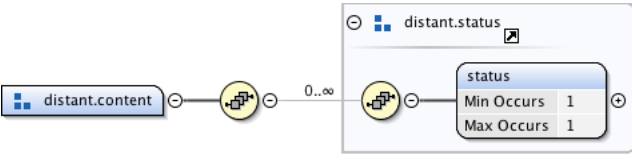
<xs:complexType>
  <xs:sequence>
    <xs:group ref="generic.content"/>
    <xs:group ref="additional.finding.content"/>
  </xs:sequence>
  <xs:attribute name="of" type="additional.finding.of.valueset"/>
  <xs:attributeGroup ref="generic.attrib"/>
</xs:complexType>
</xs:element>
</xs:sequence>
</xs:group>

```

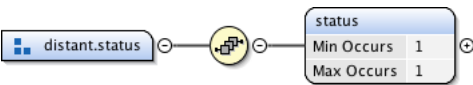
### Element Group additional.finding.content

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element additional.content/finding
Model	response
Children	response
Source	<pre> &lt;xs:group name="additional.finding.content"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="response"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt;           &lt;xs:group ref="generic.content"/&gt;         &lt;/xs:sequence&gt;         &lt;xs:attribute name="value" type="additional.finding.valueset" use="required"/&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>

### Element Group distant.content

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element distant
Model	status
Children	status
Source	<pre> &lt;xs:group name="distant.content"&gt;   &lt;xs:sequence&gt;     &lt;xs:group maxOccurs="unbounded" minOccurs="0" ref="distant.status"/&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>

### Element Group distant.status


Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element Group distant.content
Model	status
Children	status
Source	<pre> &lt;xs:group name="distant.status"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="status"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt;           &lt;xs:group ref="generic.content"/&gt;         &lt;/xs:sequence&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>

```

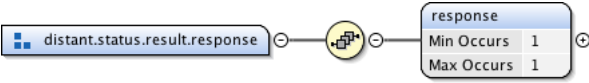
<xs:group ref="distant.status.result"/>
<xs:group minOccurs="0" ref="size"/>
</xs:sequence>
<xs:attribute name="of" type="distant.status.of.valueset" use="required"/>
<xs:attribute name="on" type="laterality.valueset"/>
</xs:complexType>
</xs:element>
</xs:sequence>
</xs:group>

```

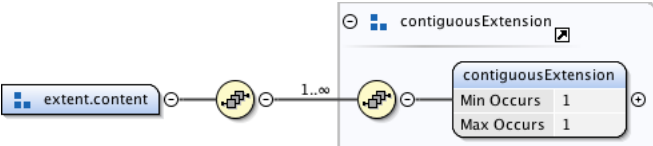
## Element Group distant.status.result

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element distant.status/status
Model	result
Children	result
Source	<pre> &lt;xs:group name="distant.status.result"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="result"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt;           &lt;xs:group ref="generic.content"/&gt;           &lt;xs:group ref="distant.status.result.response"/&gt;         &lt;/xs:sequence&gt;           &lt;xs:attribute name="for" type="distant.status.result.for.valueset"/&gt;           &lt;xs:attributeGroup ref="date"/&gt;         &lt;/xs:complexType&gt;       &lt;/xs:element&gt;     &lt;/xs:sequence&gt;   &lt;/xs:group&gt; </pre>

## Element Group distant.status.result.response

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element distant.status.result/result
Model	response
Children	response
Source	<pre> &lt;xs:group name="distant.status.result.response"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="response"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt;           &lt;xs:group ref="generic.content"/&gt;         &lt;/xs:sequence&gt;           &lt;xs:attribute name="value" use="required"&gt;             &lt;xs:simpleType&gt;               &lt;xs:union memberTypes="val:base.values val:qualitative.values"/&gt;             &lt;/xs:simpleType&gt;           &lt;/xs:attribute&gt;         &lt;/xs:complexType&gt;       &lt;/xs:element&gt;     &lt;/xs:sequence&gt;   &lt;/xs:group&gt; </pre>

## Element Group extent.content

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	

Used by	Element extent
Model	contiguousExtension
Children	contiguousExtension
Source	<pre>&lt;xs:group name="extent.content"&gt;   &lt;xs:sequence&gt;     &lt;xs:group maxOccurs="unbounded" ref="contiguousExtension"/&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt;</pre>

## Element Group contiguousExtension

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element Group extent.content
Model	contiguousExtension
Children	contiguousExtension
Source	<pre>&lt;xs:group name="contiguousExtension"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="contiguousExtension"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt;           &lt;xs:group ref="generic.content"/&gt;           &lt;xs:group maxOccurs="unbounded" minOccurs="0" ref="contiguousExtensionStatus"/&gt;         &lt;/xs:sequence&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt;</pre>

## Element Group contiguousExtensionStatus

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element contiguousExtension/contiguousExtension
Model	status
Children	status
Source	<pre>&lt;xs:group name="contiguousExtensionStatus"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="status"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence maxOccurs="unbounded"&gt;           &lt;xs:group ref="generic.content"/&gt;           &lt;xs:group ref="contiguousExtension.status.result"/&gt;           &lt;xs:group minOccurs="0" ref="contiguousExtension.status.distance"/&gt;         &lt;/xs:sequence&gt;         &lt;xs:attribute name="of" type="extent.status.of.valueset"/&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt;</pre>

## Element Group contiguousExtension.status.result

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element contiguousExtensionStatus/status
Model	result
Children	result

Source	<pre> &lt;xs:group name="contiguousExtension.status.result"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="result"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt;           &lt;xs:group ref="generic.content" /&gt;           &lt;xs:group maxOccurs="unbounded" minOccurs="0" ref="contiguousExtension.status.result.content" /&gt;         &lt;/xs:sequence&gt;         &lt;xs:attribute name="for" use="required"&gt;           &lt;xs:simpleType&gt;             &lt;xs:union memberTypes="val:base.values"&gt;               &lt;xs:simpleType&gt;                 &lt;xs:restriction base="xs:token"&gt;                   &lt;xs:enumeration value="invades" /&gt;                   &lt;xs:enumeration value="adheres" /&gt;                   &lt;xs:enumeration value="specify" /&gt;                   &lt;xs:enumeration value="inapplicable" /&gt;                   &lt;xs:enumeration value="unreported" /&gt;                 &lt;/xs:restriction&gt;               &lt;/xs:simpleType&gt;             &lt;/xs:union&gt;           &lt;/xs:simpleType&gt;         &lt;/xs:attribute&gt;         &lt;xs:attribute name="on" type="laterality.valueset" /&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>
--------	------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------

### Element Group contiguousExtension.status.result.content

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element contiguousExtension.status.result/result
Model	response
Children	response
Source	<pre> &lt;xs:group name="contiguousExtension.status.result.content"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="response"&gt;       &lt;xs:complexType&gt;         &lt;xs:group ref="generic.content" /&gt;         &lt;xs:attribute name="value" use="required"&gt;           &lt;xs:simpleType&gt;             &lt;xs:union memberTypes="val:base.values val:qualitative.values" /&gt;           &lt;/xs:simpleType&gt;         &lt;/xs:attribute&gt;         &lt;xs:attribute name="extent" type="focality.valueset" /&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>

### Element Group contiguousExtension.status.distance

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element contiguousExtensionStatus/status
Model	distance
Children	distance
Source	<pre> &lt;xs:group name="contiguousExtension.status.distance"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="distance"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt;           &lt;xs:group ref="generic.content" /&gt;           &lt;xs:group ref="length" /&gt;         &lt;/xs:sequence&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>

```

</xs:sequence>
<xs:attribute name="for">
  <xs:simpleType>
    <xs:restriction base="xs:token">
      <xs:enumeration value="invades"/>
      <xs:enumeration value="adheres"/>
    </xs:restriction>
  </xs:simpleType>
</xs:attribute>
<xs:attribute name="from" type="val:base.values"/>
</xs:complexType>
</xs:element>
</xs:sequence>
</xs:group>

```

## Element Group history.content

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	<pre> graph LR     historyContent[history.content] --- choice1(( ))     choice1 --- genericContent[generic.content]     choice1 --- choice2(( ))     choice2 --- clinicalFinding[clinicalFinding]     choice2 --- pathologicFinding[pathologicFinding]     choice2 --- priorTherapy[priorTherapy]     choice2 --- radiologicFinding[radiologicFinding]     genericContent --- specify[specify]     genericContent --- label[label]     genericContent --- comment[comment]     genericContent --- other[##other]     clinicalFinding --- clinicalFindingBox[clinicalFinding Min Occurs 1 Max Occurs 1]     pathologicFinding --- pathologicFindingBox[pathologicFinding Min Occurs 1 Max Occurs 1]     priorTherapy --- priorTherapyBox[priorTherapy Min Occurs 1 Max Occurs 1]     radiologicFinding --- radiologicFindingBox[radiologicFinding Min Occurs 1 Max Occurs 1] </pre>
Used by	Element history
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer' , ((clinicalFinding)   (pathologicFinding)   (priorTherapy)   (radiologicFinding))
Children	clinicalFinding, comment, label, pathologicFinding, priorTherapy, radiologicFinding, specify
Source	<pre> &lt;xs:group name="history.content"&gt;   &lt;xs:sequence&gt;     &lt;xs:group ref="generic.content"/&gt;     &lt;xs:choice maxOccurs="unbounded" minOccurs="0"&gt;       &lt;xs:group ref="clinicalFinding"/&gt;       &lt;xs:group ref="pathologicFinding"/&gt;       &lt;xs:group ref="priorTherapy"/&gt;       &lt;xs:group ref="radiologicFinding"/&gt;     &lt;/xs:choice&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>

## Element Group clinicalFinding

Namespace	http://purl.org/obbr/pathology/cancer/
-----------	----------------------------------------

Diagram	
Used by	Element Group history.content
Model	clinicalFinding
Children	clinicalFinding
Source	<pre> &lt;xs:group name="clinicalFinding"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="clinicalFinding"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt;           &lt;xs:group ref="generic.content" /&gt;           &lt;xs:group ref="clinicalFinding.content" /&gt;         &lt;/xs:sequence&gt;         &lt;xs:attributeGroup ref="date" /&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>

### Element Group clinicalFinding.content

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element clinicalFinding/clinicalFinding
Model	response
Children	response
Source	<pre> &lt;xs:group name="clinicalFinding.content"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="response"&gt;       &lt;xs:complexType&gt;         &lt;xs:group ref="generic.content" /&gt;         &lt;xs:attribute name="value" use="required"&gt;           &lt;xs:simpleType&gt;             &lt;xs:restriction base="xs:token"&gt;               &lt;xs:enumeration value="specify" /&gt;               &lt;xs:enumeration value="inapplicable" /&gt;               &lt;xs:enumeration value="unreported" /&gt;             &lt;/xs:restriction&gt;           &lt;/xs:simpleType&gt;         &lt;/xs:attribute&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>

### Element Group pathologicFinding

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element Group history.content
Model	pathologicFinding
Children	pathologicFinding
Source	<pre> &lt;xs:group name="pathologicFinding"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="pathologicFinding"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt;           &lt;xs:group ref="generic.content" /&gt;           &lt;xs:group ref="pathologicFinding.content" /&gt;         &lt;/xs:sequence&gt;         &lt;xs:attributeGroup ref="date" /&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>



```
</xs:sequence>
</xs:group>
```

### Element Group **pathologicFinding.content**

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element pathologicFinding/pathologicFinding
Model	response
Children	response
Source	<pre>&lt;xs:group name="pathologicFinding.content"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="response"&gt;       &lt;xs:complexType&gt;         &lt;xs:group ref="generic.content"/&gt;         &lt;xs:attribute name="value" use="required"&gt;           &lt;xs:simpleType&gt;             &lt;xs:restriction base="xs:token"&gt;               &lt;xs:enumeration value="specify"/&gt;               &lt;xs:enumeration value="inapplicable"/&gt;               &lt;xs:enumeration value="unreported"/&gt;             &lt;/xs:restriction&gt;           &lt;/xs:simpleType&gt;         &lt;/xs:attribute&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt;</pre>

### Element Group **priorTherapy**

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element Group history.content
Model	priorTherapy
Children	priorTherapy
Source	<pre>&lt;xs:group name="priorTherapy"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="priorTherapy"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt;           &lt;xs:group ref="generic.content"/&gt;           &lt;xs:group ref="priorTherapy.content"/&gt;         &lt;/xs:sequence&gt;         &lt;xs:attributeGroup ref="date"/&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt;</pre>

### Element Group **priorTherapy.content**


Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element priorTherapy/priorTherapy
Model	response
Children	response
Source	<pre>&lt;xs:group name="priorTherapy.content"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="response"&gt;       &lt;xs:complexType&gt;</pre>

```


<xs:group ref="generic.content" />
<xs:attribute name="value" use="required">
  <xs:simpleType>
    <xs:restriction base="xs:token">
      <xs:enumeration value="specify" />
      <xs:enumeration value="inapplicable" />
      <xs:enumeration value="unreported" />
      <xs:enumeration value="chemotherapy" />
      <xs:enumeration value="radiation" />
      <xs:enumeration value="surgery" />
    </xs:restriction>
  </xs:simpleType>
</xs:attribute>
</xs:complexType>
</xs:element>
</xs:sequence>
</xs:group>

```

## Element Group radiologicFinding

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element Group history.content
Model	radiologicFinding
Children	radiologicFinding
Source	<pre> &lt;xs:group name="radiologicFinding"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="radiologicFinding"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt;           &lt;xs:group ref="generic.content" /&gt;           &lt;xs:group ref="radiologicFinding.content" /&gt;         &lt;/xs:sequence&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>

## Element Group radiologicFinding.content

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element radiologicFinding/radiologicFinding
Model	response
Children	response
Source	<pre> &lt;xs:group name="radiologicFinding.content"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="response"&gt;       &lt;xs:complexType&gt;         &lt;xs:group ref="generic.content" /&gt;         &lt;xs:attribute name="value" use="required"&gt;           &lt;xs:simpleType&gt;             &lt;xs:restriction base="xs:token"&gt;               &lt;xs:enumeration value="specify" /&gt;               &lt;xs:enumeration value="inapplicable" /&gt;               &lt;xs:enumeration value="unreported" /&gt;             &lt;/xs:restriction&gt;           &lt;/xs:simpleType&gt;         &lt;/xs:attribute&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>

## Element Group margin.content

Namespace	http://purl.org/obbr/pathology/cancer/
-----------	----------------------------------------

Diagram	
Used by	Element            margin
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , status
Children	comment, label, specify, status
Source	<pre> &lt;xs:group name="margin.content"&gt;   &lt;xs:sequence&gt;     &lt;xs:group ref="generic.content" /&gt;     &lt;xs:group maxOccurs="unbounded" minOccurs="0" ref="margins.margin.status" /&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>

### Element Group margins.margin.status

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element Group    margin.content
Model	status
Children	status
Source	<pre> &lt;xs:group name="margins.margin.status"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="status"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt;           &lt;xs:group ref="generic.content" /&gt;           &lt;xs:group ref="margins.margin.status.result" /&gt;           &lt;xs:group ref="distance" /&gt;         &lt;/xs:sequence&gt;           &lt;xs:attribute name="of" type="margin.status.of.valueset" use="required" /&gt;           &lt;xs:attribute name="on" type="laterality.valueset" /&gt;         &lt;/xs:complexType&gt;       &lt;/xs:element&gt;     &lt;/xs:sequence&gt;   &lt;/xs:group&gt; </pre>

### Element Group margins.margin.status.result

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element            margins.margin.status/status
Model	result
Children	result
Source	<pre> &lt;xs:group name="margins.margin.status.result"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="result"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt; </pre>

```

<xs:group ref="margins.margin.status.result.content"/>
<xs:group ref="generic.content"/>
</xs:sequence>
<xs:attribute name="for" use="required">
  <xs:simpleType>
    <xs:restriction base="xs:token">
      <xs:enumeration value="in-situ"/>
      <xs:enumeration value="invasive"/>
      <xs:enumeration value="specify"/>
      <xs:enumeration value="inapplicable"/>
      <xs:enumeration value="unreported"/>
    </xs:restriction>
  </xs:simpleType>
</xs:attribute>
</xs:complexType>
</xs:element>
</xs:sequence>
</xs:group>

```

## Element Group margins.margin.status.result.content

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element margins.margin.status.result/result
Model	response
Children	response
Source	<pre> &lt;xs:group name="margins.margin.status.result.content"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="response"&gt;       &lt;xs:complexType&gt;         &lt;xs:group ref="generic.content"/&gt;         &lt;xs:attribute name="value" use="required"&gt;           &lt;xs:simpleType&gt;             &lt;xs:restriction base="xs:token"&gt;               &lt;xs:enumeration value="specify"/&gt;               &lt;xs:enumeration value="inapplicable"/&gt;               &lt;xs:enumeration value="unreported"/&gt;               &lt;xs:enumeration value="positive"/&gt;               &lt;xs:enumeration value="negative"/&gt;               &lt;xs:enumeration value="equivocal"/&gt;               &lt;xs:enumeration value="indeterminate"/&gt;             &lt;/xs:restriction&gt;           &lt;/xs:simpleType&gt;         &lt;/xs:attribute&gt;         &lt;xs:attribute name="extent" type="focality.valueset"/&gt;         &lt;xs:attribute name="closest"&gt;           &lt;xs:simpleType&gt;             &lt;xs:union memberTypes="val:base.values val:qualitative.values"/&gt;           &lt;/xs:simpleType&gt;         &lt;/xs:attribute&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>

## Element Group distance

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element margins.margin.status/status
Model	distance
Children	distance
Source	<pre> &lt;xs:group name="distance"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="distance"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt;           &lt;xs:group ref="generic.content"/&gt; </pre>

```

    <xs:group ref="length"/>
  </xs:sequence>
  <xs:attribute name="for" use="required">
    <xs:simpleType>
      <xs:restriction base="xs:token">
        <xs:enumeration value="in-situ"/>
        <xs:enumeration value="invasive"/>
      </xs:restriction>
    </xs:simpleType>
  </xs:attribute>
  <xs:attribute name="from">
    <xs:simpleType>
      <xs:restriction base="xs:token">
        <xs:enumeration value="margin"/>
      </xs:restriction>
    </xs:simpleType>
  </xs:attribute>
</xs:complexType>
</xs:element>
</xs:sequence>
</xs:group>

```

## Element Group nodes.content

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element nodes
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , status
Children	comment, label, specify, status
Source	<pre> &lt;xs:group name="nodes.content"&gt;   &lt;xs:sequence&gt;     &lt;xs:group ref="generic.content"/&gt;     &lt;xs:group maxOccurs="unbounded" minOccurs="0" ref="nodesStatus"/&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>

## Element Group nodesStatus

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element Group nodes.content
Model	status
Children	status
Source	<pre> &lt;xs:group name="nodesStatus"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="status"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt;           &lt;xs:group ref="generic.content"/&gt;           &lt;xs:group ref="node.status.result"/&gt;           &lt;xs:group ref="largestMetastasis"/&gt;           &lt;xs:group ref="extracapsularExtension"/&gt;         &lt;/xs:sequence&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>

```

</xs:sequence>
<xs:attribute name="of" type="nodes.status.of.valueset" use="required"/>
<xs:attribute name="on" type="laterality.valueset"/>
<xs:attribute name="technique">
  <xs:simpleType>
    <xs:restriction base="xs:token">
      <xs:enumeration value="H+E one level"/>
      <xs:enumeration value="H+E multiple level(s)"/>
      <xs:enumeration value="H+E plus immunohistochemistry"/>
      <xs:enumeration value="specify"/>
      <xs:enumeration value="inapplicable"/>
      <xs:enumeration value="unreported"/>
    </xs:restriction>
  </xs:simpleType>
</xs:attribute>
</xs:complexType>
</xs:element>
</xs:sequence>
</xs:group>

```

### Element Group node.status.result

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element nodesStatus/status
Model	result
Children	result
Source	<pre> &lt;xs:group name="node.status.result"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="result"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt;           &lt;xs:group ref="node.status.result.response"/&gt;           &lt;xs:group ref="generic.content"/&gt;         &lt;/xs:sequence&gt;         &lt;xs:attribute name="for" use="required"&gt;           &lt;xs:simpleType&gt;             &lt;xs:union&gt;               &lt;xs:simpleType&gt;                 &lt;xs:restriction base="xs:token"&gt;                   &lt;xs:enumeration value="total"/&gt;                   &lt;xs:enumeration value="negative"/&gt;                   &lt;xs:enumeration value="positive"/&gt;                 &lt;/xs:restriction&gt;               &lt;/xs:simpleType&gt;             &lt;/xs:union&gt;           &lt;/xs:simpleType&gt;         &lt;/xs:attribute&gt;         &lt;xs:attribute name="type"&gt;           &lt;xs:simpleType&gt;             &lt;xs:restriction base="xs:token"&gt;               &lt;xs:enumeration value="necroinflammatory"/&gt;             &lt;/xs:restriction&gt;           &lt;/xs:simpleType&gt;         &lt;/xs:attribute&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>

### Element Group node.status.result.response

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element node.status.result/result
Model	response
Children	response
Source	<pre> &lt;xs:group name="node.status.result.response"&gt; </pre>

```

<xs:sequence>
  <xs:element name="response">
    <xs:complexType>
      <xs:attribute name="value" use="required">
        <xs:simpleType>
          <xs:union memberTypes="xs:nonNegativeInteger">
            <xs:simpleType>
              <xs:restriction base="xs:token">
                <xs:enumeration value="specify"/>
              </xs:restriction>
            </xs:simpleType>
            <xs:simpleType>
              <xs:restriction base="xs:token">
                <xs:enumeration value="inapplicable"/>
              </xs:restriction>
            </xs:simpleType>
            <xs:simpleType>
              <xs:restriction base="xs:token">
                <xs:enumeration value="unreported"/>
              </xs:restriction>
            </xs:simpleType>
          </xs:union>
        </xs:simpleType>
      </xs:attribute>
      <xs:attribute name="relation" type="relation.valueset"/>
    </xs:complexType>
  </xs:element>
</xs:sequence>
</xs:group>

```

### Element Group largestMetastasis

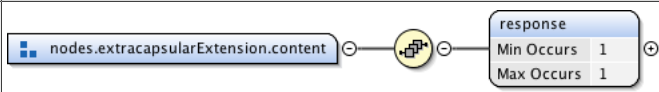
Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element nodesStatus/status
Model	largestMetastasis
Children	largestMetastasis
Source	<pre> &lt;xs:group name="largestMetastasis"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="largestMetastasis"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt;           &lt;xs:group ref="generic.content"/&gt;           &lt;xs:group ref="size"/&gt;         &lt;/xs:sequence&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>

### Element Group extracapsularExtension

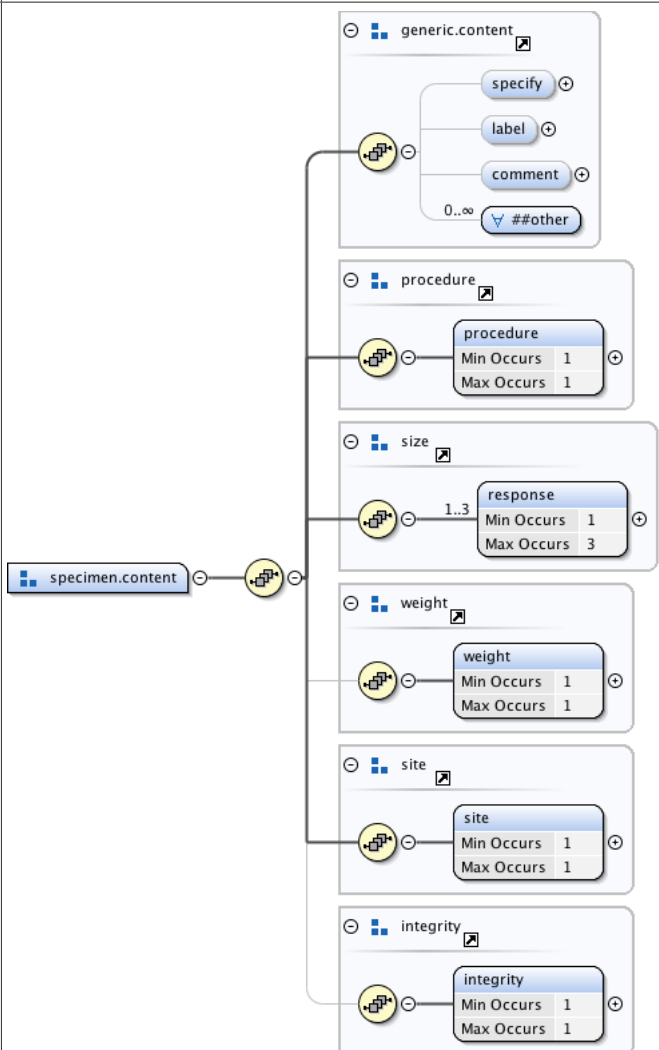
Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element nodesStatus/status
Model	extracapsularExtension
Children	extracapsularExtension
Source	<pre> &lt;xs:group name="extracapsularExtension"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="extracapsularExtension"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt;           &lt;xs:group ref="nodes.extracapsularExtension.content"/&gt;           &lt;xs:group ref="generic.content"/&gt;         &lt;/xs:sequence&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>

&lt;/xs:group&gt;

**Element Group nodes.extracapsularExtension.content**

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element extracapsularExtension/extracapsularExtension
Model	response
Children	response
Source	<pre> &lt;xs:group name="nodes.extracapsularExtension.content"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="response"&gt;       &lt;xs:complexType&gt;         &lt;xs:group ref="generic.content"/&gt;         &lt;xs:attribute name="value" use="required"&gt;           &lt;xs:simpleType&gt;             &lt;xs:union memberTypes="val:base.values val:qualitative.values"/&gt;           &lt;/xs:simpleType&gt;         &lt;/xs:attribute&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>

**Element Group specimen.content**

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	



Used by	Element specimen
Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , procedure , response{1,3} , weight , site , integrity
Children	comment, integrity, label, procedure, response, site, specify, weight
Source	<pre> &lt;xs:group name="specimen.content"&gt;   &lt;xs:sequence&gt;     &lt;xs:group ref="generic.content"/&gt;     &lt;xs:group ref="procedure"/&gt;     &lt;xs:group ref="size"/&gt;     &lt;xs:group minOccurs="0" ref="weight"/&gt;     &lt;xs:group ref="site"/&gt;     &lt;xs:group minOccurs="0" ref="integrity"/&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>

### Element Group procedure

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element Group specimen.content
Model	procedure
Children	procedure
Source	<pre> &lt;xs:group name="procedure"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="procedure"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt;           &lt;xs:group ref="generic.content"/&gt;           &lt;xs:group ref="specimen.procedure.content"/&gt;         &lt;/xs:sequence&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>

### Element Group specimen.procedure.content

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element procedure/procedure
Model	response+
Children	response
Source	<pre> &lt;xs:group name="specimen.procedure.content"&gt;   &lt;xs:sequence&gt;     &lt;xs:element maxOccurs="unbounded" name="response"&gt;       &lt;xs:complexType&gt;         &lt;xs:group ref="generic.content"/&gt;         &lt;xs:attribute name="value" type="specimen.procedure.response.valueset" use="required"/&gt;         &lt;xs:attribute name="on" type="laterality.valueset"/&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>

### Element Group weight

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	

Used by	Element Group specimen.content
Model	weight
Children	weight
Source	<pre> &lt;xs:group name="weight"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="weight"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt;           &lt;xs:group ref="generic.content"/&gt;           &lt;xs:group ref="specimen.weight.content"/&gt;         &lt;/xs:sequence&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>

### Element Group specimen.weight.content

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element weight/weight
Model	response
Children	response
Source	<pre> &lt;xs:group name="specimen.weight.content"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="response"&gt;       &lt;xs:complexType&gt;         &lt;xs:group ref="generic.content"/&gt;         &lt;xs:attributeGroup ref="weight"/&gt;         &lt;xs:attributeGroup ref="generic.attrib"/&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>

### Element Group site

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	<div>Element Group specimen.content</div> <div>Element focus/focus</div>
Model	site
Children	site
Source	<pre> &lt;xs:group name="site"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="site"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt;           &lt;xs:group ref="generic.content"/&gt;           &lt;xs:group ref="site.content"/&gt;         &lt;/xs:sequence&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>

### Element Group site.content

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	

Used by	Element site/site
Model	response
Children	response
Source	<pre> &lt;xs:group name="site.content"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="response"&gt;       &lt;xs:complexType&gt;         &lt;xs:group ref="generic.content" /&gt;         &lt;xs:attribute name="value" use="required"&gt;           &lt;xs:simpleType&gt;             &lt;xs:restriction base="xs:token"&gt;               &lt;xs:enumeration value="specify"/&gt;               &lt;xs:enumeration value="inapplicable"/&gt;               &lt;xs:enumeration value="unreported"/&gt;             &lt;/xs:restriction&gt;           &lt;/xs:simpleType&gt;         &lt;/xs:attribute&gt;         &lt;xs:attribute name="on" type="laterality.valueset"/&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>

### Element Group integrity

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element Group specimen.content
Model	integrity
Children	integrity
Source	<pre> &lt;xs:group name="integrity"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="integrity"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt;           &lt;xs:group ref="generic.content" /&gt;           &lt;xs:group ref="specimen.integrity.content" /&gt;         &lt;/xs:sequence&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>

### Element Group specimen.integrity.content

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element integrity/integrity
Model	response
Children	response
Source	<pre> &lt;xs:group name="specimen.integrity.content"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="response"&gt;       &lt;xs:complexType&gt;         &lt;xs:group ref="generic.content" /&gt;         &lt;xs:attribute name="value" use="required"&gt;           &lt;xs:simpleType&gt;             &lt;xs:restriction base="xs:token"&gt;               &lt;xs:enumeration value="specify"/&gt;               &lt;xs:enumeration value="inapplicable"/&gt;               &lt;xs:enumeration value="unreported"/&gt;               &lt;xs:enumeration value="intact"/&gt;               &lt;xs:enumeration value="fragmented"/&gt;             &lt;/xs:restriction&gt;           &lt;/xs:simpleType&gt;         &lt;/xs:attribute&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>

```

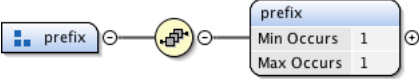
</xs:attribute>
</xs:complexType>
</xs:element>
</xs:sequence>
</xs:group>

```


## Element Group `stage.content`

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	<p>The diagram illustrates the structure of the <code>stage.content</code> element group. It is a sequence of the following components:</p> <ul style="list-style-type: none"> <li><b>generic.content</b>: Contains <code>specify</code> (0..1), <code>label</code> (0..1), <code>comment</code> (0..1), and <code>##other</code> (0..∞).</li> <li><b>prefix</b>: Contains <code>prefix</code> (Min Occurs: 1, Max Occurs: 1).</li> <li><b>multiplicity</b>: Contains <code>multiplicity</code> (Min Occurs: 1, Max Occurs: 1).</li> <li><b>pT</b>: Contains <code>pT</code> (Min Occurs: 1, Max Occurs: 1).</li> <li><b>pN</b>: Contains <code>pN</code> (Min Occurs: 1, Max Occurs: 1).</li> <li><b>pM</b>: Contains <code>pM</code> (Min Occurs: 1, Max Occurs: 1).</li> <li><b>stage.group</b>: Contains <code>group</code> (Min Occurs: 1, Max Occurs: 1).</li> </ul>
Used by	Element <code>stage</code>
Model	<code>specify{0,1}</code> , <code>label{0,1}</code> , <code>comment{0,1}</code> , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , <code>prefix</code> , <code>multiplicity</code> , <code>pT</code> , <code>pN</code> , <code>pM</code> , <code>group</code>
Children	<code>comment</code> , <code>group</code> , <code>label</code> , <code>multiplicity</code> , <code>pM</code> , <code>pN</code> , <code>pT</code> , <code>prefix</code> , <code>specify</code>
Source	<pre> &lt;xs:group name="stage.content"&gt;   &lt;xs:sequence&gt;     &lt;xs:group ref="generic.content"/&gt;     &lt;xs:group ref="prefix"/&gt;     &lt;xs:group ref="multiplicity"/&gt;     &lt;xs:group ref="pT"/&gt;     &lt;xs:group ref="pN"/&gt;     &lt;xs:group ref="pM"/&gt;     &lt;xs:group ref="stage.group"/&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>

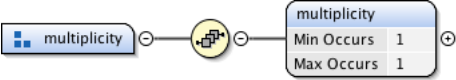
## Element Group prefix

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element Group stage.content
Model	prefix
Children	prefix
Source	<pre> &lt;xs:group name="prefix"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="prefix"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt;           &lt;xs:group ref="generic.content" /&gt;           &lt;xs:group ref="stage.prefix.response" /&gt;         &lt;/xs:sequence&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>

## Element Group stage.prefix.response

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element prefix/prefix
Model	response
Children	response
Source	<pre> &lt;xs:group name="stage.prefix.response"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="response"&gt;       &lt;xs:complexType&gt;         &lt;xs:group ref="generic.content" /&gt;         &lt;xs:attribute name="value" use="required"&gt;           &lt;xs:simpleType&gt;             &lt;xs:union memberTypes="val:base.values val:stage.prefix.values" /&gt;           &lt;/xs:simpleType&gt;         &lt;/xs:attribute&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>

## Element Group multiplicity

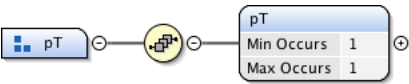
Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element Group stage.content
Model	multiplicity
Children	multiplicity
Source	<pre> &lt;xs:group name="multiplicity"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="multiplicity"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt;           &lt;xs:group ref="generic.content" /&gt;           &lt;xs:element name="response"&gt;             &lt;xs:complexType&gt;               &lt;xs:group ref="generic.content" /&gt;               &lt;xs:attribute name="value" use="required"&gt;                 &lt;xs:simpleType&gt; </pre>

```

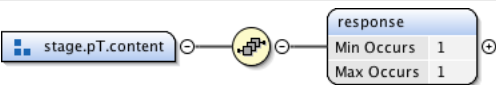
        <xs:union memberTypes="val:base.values multiplicity.valueset"/>
      </xs:simpleType>
    </xs:attribute>
  </xs:complexType>
</xs:element>
</xs:sequence>
</xs:complexType>
</xs:element>
</xs:sequence>
</xs:group>

```


## Element Group pT

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element Group stage.content
Model	pT
Children	pT
Source	<pre> &lt;xs:group name="pT"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="pT"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt;           &lt;xs:group ref="generic.content"/&gt;           &lt;xs:group ref="stage.pT.content"/&gt;         &lt;/xs:sequence&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>

## Element Group stage.pT.content

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element pT/pT
Model	response
Children	response
Source	<pre> &lt;xs:group name="stage.pT.content"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="response"&gt;       &lt;xs:complexType&gt;         &lt;xs:group ref="generic.content"/&gt;         &lt;xs:attribute name="value" use="required"&gt;           &lt;xs:simpleType&gt;             &lt;xs:restriction base="xs:token"&gt;               &lt;xs:enumeration value="specify"/&gt;               &lt;xs:enumeration value="inapplicable"/&gt;               &lt;xs:enumeration value="unreported"/&gt;             &lt;/xs:restriction&gt;           &lt;/xs:simpleType&gt;         &lt;/xs:attribute&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>

## Element Group pN

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element Group stage.content

Model	pN
Children	pN
Source	<pre> &lt;xs:group name="pN"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="pN"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt;           &lt;xs:group ref="generic.content"/&gt;           &lt;xs:group ref="stage.pN.content"/&gt;         &lt;/xs:sequence&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>

### Element Group stage.pN.content

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element pN/pN
Model	response
Children	response
Source	<pre> &lt;xs:group name="stage.pN.content"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="response"&gt;       &lt;xs:complexType&gt;         &lt;xs:group ref="generic.content"/&gt;         &lt;xs:attribute name="value" use="required"&gt;           &lt;xs:simpleType&gt;             &lt;xs:restriction base="xs:token"&gt;               &lt;xs:enumeration value="specify"/&gt;               &lt;xs:enumeration value="inapplicable"/&gt;               &lt;xs:enumeration value="unreported"/&gt;             &lt;/xs:restriction&gt;           &lt;/xs:simpleType&gt;         &lt;/xs:attribute&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>

### Element Group pM

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element Group stage.content
Model	pM
Children	pM
Source	<pre> &lt;xs:group name="pM"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="pM"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt;           &lt;xs:group ref="generic.content"/&gt;           &lt;xs:group ref="stage.pM.content"/&gt;         &lt;/xs:sequence&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>

### Element Group stage.pM.content

Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram	
Used by	Element pM/pM
Model	response
Children	response
Source	<pre> &lt;xs:group name="stage.pM.content"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="response"&gt;       &lt;xs:complexType&gt;         &lt;xs:group ref="generic.content" /&gt;         &lt;xs:attribute name="value" use="required"&gt;           &lt;xs:simpleType&gt;             &lt;xs:restriction base="xs:token"&gt;               &lt;xs:enumeration value="specify"/&gt;               &lt;xs:enumeration value="inapplicable"/&gt;               &lt;xs:enumeration value="unreported"/&gt;             &lt;/xs:restriction&gt;           &lt;/xs:simpleType&gt;         &lt;/xs:attribute&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>

### Element Group stage.group

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element Group stage.content
Model	group
Children	group
Source	<pre> &lt;xs:group name="stage.group"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="group"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt;           &lt;xs:group ref="generic.content" /&gt;           &lt;xs:group ref="stage.group.response" /&gt;         &lt;/xs:sequence&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>

### Element Group stage.group.response

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element stage.group/group
Model	response
Children	response
Source	<pre> &lt;xs:group name="stage.group.response"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="response"&gt;       &lt;xs:complexType&gt;         &lt;xs:group ref="generic.content" /&gt;         &lt;xs:attribute name="value" type="stage.group.valueset" use="required"/&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>



## Element Group `studies.content`

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element <code>studies</code>
Model	<code>specify{0,1}</code> , <code>label{0,1}</code> , <code>comment{0,1}</code> , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , <code>study</code>
Children	<code>comment</code> , <code>label</code> , <code>specify</code> , <code>study</code>
Source	<pre> &lt;xs:group name="studies.content"&gt;   &lt;xs:sequence&gt;     &lt;xs:group ref="generic.content"/&gt;     &lt;xs:element name="study"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt;           &lt;xs:group ref="generic.content"/&gt;           &lt;xs:group ref="studies.study.content"/&gt;         &lt;/xs:sequence&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>

## Element Group `studies.study.content`

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element <code>studies.content/study</code>
Model	
Source	<pre> &lt;xs:group name="studies.study.content"&gt;   &lt;xs:sequence/&gt; &lt;/xs:group&gt; </pre>

## Element Group `tumor.content`

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element <code>tumor</code>

Model	specify{0,1} , label{0,1} , comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , focus
Children	comment, focus, label, specify
Source	<pre> &lt;xs:group name="tumor.content"&gt;   &lt;xs:sequence&gt;     &lt;xs:group ref="generic.content" /&gt;     &lt;xs:group maxOccurs="unbounded" ref="focus" /&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>

### Element Group focus

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element Group tumor.content
Model	focus
Children	focus
Source	<pre> &lt;xs:group name="focus"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="focus"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt;           &lt;xs:group ref="generic.content" /&gt;           &lt;xs:group ref="site" /&gt;           &lt;xs:group ref="size" /&gt;           &lt;xs:group ref="histologicType" /&gt;           &lt;xs:group ref="grade" /&gt;         &lt;/xs:sequence&gt;         &lt;xs:attribute name="type" use="required"&gt;           &lt;xs:simpleType&gt;             &lt;xs:restriction base="xs:token"&gt;               &lt;xs:enumeration value="principal" /&gt;               &lt;xs:enumeration value="sole" /&gt;               &lt;xs:enumeration value="additional" /&gt;             &lt;/xs:restriction&gt;           &lt;/xs:simpleType&gt;         &lt;/xs:attribute&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>

### Element Group histologicType

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element focus/focus
Model	histologicType
Children	histologicType
Source	<pre> &lt;xs:group name="histologicType"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="histologicType"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt;           &lt;xs:group ref="tumor.histologicType.content" /&gt;           &lt;xs:group ref="generic.content" /&gt;         &lt;/xs:sequence&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>

### Element Group tumor.histologicType.content

Namespace	http://purl.org/obbr/pathology/cancer/
-----------	----------------------------------------

Diagram	
Used by	Element histologicType/histologicType
Model	response
Children	response
Source	<pre> &lt;xs:group name="tumor.histologicType.content"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="response"&gt;       &lt;xs:complexType&gt;         &lt;xs:group ref="generic.content" /&gt;         &lt;xs:attribute name="value" use="required"&gt;           &lt;xs:simpleType&gt;             &lt;xs:union memberTypes="val:base.values tumor.histologicType.valueset" /&gt;           &lt;/xs:simpleType&gt;         &lt;/xs:attribute&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>

## Element Group grade

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element focus/focus
Model	grade
Children	grade
Source	<pre> &lt;xs:group name="grade"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="grade"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt;           &lt;xs:group ref="generic.content" /&gt;           &lt;xs:group ref="tumor.grade.response" /&gt;         &lt;/xs:sequence&gt;         &lt;xs:attribute name="system" type="tumor.gradeSystem.valueset" use="required" /&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>

## Element Group tumor.grade.response

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element grade/grade
Model	response
Children	response
Source	<pre> &lt;xs:group name="tumor.grade.response"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="response"&gt;       &lt;xs:complexType&gt;         &lt;xs:group ref="generic.content" /&gt;         &lt;xs:attribute name="value" use="required"&gt;           &lt;xs:simpleType&gt;             &lt;xs:union memberTypes="val:base.values tumor.grade.valueset" /&gt;           &lt;/xs:simpleType&gt;         &lt;/xs:attribute&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>

```
</xs:sequence>
</xs:group>
```

## Attribute Groups

### Attribute Group `generic.attrib`

Namespace	http://purl.org/obbr/pathology/cancer/				
Diagram					
Used by	Elements accessory, accessory.lymphvascular.content/response, accessory.lymphvascular/lymphvascularInvasion, accessory.perineural.content/response, accessory.perineural/perineuralInvasion, accessory.treatment.content/response, accessory.treatment/treatmentResponse, additional, additional.content/finding, diagnosis, distant, extent, history, margin, nodes, specimen, specimen.weight.content/response, stage, studies, tumor				
Attributes	QName	Type	Fixed	Default	Use
	ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'				
	id	xs:ID			optional
Source	<pre>&lt;xs:attributeGroup name="generic.attrib"&gt;   &lt;xs:attribute name="id" type="xs:ID" xmlns="http://www.w3.org/XML/1998/namespace/" /&gt;   &lt;xs:anyAttribute namespace="##other" /&gt; &lt;/xs:attributeGroup&gt;</pre>				

### Attribute Group `date`

Namespace	http://purl.org/obbr/pathology/cancer/				
Diagram					
Used by	Elements clinicalFinding/clinicalFinding, distant.status.result/result, pathologicFinding/pathologicFinding, priorTherapy/priorTherapy, radiologicFinding/radiologicFinding				
Attributes	QName	Type	Fixed	Default	Use
	date	date.valueset			optional
	duration	xs:duration			optional
	end	date.valueset			optional
	start	date.valueset			optional
Source	<pre>&lt;xs:attributeGroup name="date"&gt;   &lt;xs:attribute name="date" type="date.valueset" /&gt;   &lt;xs:attribute name="start" type="date.valueset" /&gt;   &lt;xs:attribute name="end" type="date.valueset" /&gt;   &lt;xs:attribute name="duration" type="xs:duration" /&gt; &lt;/xs:attributeGroup&gt;</pre>				

### Attribute Group `dimensionalQuantity`

Namespace	http://purl.org/obbr/pathology/cancer/
-----------	----------------------------------------

Diagram					
Used by	Elements	length/response, size/response			
Attributes	QName	Type	Fixed	Default	Use
	relation	relation.valueset			optional
	unit	lengthUnit.valueset			required
	value	positiveDecimal			required
Source	<pre>&lt;xs:attributeGroup name="dimensionalQuantity"&gt;   &lt;xs:attribute name="value" type="positiveDecimal" use="required"/&gt;   &lt;xs:attribute name="unit" type="lengthUnit.valueset" use="required"/&gt;   &lt;xs:attribute name="relation" type="relation.valueset"/&gt; &lt;/xs:attributeGroup&gt;</pre>				

### Attribute Group weight

Namespace	http://purl.org/obbr/pathology/cancer/				
Diagram					
Used by	Element	specimen.weight.content/response			
Attributes	QName	Type	Fixed	Default	Use
	relation	relation.valueset			optional
	unit	weightUnit.valueset			required
	value	weight.valueset			required
Source	<pre>&lt;xs:attributeGroup name="weight"&gt;   &lt;xs:attribute name="value" type="weight.valueset" use="required"/&gt;   &lt;xs:attribute name="unit" type="weightUnit.valueset" use="required"/&gt;   &lt;xs:attribute name="relation" type="relation.valueset"/&gt; &lt;/xs:attributeGroup&gt;</pre>				

Namespace: "http://purl.org/obbr/pathology/cancer/values/"

## Schemas

### Imported schema values.xsd

Namespace	http://purl.org/obbr/pathology/cancer/values/
Properties	attribute form default: unqualified
	element form default: unqualified

## Simple Types

### Simple Type val:base.values

Namespace	http://purl.org/obbr/pathology/cancer/values/
Diagram	
Type	restriction of xs:token
Facets	enumeration specify

	enumeration	inapplicable
	enumeration	unreported
Used by	Attribute	contiguousExtension.status.distance/distance/@from
Source	<pre>&lt;xs:simpleType name="base.values"&gt;   &lt;xs:restriction base="xs:token"&gt;     &lt;xs:enumeration value="specify"/&gt;     &lt;xs:enumeration value="inapplicable"/&gt;     &lt;xs:enumeration value="unreported"/&gt;   &lt;/xs:restriction&gt; &lt;/xs:simpleType&gt;</pre>	

### Simple Type val:qualitative.values

Namespace	http://purl.org/obbr/pathology/cancer/values/	
Diagram	<pre> graph LR     A[qualitative.values] --- B(( ))     B --- C[xs.token]   </pre>	
Type	restriction of xs:token	
Facets	enumeration	positive
	enumeration	negative
	enumeration	equivocal
	enumeration	indeterminate
Source	<pre>&lt;xs:simpleType name="qualitative.values"&gt;   &lt;xs:restriction base="xs:token"&gt;     &lt;xs:enumeration value="positive"/&gt;     &lt;xs:enumeration value="negative"/&gt;     &lt;xs:enumeration value="equivocal"/&gt;     &lt;xs:enumeration value="indeterminate"/&gt;   &lt;/xs:restriction&gt; &lt;/xs:simpleType&gt;</pre>	

### Simple Type val:lengthUnit.values

Namespace	http://purl.org/obbr/pathology/cancer/values/	
Diagram	<pre> graph LR     A[lengthUnit.values] --- B(( ))     B --- C[xs.token]   </pre>	
Type	restriction of xs:token	
Facets	enumeration	cm
	enumeration	mm
	enumeration	in
Source	<pre>&lt;xs:simpleType name="lengthUnit.values"&gt;   &lt;xs:restriction base="xs:token"&gt;     &lt;xs:enumeration value="cm"/&gt;     &lt;xs:enumeration value="mm"/&gt;     &lt;xs:enumeration value="in"/&gt;   &lt;/xs:restriction&gt; &lt;/xs:simpleType&gt;</pre>	

### Simple Type val:weightUnit.values

Namespace	http://purl.org/obbr/pathology/cancer/values/	
Diagram	<pre> graph LR     A[weightUnit.values] --- B(( ))     B --- C[xs.token]   </pre>	
Type	restriction of xs:token	
Facets	enumeration	mg
	enumeration	g
	enumeration	kg
Source	<pre>&lt;xs:simpleType name="weightUnit.values"&gt;   &lt;xs:restriction base="xs:token"&gt;     &lt;xs:enumeration value="mg"/&gt;     &lt;xs:enumeration value="g"/&gt;     &lt;xs:enumeration value="kg"/&gt;   &lt;/xs:restriction&gt; &lt;/xs:simpleType&gt;</pre>	

	</xs:simpleType>
--	------------------

### Simple Type val:scalar.values

Namespace	http://purl.org/obbr/pathology/cancer/values/
Diagram	<pre> graph LR     scalar.values --- val.base.values     scalar.values --- restricts_xs_float["restricts: xs:float"]   </pre>
Type	union of(val:base.values, restriction of xs:float)
Source	<pre> &lt;xs:simpleType name="scalar.values"&gt;   &lt;xs:union memberTypes="val:base.values"&gt;     &lt;xs:simpleType&gt;       &lt;xs:restriction base="xs:float"&gt;         &lt;xs:minExclusive value="0.0"/&gt;       &lt;/xs:restriction&gt;     &lt;/xs:simpleType&gt;   &lt;/xs:union&gt; &lt;/xs:simpleType&gt; </pre>


### Simple Type val:relation.values

Namespace	http://purl.org/obbr/pathology/cancer/values/												
Diagram	<pre> graph LR     relation.values --- xs.token   </pre>												
Type	restriction of xs:token												
Facets	<table> <tr><td>enumeration</td><td>less than</td></tr> <tr><td>enumeration</td><td>less than or equal</td></tr> <tr><td>enumeration</td><td>equal</td></tr> <tr><td>enumeration</td><td>approximately equal</td></tr> <tr><td>enumeration</td><td>greater than or equal</td></tr> <tr><td>enumeration</td><td>greater than</td></tr> </table>	enumeration	less than	enumeration	less than or equal	enumeration	equal	enumeration	approximately equal	enumeration	greater than or equal	enumeration	greater than
enumeration	less than												
enumeration	less than or equal												
enumeration	equal												
enumeration	approximately equal												
enumeration	greater than or equal												
enumeration	greater than												
Source	<pre> &lt;xs:simpleType name="relation.values"&gt;   &lt;xs:restriction base="xs:token"&gt;     &lt;xs:enumeration value="less than"/&gt;     &lt;xs:enumeration value="less than or equal"/&gt;     &lt;xs:enumeration value="equal"/&gt;     &lt;xs:enumeration value="approximately equal"/&gt;     &lt;xs:enumeration value="greater than or equal"/&gt;     &lt;xs:enumeration value="greater than"/&gt;   &lt;/xs:restriction&gt; &lt;/xs:simpleType&gt; </pre>												


### Simple Type val:laterality.values

Namespace	http://purl.org/obbr/pathology/cancer/values/												
Diagram	<pre> graph LR     laterality.values --- xs.token   </pre>												
Type	restriction of xs:token												
Facets	<table> <tr><td>enumeration</td><td>ipsilateral</td></tr> <tr><td>enumeration</td><td>contralateral</td></tr> <tr><td>enumeration</td><td>right</td></tr> <tr><td>enumeration</td><td>left</td></tr> <tr><td>enumeration</td><td>midline</td></tr> <tr><td>enumeration</td><td>bilateral</td></tr> </table>	enumeration	ipsilateral	enumeration	contralateral	enumeration	right	enumeration	left	enumeration	midline	enumeration	bilateral
enumeration	ipsilateral												
enumeration	contralateral												
enumeration	right												
enumeration	left												
enumeration	midline												
enumeration	bilateral												
Source	<pre> &lt;xs:simpleType name="laterality.values"&gt;   &lt;xs:restriction base="xs:token"&gt;     &lt;xs:enumeration value="ipsilateral"/&gt;     &lt;xs:enumeration value="contralateral"/&gt;     &lt;xs:enumeration value="right"/&gt;     &lt;xs:enumeration value="left"/&gt;     &lt;xs:enumeration value="midline"/&gt;     &lt;xs:enumeration value="bilateral"/&gt;   &lt;/xs:restriction&gt; &lt;/xs:simpleType&gt; </pre>												


**Simple Type val:focality.values**

Namespace	http://purl.org/obbr/pathology/cancer/values/								
Diagram									
Type	restriction of xs:token								
Facets	<table border="1"> <tr><td>enumeration</td><td>microfocal</td></tr> <tr><td>enumeration</td><td>unifocal</td></tr> <tr><td>enumeration</td><td>multifocal</td></tr> <tr><td>enumeration</td><td>extensive</td></tr> </table>	enumeration	microfocal	enumeration	unifocal	enumeration	multifocal	enumeration	extensive
enumeration	microfocal								
enumeration	unifocal								
enumeration	multifocal								
enumeration	extensive								
Source	<pre>&lt;xs:simpleType name="focality.values"&gt;   &lt;xs:restriction base="xs:token"&gt;     &lt;xs:enumeration value="microfocal"/&gt;     &lt;xs:enumeration value="unifocal"/&gt;     &lt;xs:enumeration value="multifocal"/&gt;     &lt;xs:enumeration value="extensive"/&gt;   &lt;/xs:restriction&gt; &lt;/xs:simpleType&gt;</pre>								

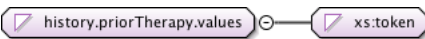
**Simple Type val:accessory.treatmentResponse.values**

Namespace	http://purl.org/obbr/pathology/cancer/values/								
Diagram									
Type	restriction of xs:token								
Facets	<table border="1"> <tr><td>enumeration</td><td>none</td></tr> <tr><td>enumeration</td><td>slight</td></tr> <tr><td>enumeration</td><td>marked</td></tr> <tr><td>enumeration</td><td>complete</td></tr> </table>	enumeration	none	enumeration	slight	enumeration	marked	enumeration	complete
enumeration	none								
enumeration	slight								
enumeration	marked								
enumeration	complete								
Source	<pre>&lt;xs:simpleType name="accessory.treatmentResponse.values"&gt;   &lt;xs:restriction base="xs:token"&gt;     &lt;xs:enumeration value="none"/&gt;     &lt;xs:enumeration value="slight"/&gt;     &lt;xs:enumeration value="marked"/&gt;     &lt;xs:enumeration value="complete"/&gt;   &lt;/xs:restriction&gt; &lt;/xs:simpleType&gt;</pre>								

**Simple Type val:extent.contiguousExtension.status.result.for.values**

Namespace	http://purl.org/obbr/pathology/cancer/values/				
Diagram					
Type	restriction of xs:token				
Facets	<table border="1"> <tr><td>enumeration</td><td>invades</td></tr> <tr><td>enumeration</td><td>adheres</td></tr> </table>	enumeration	invades	enumeration	adheres
enumeration	invades				
enumeration	adheres				
Source	<pre>&lt;xs:simpleType name="extent.contiguousExtension.status.result.for.values"&gt;   &lt;xs:restriction base="xs:token"&gt;     &lt;xs:enumeration value="invades"/&gt;     &lt;xs:enumeration value="adheres"/&gt;   &lt;/xs:restriction&gt; &lt;/xs:simpleType&gt;</pre>				


**Simple Type val:history.priorTherapy.values**

Namespace	http://purl.org/obbr/pathology/cancer/values/						
Diagram							
Type	restriction of xs:token						
Facets	<table border="1"> <tr><td>enumeration</td><td>chemotherapy</td></tr> <tr><td>enumeration</td><td>radiation</td></tr> <tr><td>enumeration</td><td>surgery</td></tr> </table>	enumeration	chemotherapy	enumeration	radiation	enumeration	surgery
enumeration	chemotherapy						
enumeration	radiation						
enumeration	surgery						




Source	<pre> &lt;xs:simpleType name="history.priorTherapy.values"&gt;   &lt;xs:restriction base="xs:token"&gt;     &lt;xs:enumeration value="chemotherapy"/&gt;     &lt;xs:enumeration value="radiation"/&gt;     &lt;xs:enumeration value="surgery"/&gt;   &lt;/xs:restriction&gt; &lt;/xs:simpleType&gt; </pre>
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### Simple Type val:margin.status.result.for.values

Namespace	http://purl.org/obbr/pathology/cancer/values/				
Diagram					
Type	restriction of xs:token				
Facets	<table> <tr> <td>enumeration</td><td>in-situ</td></tr> <tr> <td>enumeration</td><td>invasive</td></tr> </table>	enumeration	in-situ	enumeration	invasive
enumeration	in-situ				
enumeration	invasive				
Source	<pre> &lt;xs:simpleType name="margin.status.result.for.values"&gt;   &lt;xs:restriction base="xs:token"&gt;     &lt;xs:enumeration value="in-situ"/&gt;     &lt;xs:enumeration value="invasive"/&gt;   &lt;/xs:restriction&gt; &lt;/xs:simpleType&gt; </pre>				

### Simple Type val:distant.status.of.values


Namespace	http://purl.org/obbr/pathology/cancer/values/								
Diagram									
Type	restriction of xs:token								
Facets	<table> <tr> <td>enumeration</td><td>liver</td></tr> <tr> <td>enumeration</td><td>bone</td></tr> <tr> <td>enumeration</td><td>lung</td></tr> <tr> <td>enumeration</td><td>brain</td></tr> </table>	enumeration	liver	enumeration	bone	enumeration	lung	enumeration	brain
enumeration	liver								
enumeration	bone								
enumeration	lung								
enumeration	brain								
Source	<pre> &lt;xs:simpleType name="distant.status.of.values"&gt;   &lt;xs:restriction base="xs:token"&gt;     &lt;xs:enumeration value="liver"/&gt;     &lt;xs:enumeration value="bone"/&gt;     &lt;xs:enumeration value="lung"/&gt;     &lt;xs:enumeration value="brain"/&gt;   &lt;/xs:restriction&gt; &lt;/xs:simpleType&gt; </pre>								

### Simple Type val:distant.status.result.for.values


Namespace	http://purl.org/obbr/pathology/cancer/values/								
Diagram									
Type	restriction of xs:token								
Facets	<table> <tr> <td>enumeration</td><td>this accession</td></tr> <tr> <td>enumeration</td><td>prior accession</td></tr> <tr> <td>enumeration</td><td>outside report</td></tr> <tr> <td>enumeration</td><td>clinical history</td></tr> </table>	enumeration	this accession	enumeration	prior accession	enumeration	outside report	enumeration	clinical history
enumeration	this accession								
enumeration	prior accession								
enumeration	outside report								
enumeration	clinical history								
Source	<pre> &lt;xs:simpleType name="distant.status.result.for.values"&gt;   &lt;xs:restriction base="xs:token"&gt;     &lt;xs:enumeration value="this accession"/&gt;     &lt;xs:enumeration value="prior accession"/&gt;     &lt;xs:enumeration value="outside report"/&gt;     &lt;xs:enumeration value="clinical history"/&gt;   &lt;/xs:restriction&gt; &lt;/xs:simpleType&gt; </pre>								

### Simple Type val:nodes.status.result.for.values

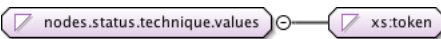
Namespace	http://purl.org/obbr/pathology/cancer/values/
-----------	-----------------------------------------------

Diagram					
Type	restriction of xs:token				
Facets	<table border="1"> <tr> <td>enumeration</td><td>negative</td></tr> <tr> <td>enumeration</td><td>positive</td></tr> </table>	enumeration	negative	enumeration	positive
enumeration	negative				
enumeration	positive				
Source	<pre>&lt;xs:simpleType name="nodes.status.result.for.values"&gt;   &lt;xs:restriction base="xs:token"&gt;     &lt;xs:enumeration value="negative"/&gt;     &lt;xs:enumeration value="positive"/&gt;   &lt;/xs:restriction&gt; &lt;/xs:simpleType&gt;</pre>				


### Simple Type val:nodes.status.result.type.values

Namespace	http://purl.org/obbr/pathology/cancer/values/		
Diagram			
Type	restriction of xs:token		
Facets	<table border="1"> <tr> <td>enumeration</td><td>necroinflammatory</td></tr> </table>	enumeration	necroinflammatory
enumeration	necroinflammatory		
Source	<pre>&lt;xs:simpleType name="nodes.status.result.type.values"&gt;   &lt;xs:restriction base="xs:token"&gt;     &lt;xs:enumeration value="necroinflammatory"/&gt;   &lt;/xs:restriction&gt; &lt;/xs:simpleType&gt;</pre>		


### Simple Type val:nodes.status.technique.values

Namespace	http://purl.org/obbr/pathology/cancer/values/						
Diagram							
Type	restriction of xs:token						
Facets	<table border="1"> <tr> <td>enumeration</td><td>H+E one level</td></tr> <tr> <td>enumeration</td><td>H+E multiple level(s)</td></tr> <tr> <td>enumeration</td><td>H+E plus immunohistochemistry</td></tr> </table>	enumeration	H+E one level	enumeration	H+E multiple level(s)	enumeration	H+E plus immunohistochemistry
enumeration	H+E one level						
enumeration	H+E multiple level(s)						
enumeration	H+E plus immunohistochemistry						
Source	<pre>&lt;xs:simpleType name="nodes.status.technique.values"&gt;   &lt;xs:restriction base="xs:token"&gt;     &lt;xs:enumeration value="H+E one level"/&gt;     &lt;xs:enumeration value="H+E multiple level(s)"/&gt;     &lt;xs:enumeration value="H+E plus immunohistochemistry"/&gt;   &lt;/xs:restriction&gt; &lt;/xs:simpleType&gt;</pre>						

### Simple Type val:specimen.integrity.values


Namespace	http://purl.org/obbr/pathology/cancer/values/				
Diagram					
Type	restriction of xs:token				
Facets	<table border="1"> <tr> <td>enumeration</td><td>intact</td></tr> <tr> <td>enumeration</td><td>fragmented</td></tr> </table>	enumeration	intact	enumeration	fragmented
enumeration	intact				
enumeration	fragmented				
Source	<pre>&lt;xs:simpleType name="specimen.integrity.values"&gt;   &lt;xs:restriction base="xs:token"&gt;     &lt;xs:enumeration value="intact"/&gt;     &lt;xs:enumeration value="fragmented"/&gt;   &lt;/xs:restriction&gt; &lt;/xs:simpleType&gt;</pre>				

### Simple Type val:stage.prefix.values


Namespace	http://purl.org/obbr/pathology/cancer/values/
Diagram	

Type	restriction of xs:token
Facets	enumeration y
	enumeration r
	enumeration a
	enumeration none
Source	<pre>&lt;xs:simpleType name="stage.prefix.values"&gt;   &lt;xs:restriction base="xs:token"&gt;     &lt;xs:enumeration value="y"/&gt;     &lt;xs:enumeration value="r"/&gt;     &lt;xs:enumeration value="a"/&gt;     &lt;xs:enumeration value="none"/&gt;   &lt;/xs:restriction&gt; &lt;/xs:simpleType&gt;</pre>


### Simple Type val:multiplicity.values

Namespace	http://purl.org/obbr/pathology/cancer/values/
Diagram	
Type	restriction of xs:token
Facets	enumeration 1
	enumeration 2
	enumeration 3
	enumeration 4
	enumeration m
Source	<pre>&lt;xs:simpleType name="multiplicity.values"&gt;   &lt;xs:restriction base="xs:token"&gt;     &lt;xs:enumeration value="1"/&gt;     &lt;xs:enumeration value="2"/&gt;     &lt;xs:enumeration value="3"/&gt;     &lt;xs:enumeration value="4"/&gt;     &lt;xs:enumeration value="m"/&gt;   &lt;/xs:restriction&gt; &lt;/xs:simpleType&gt;</pre>

### Simple Type val:tumor.grade.values

Namespace	http://purl.org/obbr/pathology/cancer/values/
Diagram	
Type	restriction of xs:token
Facets	enumeration 1
	enumeration 2
	enumeration 3
	enumeration 4
	enumeration X
Source	<pre>&lt;xs:simpleType name="tumor.grade.values"&gt;   &lt;xs:restriction base="xs:token"&gt;     &lt;xs:enumeration value="1"/&gt;     &lt;xs:enumeration value="2"/&gt;     &lt;xs:enumeration value="3"/&gt;     &lt;xs:enumeration value="4"/&gt;     &lt;xs:enumeration value="X"/&gt;   &lt;/xs:restriction&gt; &lt;/xs:simpleType&gt;</pre>

### Simple Type val:tumor.gradeSystem.values


Namespace	http://purl.org/obbr/pathology/cancer/values/
Diagram	
Type	restriction of xs:token

Facets	enumeration 4-grade
Source	<pre>&lt;xs:simpleType name="tumor.gradeSystem.values"&gt;   &lt;xs:restriction base="xs:token"&gt;     &lt;xs:enumeration value="4-grade"/&gt;   &lt;/xs:restriction&gt; &lt;/xs:simpleType&gt;</pre>


**Namespace:** ""

## Elements

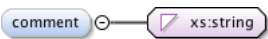
### Element `generic.content / specify`

Namespace	No namespace				
Diagram					
Type	xs:string				
Properties	<table> <tr> <td>content:</td><td>simple</td></tr> <tr> <td>minOccurs:</td><td>0</td></tr> </table>	content:	simple	minOccurs:	0
content:	simple				
minOccurs:	0				
Source	<pre>&lt;xs:element minOccurs="0" name="specify" type="xs:string"/&gt;</pre>				

### Element `generic.content / label`

Namespace	No namespace				
Diagram					
Type	xs:string				
Properties	<table> <tr> <td>content:</td><td>simple</td></tr> <tr> <td>minOccurs:</td><td>0</td></tr> </table>	content:	simple	minOccurs:	0
content:	simple				
minOccurs:	0				
Source	<pre>&lt;xs:element minOccurs="0" name="label" type="xs:string"/&gt;</pre>				

### Element `generic.content / comment`

Namespace	No namespace				
Diagram					
Type	xs:string				
Properties	<table> <tr> <td>content:</td><td>simple</td></tr> <tr> <td>minOccurs:</td><td>0</td></tr> </table>	content:	simple	minOccurs:	0
content:	simple				
minOccurs:	0				
Source	<pre>&lt;xs:element minOccurs="0" name="comment" type="xs:string"/&gt;</pre>				

## Attributes

### Attribute `generic.attrib / @id`

Namespace	No namespace		
Type	xs:ID		
Properties	<table> <tr> <td>content:</td><td>simple</td></tr> </table>	content:	simple
content:	simple		
Used by	Attribute Group generic.attrib		
Source	<pre>&lt;xs:attribute name="id" type="xs:ID" xmlns="http://www.w3.org/XML/1998/namespace/" /&gt;</pre>		

### Attribute `date / @date`

Namespace	No namespace		
Type	date.valueset		
Properties	<table> <tr> <td>content:</td><td>simple</td></tr> </table>	content:	simple
content:	simple		

Used by	Attribute Group <code>date</code>
Source	<code>&lt;xs:attribute name="date" type="date.valueset"/&gt;</code>

**Attribute `date` / `@start`**

Namespace	No namespace
Type	<code>date.valueset</code>
Properties	content: <code>simple</code>
Used by	Attribute Group <code>date</code>
Source	<code>&lt;xs:attribute name="start" type="date.valueset"/&gt;</code>

**Attribute `date` / `@end`**

Namespace	No namespace
Type	<code>date.valueset</code>
Properties	content: <code>simple</code>
Used by	Attribute Group <code>date</code>
Source	<code>&lt;xs:attribute name="end" type="date.valueset"/&gt;</code>

**Attribute `date` / `@duration`**

Namespace	No namespace
Type	<code>xs:duration</code>
Properties	content: <code>simple</code>
Used by	Attribute Group <code>date</code>
Source	<code>&lt;xs:attribute name="duration" type="xs:duration"/&gt;</code>

**Attribute `dimensionalQuantity` / `@value`**

Namespace	No namespace
Type	<code>positiveDecimal</code>
Properties	use: <code>required</code>
Facets	<code>minInclusive</code> <code>0.0</code>
Used by	Attribute Group <code>dimensionalQuantity</code>
Source	<code>&lt;xs:attribute name="value" type="positiveDecimal" use="required"/&gt;</code>

**Attribute `dimensionalQuantity` / `@unit`**

Namespace	No namespace
Type	<code>lengthUnit.valueset</code>
Properties	use: <code>required</code>
Used by	Attribute Group <code>dimensionalQuantity</code>
Source	<code>&lt;xs:attribute name="unit" type="lengthUnit.valueset" use="required"/&gt;</code>

**Attribute `dimensionalQuantity` / `@relation`**

Namespace	No namespace
Type	<code>relation.valueset</code>
Properties	content: <code>simple</code>
Used by	Attribute Group <code>dimensionalQuantity</code>
Source	<code>&lt;xs:attribute name="relation" type="relation.valueset"/&gt;</code>

**Attribute weight / @value**

Namespace	No namespace
Type	weight.valueset
Properties	use: required
Used by	Attribute Group weight
Source	<xs:attribute name="value" type="weight.valueset" use="required"/>

**Attribute weight / @unit**

Namespace	No namespace
Type	weightUnit.valueset
Properties	use: required
Used by	Attribute Group weight
Source	<xs:attribute name="unit" type="weightUnit.valueset" use="required"/>

**Attribute weight / @relation**

Namespace	No namespace
Type	relation.valueset
Properties	content: simple
Used by	Attribute Group weight
Source	<xs:attribute name="relation" type="relation.valueset"/>

**Attribute accessory.treatment.content / response / @value**

Namespace	No namespace
Type	accessory.treatmentResponse.valueset
Properties	use: required
Used by	Element accessory.treatment.content/response
Source	<xs:attribute name="value" type="accessory.treatmentResponse.valueset" use="required"/>

**Attribute accessory.lymphvascular.content / response / @value**

Namespace	No namespace
Type	accessory.lymphvascular.valueset
Properties	use: required
Used by	Element accessory.lymphvascular.content/response
Source	<xs:attribute name="value" type="accessory.lymphvascular.valueset" use="required"/>

**Attribute accessory.perineural.content / response / @value**

Namespace	No namespace
Type	accessory.perineural.valueset
Properties	use: required
Used by	Element accessory.perineural.content/response
Source	<xs:attribute name="value" type="accessory.perineural.valueset" use="required"/>

**Attribute additional.finding.content / response / @value**

Namespace	No namespace
-----------	--------------

Type	additional.finding.valueset
Properties	use: required
Used by	Element additional.finding.content/response
Source	<xs:attribute name="value" type="additional.finding.valueset" use="required"/>

**Attribute additional.content / finding / @of**

Namespace	No namespace
Type	additional.finding.of.valueset
Properties	content: simple
Used by	Element additional.content/finding
Source	<xs:attribute name="of" type="additional.finding.of.valueset"/>

**Attribute distant.status.result.response / response / @value**

Namespace	No namespace
Type	union of(val:base.values, val:qualitative.values)
Properties	use: required
Used by	Element distant.status.result.response/response
Source	<pre>&lt;xs:attribute name="value" use="required"&gt;   &lt;xs:simpleType&gt;     &lt;xs:union memberTypes="val:base.values val:qualitative.values"/&gt;   &lt;/xs:simpleType&gt; &lt;/xs:attribute&gt;</pre>

**Attribute distant.status.result / result / @for**

Namespace	No namespace
Type	distant.status.result.for.valueset
Properties	content: simple
Used by	Element distant.status.result/result
Source	<xs:attribute name="for" type="distant.status.result.for.valueset"/>

**Attribute distant.status / status / @of**

Namespace	No namespace
Type	distant.status.of.valueset
Properties	use: required
Used by	Element distant.status/status
Source	<xs:attribute name="of" type="distant.status.of.valueset" use="required"/>

**Attribute distant.status / status / @on**

Namespace	No namespace
Type	laterality.valueset
Properties	content: simple
Used by	Element distant.status/status
Source	<xs:attribute name="on" type="laterality.valueset"/>

**Attribute generic.content / @value**

Namespace	No namespace
Type	union of(val:base.values, val:qualitative.values)

Properties	use: required
Used by	Elements clinicalFinding.content/response, contiguousExtension.status.result.content/response, margins.margin.status.result.content/response, multiplicity/multiplicity/response, nodes.extracapsularExtension.content/response, pathologicFinding.content/response, priorTherapy.content/response, radiologicFinding.content/response, site.content/response, specimen.integrity.content/response, specimen.procedure.content/response, stage.group.response/response, stage.pM.content/response, stage.pN.content/response, stage.pT.content/response, stage.prefix.response/response, tumor.grade.response/response, tumor.histologicType.content/response
Source	<pre>&lt;xs:attribute name="value" use="required"&gt;   &lt;xs:simpleType&gt;     &lt;xs:union memberTypes="val:base.values val:qualitative.values"/&gt;   &lt;/xs:simpleType&gt; &lt;/xs:attribute&gt;</pre>

**Attribute generic.content / @extent**

Namespace	No namespace
Type	focality.valueset
Properties	content: simple
Used by	Elements contiguousExtension.status.result.content/response, margins.margin.status.result.content/response
Source	<pre>&lt;xs:attribute name="extent" type="focality.valueset"/&gt;</pre>

**Attribute contiguousExtension.status.result.content / response / @extent**

Namespace	No namespace
Type	focality.valueset
Properties	content: simple
Source	<pre>&lt;xs:attribute name="extent" type="focality.valueset"/&gt;</pre>

**Attribute contiguousExtension.status.result.content / response / @value**

Namespace	No namespace
Type	union of(val:base.values, val:qualitative.values)
Properties	use: required
Source	<pre>&lt;xs:attribute name="value" use="required"&gt;   &lt;xs:simpleType&gt;     &lt;xs:union memberTypes="val:base.values val:qualitative.values"/&gt;   &lt;/xs:simpleType&gt; &lt;/xs:attribute&gt;</pre>

**Attribute contiguousExtension.status.result / result / @for**

Namespace	No namespace
Type	union of(val:base.values, restriction of xs:token)
Properties	use: required
Used by	Element contiguousExtension.status.result/result
Source	<pre>&lt;xs:attribute name="for" use="required"&gt;   &lt;xs:simpleType&gt;     &lt;xs:union memberTypes="val:base.values"&gt;       &lt;xs:simpleType&gt;         &lt;xs:restriction base="xs:token"&gt;           &lt;xs:enumeration value="invades"/&gt;           &lt;xs:enumeration value="adheres"/&gt;           &lt;xs:enumeration value="specify"/&gt;           &lt;xs:enumeration value="inapplicable"/&gt;           &lt;xs:enumeration value="unreported"/&gt;         &lt;/xs:restriction&gt;       &lt;/xs:simpleType&gt;     &lt;/xs:union&gt;   &lt;/xs:simpleType&gt;</pre>



	</xs:attribute>
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**Attribute contiguousExtension.status.result / result / @on**

Namespace	No namespace
Type	laterality.valueset
Properties	content: simple
Used by	Element contiguousExtension.status.result/result
Source	<xs:attribute name="on" type="laterality.valueset"/>

**Attribute contiguousExtension.status.distance / distance / @for**

Namespace	No namespace
Type	restriction of xs:token
Properties	content: simple
Facets	enumeration invades enumeration adheres
Used by	Element contiguousExtension.status.distance/distance
Source	<pre>&lt;xs:attribute name="for"&gt;   &lt;xs:simpleType&gt;     &lt;xs:restriction base="xs:token"&gt;       &lt;xs:enumeration value="invades"/&gt;       &lt;xs:enumeration value="adheres"/&gt;     &lt;/xs:restriction&gt;   &lt;/xs:simpleType&gt; &lt;/xs:attribute&gt;</pre>

**Attribute contiguousExtension.status.distance / distance / @from**

Namespace	No namespace
Type	val:base.values
Properties	content: simple
Facets	enumeration specify enumeration inapplicable enumeration unreported
Used by	Element contiguousExtension.status.distance/distance
Source	<xs:attribute name="from" type="val:base.values"/>

**Attribute contiguousExtensionStatus / status / @of**

Namespace	No namespace
Type	extent.status.of.valueset
Properties	content: simple
Used by	Element contiguousExtensionStatus/status
Source	<xs:attribute name="of" type="extent.status.of.valueset"/>

**Attribute clinicalFinding.content / response / @value**

Namespace	No namespace
Type	restriction of xs:token
Properties	use: required
Facets	enumeration specify enumeration inapplicable enumeration unreported

Source	<pre> &lt;xs:attribute name="value" use="required"&gt;   &lt;xs:simpleType&gt;     &lt;xs:restriction base="xs:token"&gt;       &lt;xs:enumeration value="specify"/&gt;       &lt;xs:enumeration value="inapplicable"/&gt;       &lt;xs:enumeration value="unreported"/&gt;     &lt;/xs:restriction&gt;   &lt;/xs:simpleType&gt; &lt;/xs:attribute&gt; </pre>
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### Attribute **pathologicFinding.content** / **response** / **@value**

Namespace	No namespace
Type	restriction of xs:token
Properties	use: required
Facets	enumeration specify
	enumeration inapplicable
	enumeration unreported
Source	<pre> &lt;xs:attribute name="value" use="required"&gt;   &lt;xs:simpleType&gt;     &lt;xs:restriction base="xs:token"&gt;       &lt;xs:enumeration value="specify"/&gt;       &lt;xs:enumeration value="inapplicable"/&gt;       &lt;xs:enumeration value="unreported"/&gt;     &lt;/xs:restriction&gt;   &lt;/xs:simpleType&gt; &lt;/xs:attribute&gt; </pre>

### Attribute **priorTherapy.content** / **response** / **@value**

Namespace	No namespace
Type	restriction of xs:token
Properties	use: required
Facets	enumeration specify
	enumeration inapplicable
	enumeration unreported
	enumeration chemotherapy
	enumeration radiation
	enumeration surgery
Source	<pre> &lt;xs:attribute name="value" use="required"&gt;   &lt;xs:simpleType&gt;     &lt;xs:restriction base="xs:token"&gt;       &lt;xs:enumeration value="specify"/&gt;       &lt;xs:enumeration value="inapplicable"/&gt;       &lt;xs:enumeration value="unreported"/&gt;       &lt;xs:enumeration value="chemotherapy"/&gt;       &lt;xs:enumeration value="radiation"/&gt;       &lt;xs:enumeration value="surgery"/&gt;     &lt;/xs:restriction&gt;   &lt;/xs:simpleType&gt; &lt;/xs:attribute&gt; </pre>

### Attribute **radiologicFinding.content** / **response** / **@value**

Namespace	No namespace
Type	restriction of xs:token
Properties	use: required
Facets	enumeration specify
	enumeration inapplicable
	enumeration unreported
Source	<pre> &lt;xs:attribute name="value" use="required"&gt;   &lt;xs:simpleType&gt;     &lt;xs:restriction base="xs:token"&gt; </pre>

```

<xs:enumeration value="specify"/>
<xs:enumeration value="inapplicable"/>
<xs:enumeration value="unreported"/>
</xs:restriction>
</xs:simpleType>
</xs:attribute>

```

**Attribute generic.content / @closest**

Namespace	No namespace
Type	union of(val:base.values, val:qualitative.values)
Properties	content: simple
Used by	Element margins.margin.status.result.content/response
Source	<pre> &lt;xs:attribute name="closest"&gt;   &lt;xs:simpleType&gt;     &lt;xs:union memberTypes="val:base.values val:qualitative.values"/&gt;   &lt;/xs:simpleType&gt; &lt;/xs:attribute&gt; </pre>

**Attribute margins.margin.status.result.content / response / @closest**

Namespace	No namespace
Type	union of(val:base.values, val:qualitative.values)
Properties	content: simple
Source	<pre> &lt;xs:attribute name="closest"&gt;   &lt;xs:simpleType&gt;     &lt;xs:union memberTypes="val:base.values val:qualitative.values"/&gt;   &lt;/xs:simpleType&gt; &lt;/xs:attribute&gt; </pre>

**Attribute margins.margin.status.result.content / response / @extent**

Namespace	No namespace
Type	focality.valueset
Properties	content: simple
Source	<pre> &lt;xs:attribute name="extent" type="focality.valueset"/&gt; </pre>

**Attribute margins.margin.status.result.content / response / @value**

Namespace	No namespace														
Type	restriction of xs:token														
Properties	use: required														
Facets	<table> <tr><td>enumeration</td><td>specify</td></tr> <tr><td>enumeration</td><td>inapplicable</td></tr> <tr><td>enumeration</td><td>unreported</td></tr> <tr><td>enumeration</td><td>positive</td></tr> <tr><td>enumeration</td><td>negative</td></tr> <tr><td>enumeration</td><td>equivocal</td></tr> <tr><td>enumeration</td><td>indeterminate</td></tr> </table>	enumeration	specify	enumeration	inapplicable	enumeration	unreported	enumeration	positive	enumeration	negative	enumeration	equivocal	enumeration	indeterminate
enumeration	specify														
enumeration	inapplicable														
enumeration	unreported														
enumeration	positive														
enumeration	negative														
enumeration	equivocal														
enumeration	indeterminate														
Source	<pre> &lt;xs:attribute name="value" use="required"&gt;   &lt;xs:simpleType&gt;     &lt;xs:restriction base="xs:token"&gt;       &lt;xs:enumeration value="specify"/&gt;       &lt;xs:enumeration value="inapplicable"/&gt;       &lt;xs:enumeration value="unreported"/&gt;       &lt;xs:enumeration value="positive"/&gt;       &lt;xs:enumeration value="negative"/&gt;       &lt;xs:enumeration value="equivocal"/&gt;       &lt;xs:enumeration value="indeterminate"/&gt;     &lt;/xs:restriction&gt;   &lt;/xs:simpleType&gt; &lt;/xs:attribute&gt; </pre>														

**Attribute margins.margin.status.result / result / @for**

Namespace	No namespace
Type	restriction of xs:token
Properties	use: required
Facets	enumeration in-situ
	enumeration invasive
	enumeration specify
	enumeration inapplicable
	enumeration unreported
Used by	Element margins.margin.status.result/result
Source	<pre>&lt;xs:attribute name="for" use="required"&gt;   &lt;xs:simpleType&gt;     &lt;xs:restriction base="xs:token"&gt;       &lt;xs:enumeration value="in-situ"/&gt;       &lt;xs:enumeration value="invasive"/&gt;       &lt;xs:enumeration value="specify"/&gt;       &lt;xs:enumeration value="inapplicable"/&gt;       &lt;xs:enumeration value="unreported"/&gt;     &lt;/xs:restriction&gt;   &lt;/xs:simpleType&gt; &lt;/xs:attribute&gt;</pre>

**Attribute distance / distance / @for**

Namespace	No namespace
Type	restriction of xs:token
Properties	use: required
Facets	enumeration in-situ
	enumeration invasive
Used by	Element distance/distance
Source	<pre>&lt;xs:attribute name="for" use="required"&gt;   &lt;xs:simpleType&gt;     &lt;xs:restriction base="xs:token"&gt;       &lt;xs:enumeration value="in-situ"/&gt;       &lt;xs:enumeration value="invasive"/&gt;     &lt;/xs:restriction&gt;   &lt;/xs:simpleType&gt; &lt;/xs:attribute&gt;</pre>

**Attribute distance / distance / @from**

Namespace	No namespace
Type	restriction of xs:token
Properties	content: simple
Facets	enumeration margin
Used by	Element distance/distance
Source	<pre>&lt;xs:attribute name="from"&gt;   &lt;xs:simpleType&gt;     &lt;xs:restriction base="xs:token"&gt;       &lt;xs:enumeration value="margin"/&gt;     &lt;/xs:restriction&gt;   &lt;/xs:simpleType&gt; &lt;/xs:attribute&gt;</pre>

**Attribute margins.margin.status / status / @of**

Namespace	No namespace
Type	margin.status.of.valueset
Properties	use: required

Used by	Element margins.margin.status/status
Source	<code>&lt;xs:attribute name="of" type="margin.status.of.valueset" use="required"/&gt;</code>

**Attribute margins.margin.status / status / @on**

Namespace	No namespace
Type	laterality.valueset
Properties	content: simple
Used by	Element margins.margin.status/status
Source	<code>&lt;xs:attribute name="on" type="laterality.valueset"/&gt;</code>

**Attribute node.status.result.response / response / @value**

Namespace	No namespace
Type	union of(xs:nonNegativeInteger, restriction of xs:token, restriction of xs:token, restriction of xs:token)
Properties	use: required
Used by	Element node.status.result.response/response
Source	<pre> &lt;xs:attribute name="value" use="required"&gt;   &lt;xs:simpleType&gt;     &lt;xs:union memberTypes="xs:nonNegativeInteger"&gt;       &lt;xs:simpleType&gt;         &lt;xs:restriction base="xs:token"&gt;           &lt;xs:enumeration value="specify"/&gt;         &lt;/xs:restriction&gt;       &lt;/xs:simpleType&gt;       &lt;xs:simpleType&gt;         &lt;xs:restriction base="xs:token"&gt;           &lt;xs:enumeration value="inapplicable"/&gt;         &lt;/xs:restriction&gt;       &lt;/xs:simpleType&gt;       &lt;xs:simpleType&gt;         &lt;xs:restriction base="xs:token"&gt;           &lt;xs:enumeration value="unreported"/&gt;         &lt;/xs:restriction&gt;       &lt;/xs:simpleType&gt;     &lt;/xs:union&gt;   &lt;/xs:simpleType&gt; &lt;/xs:attribute&gt; </pre>

**Attribute node.status.result.response / response / @relation**

Namespace	No namespace
Type	relation.valueset
Properties	content: simple
Used by	Element node.status.result.response/response
Source	<code>&lt;xs:attribute name="relation" type="relation.valueset"/&gt;</code>

**Attribute node.status.result / result / @for**

Namespace	No namespace
Type	union of(restriction of xs:token)
Properties	use: required
Used by	Element node.status.result/result
Source	<pre> &lt;xs:attribute name="for" use="required"&gt;   &lt;xs:simpleType&gt;     &lt;xs:union&gt;       &lt;xs:simpleType&gt;         &lt;xs:restriction base="xs:token"&gt;           &lt;xs:enumeration value="total"/&gt;           &lt;xs:enumeration value="negative"/&gt;           &lt;xs:enumeration value="positive"/&gt;         &lt;/xs:restriction&gt;       &lt;/xs:simpleType&gt;     &lt;/xs:union&gt;   &lt;/xs:simpleType&gt; &lt;/xs:attribute&gt; </pre>

```

    </xs:union>
  </xs:simpleType>
</xs:attribute>

```

**Attribute node.status.result / result / @type**

Namespace	No namespace
Type	restriction of xs:token
Properties	content: simple
Facets	enumeration necroinflammatory
Used by	Element node.status.result/result
Source	<pre> &lt;xs:attribute name="type"&gt;   &lt;xs:simpleType&gt;     &lt;xs:restriction base="xs:token"&gt;       &lt;xs:enumeration value="necroinflammatory"/&gt;     &lt;/xs:restriction&gt;   &lt;/xs:simpleType&gt; &lt;/xs:attribute&gt; </pre>

**Attribute nodes.extracapsularExtension.content / response / @value**

Namespace	No namespace
Type	union of(val:base.values, val:qualitative.values)
Properties	use: required
Source	<pre> &lt;xs:attribute name="value" use="required"&gt;   &lt;xs:simpleType&gt;     &lt;xs:union memberTypes="val:base.values val:qualitative.values"/&gt;   &lt;/xs:simpleType&gt; &lt;/xs:attribute&gt; </pre>

**Attribute nodesStatus / status / @of**

Namespace	No namespace
Type	nodes.status.of.valueset
Properties	use: required
Used by	Element nodesStatus/status
Source	<pre> &lt;xs:attribute name="of" type="nodes.status.of.valueset" use="required"/&gt; </pre>

**Attribute nodesStatus / status / @on**

Namespace	No namespace
Type	laterality.valueset
Properties	content: simple
Used by	Element nodesStatus/status
Source	<pre> &lt;xs:attribute name="on" type="laterality.valueset"/&gt; </pre>

**Attribute nodesStatus / status / @technique**

Namespace	No namespace
Type	restriction of xs:token
Properties	content: simple
Facets	enumeration H+E one level
	enumeration H+E multiple level(s)
	enumeration H+E plus immunohistochemistry
	enumeration specify
	enumeration inapplicable

	enumeration unreported
Used by	Element nodesStatus/status
Source	<pre>&lt;xs:attribute name="technique"&gt;   &lt;xs:simpleType&gt;     &lt;xs:restriction base="xs:token"&gt;       &lt;xs:enumeration value="H+E one level"/&gt;       &lt;xs:enumeration value="H+E multiple level(s)"/&gt;       &lt;xs:enumeration value="H+E plus immunohistochemistry"/&gt;       &lt;xs:enumeration value="specify"/&gt;       &lt;xs:enumeration value="inapplicable"/&gt;       &lt;xs:enumeration value="unreported"/&gt;     &lt;/xs:restriction&gt;   &lt;/xs:simpleType&gt; &lt;/xs:attribute&gt;</pre>

**Attribute generic.content / @on**

Namespace	No namespace
Type	laterality.valueset
Properties	content: simple
Used by	Elements site.content/response, specimen.procedure.content/response
Source	<pre>&lt;xs:attribute name="on" type="laterality.valueset"/&gt;</pre>

**Attribute specimen.procedure.content / response / @on**

Namespace	No namespace
Type	laterality.valueset
Properties	content: simple
Source	<pre>&lt;xs:attribute name="on" type="laterality.valueset"/&gt;</pre>

**Attribute specimen.procedure.content / response / @value**

Namespace	No namespace
Type	specimen.procedure.response.valueset
Properties	use: required
Source	<pre>&lt;xs:attribute name="value" type="specimen.procedure.response.valueset" use="required"/&gt;</pre>

**Attribute site.content / response / @on**

Namespace	No namespace
Type	laterality.valueset
Properties	content: simple
Source	<pre>&lt;xs:attribute name="on" type="laterality.valueset"/&gt;</pre>

**Attribute site.content / response / @value**

Namespace	No namespace
Type	restriction of xs:token
Properties	use: required
Facets	enumeration specify enumeration inapplicable enumeration unreported
Source	<pre>&lt;xs:attribute name="value" use="required"&gt;   &lt;xs:simpleType&gt;     &lt;xs:restriction base="xs:token"&gt;       &lt;xs:enumeration value="specify"/&gt;       &lt;xs:enumeration value="inapplicable"/&gt;       &lt;xs:enumeration value="unreported"/&gt;     &lt;/xs:restriction&gt;   &lt;/xs:simpleType&gt; &lt;/xs:attribute&gt;</pre>

```
</xs:simpleType>
</xs:attribute>
```

**Attribute specimen.integrity.content / response / @value**

Namespace	No namespace
Type	restriction of xs:token
Properties	use: required
Facets	enumeration specify
	enumeration inapplicable
	enumeration unreported
	enumeration intact
	enumeration fragmented
Source	<pre>&lt;xs:attribute name="value" use="required"&gt;   &lt;xs:simpleType&gt;     &lt;xs:restriction base="xs:token"&gt;       &lt;xs:enumeration value="specify"/&gt;       &lt;xs:enumeration value="inapplicable"/&gt;       &lt;xs:enumeration value="unreported"/&gt;       &lt;xs:enumeration value="intact"/&gt;       &lt;xs:enumeration value="fragmented"/&gt;     &lt;/xs:restriction&gt;   &lt;/xs:simpleType&gt; &lt;/xs:attribute&gt;</pre>

**Attribute stage.prefix.response / response / @value**

Namespace	No namespace
Type	union of(val:base.values, val:stage.prefix.values)
Properties	use: required
Source	<pre>&lt;xs:attribute name="value" use="required"&gt;   &lt;xs:simpleType&gt;     &lt;xs:union memberTypes="val:base.values val:stage.prefix.values"/&gt;   &lt;/xs:simpleType&gt; &lt;/xs:attribute&gt;</pre>

**Attribute multiplicity / multiplicity / response / @value**

Namespace	No namespace
Type	union of(val:base.values, multiplicity.valueset)
Properties	use: required
Source	<pre>&lt;xs:attribute name="value" use="required"&gt;   &lt;xs:simpleType&gt;     &lt;xs:union memberTypes="val:base.values multiplicity.valueset"/&gt;   &lt;/xs:simpleType&gt; &lt;/xs:attribute&gt;</pre>

**Attribute stage.pT.content / response / @value**

Namespace	No namespace
Type	restriction of xs:token
Properties	use: required
Facets	enumeration specify
	enumeration inapplicable
	enumeration unreported
Source	<pre>&lt;xs:attribute name="value" use="required"&gt;   &lt;xs:simpleType&gt;     &lt;xs:restriction base="xs:token"&gt;       &lt;xs:enumeration value="specify"/&gt;       &lt;xs:enumeration value="inapplicable"/&gt;       &lt;xs:enumeration value="unreported"/&gt;     &lt;/xs:restriction&gt;   &lt;/xs:simpleType&gt; &lt;/xs:attribute&gt;</pre>



	</xs:attribute>
--	-----------------

**Attribute stage.pN.content / response / @value**

Namespace	No namespace
Type	restriction of xs:token
Properties	use: required
Facets	enumeration specify
	enumeration inapplicable
	enumeration unreported
Source	<pre>&lt;xs:attribute name="value" use="required"&gt;   &lt;xs:simpleType&gt;     &lt;xs:restriction base="xs:token"&gt;       &lt;xs:enumeration value="specify"/&gt;       &lt;xs:enumeration value="inapplicable"/&gt;       &lt;xs:enumeration value="unreported"/&gt;     &lt;/xs:restriction&gt;   &lt;/xs:simpleType&gt; &lt;/xs:attribute&gt;</pre>

**Attribute stage.pM.content / response / @value**

Namespace	No namespace
Type	restriction of xs:token
Properties	use: required
Facets	enumeration specify
	enumeration inapplicable
	enumeration unreported
Source	<pre>&lt;xs:attribute name="value" use="required"&gt;   &lt;xs:simpleType&gt;     &lt;xs:restriction base="xs:token"&gt;       &lt;xs:enumeration value="specify"/&gt;       &lt;xs:enumeration value="inapplicable"/&gt;       &lt;xs:enumeration value="unreported"/&gt;     &lt;/xs:restriction&gt;   &lt;/xs:simpleType&gt; &lt;/xs:attribute&gt;</pre>

**Attribute stage.group.response / response / @value**

Namespace	No namespace
Type	stage.group.valueset
Properties	use: required
Source	<xs:attribute name="value" type="stage.group.valueset" use="required"/>

**Attribute tumor.histologicType.content / response / @value**

Namespace	No namespace
Type	union of(val:base.values, tumor.histologicType.valueset)
Properties	use: required
Source	<pre>&lt;xs:attribute name="value" use="required"&gt;   &lt;xs:simpleType&gt;     &lt;xs:union memberTypes="val:base.values tumor.histologicType.valueset"/&gt;   &lt;/xs:simpleType&gt; &lt;/xs:attribute&gt;</pre>

**Attribute tumor.grade.response / response / @value**

Namespace	No namespace
Type	union of(val:base.values, tumor.grade.valueset)
Properties	use: required

Source	<pre>&lt;xs:attribute name="value" use="required"&gt;   &lt;xs:simpleType&gt;     &lt;xs:union memberTypes="val:base.values tumor.grade.valueset"/&gt;   &lt;/xs:simpleType&gt; &lt;/xs:attribute&gt;</pre>
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**Attribute grade / grade / @system**

Namespace	No namespace
Type	tumor.gradeSystem.valueset
Properties	use: required
Used by	Element grade/grade
Source	<pre>&lt;xs:attribute name="system" type="tumor.gradeSystem.valueset" use="required"/&gt;</pre>

**Attribute focus / focus / @type**

Namespace	No namespace						
Type	restriction of xs:token						
Properties	use: required						
Facets	<table> <tr> <td>enumeration</td><td>principal</td></tr> <tr> <td>enumeration</td><td>sole</td></tr> <tr> <td>enumeration</td><td>additional</td></tr> </table>	enumeration	principal	enumeration	sole	enumeration	additional
enumeration	principal						
enumeration	sole						
enumeration	additional						
Used by	Element focus/focus						
Source	<pre>&lt;xs:attribute name="type" use="required"&gt;   &lt;xs:simpleType&gt;     &lt;xs:restriction base="xs:token"&gt;       &lt;xs:enumeration value="principal"/&gt;       &lt;xs:enumeration value="sole"/&gt;       &lt;xs:enumeration value="additional"/&gt;     &lt;/xs:restriction&gt;   &lt;/xs:simpleType&gt; &lt;/xs:attribute&gt;</pre>						

**Attribute diagnosis / @schemaLocation**

Namespace	No namespace
Type	xs:anyURI
Properties	use: required
Used by	Element diagnosis
Source	<pre>&lt;xs:attribute name="schemaLocation" type="xs:anyURI" use="required"/&gt;</pre>

**Attribute diagnosis / @version**

Namespace	No namespace
Type	xs:string
Properties	use: required
Used by	Element diagnosis
Source	<pre>&lt;xs:attribute name="version" type="xs:string" use="required"/&gt;</pre>